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Description

FIELD OF THE INVENTION

[0001] The invention relates to vaccines, specifically nucleic acid vaccines (NAVs). In particular, the invention relates to ribonucleic acid (RNA) vaccines, *i.e.*., mRNA vaccines.

BACKGROUND OF THE INVENTION

[0002] Vaccination is an effective way to provide phrophylactic protection against infectious diseases, including, but not limited to, viral, bacterial, and/or parasitic diseases, such as influenza, AIDS, hepatotisis virus infection, cholera, malaria and tuberculosis, and many other diseases. For example, influenza infections are the seventh leading cause of death in the United States with 200, 000 hospitalizations and 40,000 deaths seen in the United States per year and cause about 3-5 million hospitalizations and about 300,000 to 500,000 deaths worldwide per year. Millions of people receive flu vaccines to protect them from seasonal flu each year. Vaccination can also rapidly prevent the spread of an emerging influenze pandemic.

[0003] A typical vaccine contains an agent that resembles a weakened or dead form of the disease-causing agent, which could be a microorganism, such as bacteria, virus, fungi, parasites, or one or more toxins and/or one or more proteins, for example, surface proteins, (i.e., antigens) of such a microorganism. The antigen or agent in the vaccine can stimulate the body's immune system to recognize the agent as a foreign invader, generate antibodies against it, destroy it and develop a memory of it. The vaccine-induced memory enables the immune system to act quickly to protect the body from any of these agents that it later encounters.

[0004] Vaccine production used in the art *e.g.*, antigen vaccine production, has several stages, including the generation of antigens, antigen purification and inactivation, and vaccine formulation. First, the antigen is generated through culturing viruses in cell lines, growing bacteria in bioreactors, or producing recombinant proteins derived from viruses and bacteria in cell cultures, yeast or bacteria. Recombinant proteins are then purified and the viruses and bacteria are inactivated before they are formulated with adjuvants in vaccines. It has been a challenge to drastically reduce the time and expense associated with current technologies in vaccine development.

[0005] Another obstacle to the development of new vaccine is the constant evolution of most infectious agents, such as viruses and bacteria. Viruses often mutate their surface proteins to generate new antigens which can help them skipping the active immune system that has been immunized by vaccines containing the viruses. In contrast, bacteria acquire and mutate key proteins to evade host defense and effective antibiotic applications.

[0006] For example, influenza A, B and C viruses are the etiological agents of influenza. Hemagglutinin (HA), the major envelop glycoprotein of influenza A and B viruses, or its homologue, hemagglutinin-esterase (HE) in influenza C virus, is the natural reservoir of the viruses. The rapid evolution of the hemagglutinin (HA) protein of the influenza virus results in the constant emergence of new strains, rendering the adaptive immune response of the host only partially protective to new infections. The biggest challenge for therapy and prophylaxis against influenza and other infections using traditional vaccines is the limitation of vaccines in breadth, providing protection only against closely related subtypes. In addition, today's length of the production process inhibits any fast reaction to develop and produce an adapted vaccine in a pandemic situation.

[0007] WO2013/143555 discloses delivery HA-RNA complexed with F4 liposomes which contain DOTMA/DOPE 1:1.

[0008] Petsch et al. (2012) "Protective efficacy of in vitro synthesized, specific mRNA vaccines against influenza A virus infection", Nature Biotechnology , vol. 30, no. 12, pages 1210-1216, and WO2012116810 disclose HA mRNA vaccines. In example 1 of WO2012116810, all mRNAs were complexed with protamine as carrier.

[0009] WO2012/030901 discloses liposomes for delivery of immunogen-encoding RNA which comprise 40mol% of DLinDMA, 10mol% of DSPC, 48% of cholesterol and 2mol% of PEG-DMG.

[0010] It is of great interest to develop new vaccines as well as new approaches to combatting infectious disease and infectious agents.

SUMMARY OF THE INVENTION

[0011] Based on the disclosure that is contained herein, the present invention provides a nucleic acid vaccine, comprising:

one or more messenger RNA (mRNA) polynucleotides having an open reading frame encoding at least one viral antigenic polypeptide,
wherein the one or more mRNA polynucleotides comprise at least one chemical modification, wherein the at least one chemical modification is selected from the group consisting of pseudouridine, N1-methylpseudouridine, 2-

thiouridine, 4-thiouridine, 5-methylcytosine, 2-thio-1-methyl-1-deaza-pseudouridine, 2-thio-1-methyl-pseudouridine, 2-thio-5-aza-uridine , 2-thio-dihydropseudouridine, 2-thio-dihydrouridine, 2-thio-pseudouridine, 4-methoxy-2-thio-pseudouridine, 4-methoxy-pseudouridine, 4-thio-1-methyl-pseudouridine, 4-thio-pseudouridine, 5-aza-uridine, dihydropseudouridine, 5-methoxyuridine, and 2'-O-methyl uridine; and

wherein the one or more mRNA polynucleotides are formulated in a cationic lipid nanoparticle having a molar ratio of 20-60 % cationic lipid, 0.5-15 % PEG-modified lipid, 25-55 % sterol and 5-25 % non-cationic lipid.

[0012] In a further aspect, the present invention provides said vaccine for use as a therapeutic or prophylactic agent.

[0013] In a further aspect, the present invention provides said vaccine for use in a method of preventing or treating a viral infection.

[0014] The present invention and some preferred embodiments thereof are set out in the appended claims.

[0015] In some embodiments, the cationic lipid nanoparticle comprises a molar ratio of about 50% cationic lipid, about 1.5% PEG-modified lipid, about 38.5% cholesterol and about 10% non-cationic lipid. In some embodiments, the cationic lipid nanoparticle comprises a molar ratio of about 55% cationic lipid, about 2.5% PEG lipid, about 32.5% cholesterol and about 10% non-cationic lipid. In some embodiments,the cationic lipid is an ionizable cationic lipid and the non-cationic lipid is a neutral lipid, and the sterol is a cholesterol. In some embodiments, the cationic lipid nanoparticle has a molar ratio of 50:38.5:10:1.5 of cationic lipid: cholesterol: PEG2000-DMG:DSPC.

[0016] In some embodiments, the cationic lipid nanoparticle has a mean diameter of 50-150 nm. In some embodiments, the cationic lipid nanoparticle has a mean diameter of 80-100 nm. In some embodiments, the vaccine includes 1.5mg/mL of RNA polynucleotide and 35-45 mg/mL lipids. In some embodiments, the NAV includes about 2 mg/mL of RNA polynucleotide and about 40 mg/mL lipids.

[0017] In some embodiments, the open reading frame is codon-optimized. In some embodiments, the one or more RNA polynucleotides encode a further antigenic polypeptide. In some embodiments, the further RNA polynucleotide comprises at least one chemical modification and a codon-optimized open reading frame, said open reading frame encoding an antigenic polypeptide.

[0018] In some embodiments, the one or more antigenic polypeptide is selected from those proteins listed in Tables 6-16, Tables 29-30, or fragments thereof. In some embodiments, the open reading frame of the one or more RNA polynucleotides and/or the open reading frame of the second RNA polynucleotide each, independently, encodes an antigenic polypeptide selected from Tables 6-16, Tables 29-30, or fragments thereof. In some embodiments, each of the open reading frame of the one or more RNA polynucleotides is selected from any of the RNA sequences Table 28, or fragments thereof.

[0019] In any of the embodiments provided herein, the infectious agent is a strain of virus selected from the group consisting of adenovirus; Herpes simplex, type 1; Herpes simplex, type 2; encephalitis virus, papillomavirus, Varicella-zoster virus; Epstein-barr virus; Human cytomegalovirus; Human herpes virus, type 8; Human papillomavirus; BK virus; JC virus; Smallpox; polio virus; Hepatitis B virus; Human bocavirus; Parvovirus B19; Human astrovirus; Norwalk virus; coxsackievirus; hepatitis A virus; poliovirus; rhinovirus; Severe acute respiratory syndrome virus; Hepatitis C virus; Yellow Fever virus; Dengue virus; West Nile virus; Rubella virus; Hepatitis E virus; Human Immunodeficiency virus (HIV); Influenza virus; Guanarito virus; Junin virus; Lassa virus; Machupo virus; Sabia virus; Crimean-Congo hemorrhagic fever virus; Ebola virus; Marburg virus; Measles virus; Mumps virus; Parainfluenza virus; Respiratory syncytial virus; Human metapneumovirus; Hendra virus; Nipah virus; Rabies virus; Hepatitis D; Rotavirus; Orbivirus; Colfivirus; Banna virus; Human Enterovirus; Hanta virus; West Nile virus; Middle East Respiratory Syndrome Corona Virus; Japanese encephalitis virus; Vesicular exanthernavirus; and Eastern equine encephalitis.

[0020] In some embodiments, the virus is a strain of Influenza A or Influenza B or combinations thereof. In some embodiments, the strain of Influenza A or Influenza B is associated with birds, pigs, horses, dogs, humans or non-human primates. In some embodiments, the antigenic polypeptide encodes a hemagglutinin protein or fragment thereof. In some embodiments, the hemagglutinin protein is H1, H2, H3, H4, H5, H6, H7, H8, H9, H10, H11, H12, H13, H14, H15, H16, H17, H18, or a fragment thereof. In some embodiments, the hemagglutinin protein does not comprise a head domain (HA1). In some embodiments, the hemagglutinin protein comprises a portion of the head domain (HA1). In some embodiments, the hemagglutinin protein does not comprise a cytoplasmic domain. In some embodiments, the hemagglutinin protein comprises a portion of the cytoplasmic domain. In some embodiments, the truncated hemagglutinin protein. In some embodiments, the truncated hemagglutinin protein comprises a portion of the transmembrane domain. In some embodiments, the amino acid sequence of the hemagglutinin protein or fragment thereof comprises at least 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97% 98%, or 99% identify with any of the amino acid sequences provided in Table 6-14. In some embodiments, the virus is selected from the group consisting of H1N1, H3N2, H7N9, and H10N8. In some embodiments, the antigenic polypeptide is selected from those proteins listed in Tables 6-14, or fragments thereof.

[0021] In some embodiments, the NAV is multivalent. In some embodiments, the open reading frame of the one or more RNA polynucleotides encode at least 2, 3, 4, 5, 6, 7, 8, 9 or 10 antigenic polypeptides. In some embodiments, the open reading frame of the one or more RNA polynucleotides encode at least 10, 15, 20 or 50 antigenic polypeptides. In

some embodiments, the open reading frame of the one or more RNA polynucleotides encode 2-10, 10-15, 15-20, 20-50, 50-100 or 100-200 antigenic polypeptides.

[0022] In the invention, the chemical modification is selected from the group consisting of pseudouridine, N1-methylpseudouridine, 2-thiouridine, 4'-thiouridine, 5-methylcytosine, 2-thio-1-methyl-1-deaza-pseudouridine, 2-thio-1-methyl-pseudouridine, 2-thio-5-aza-uridine , 2-thio-dihydropseudouridine, 2-thio-dihydrouridine, 2-thio-pseudouridine, 4-methoxy-2-thio-pseudouridine, 4-methoxy-pseudouridine, 4-thio-1-methyl-pseudouridine, 4-thio-pseudouridine, 5-aza-uridine, Dihydropseudouridine, 5-methoxyuridine and 2'-O-methyl uridine.. In some embodiments, the RNA polynucleotide includes a second chemical modification wherein said second chemical modification is selected from any of those listed in Tables 22 and 23.

[0023] In some embodiments, the RNA polynucleotide comprises SEQ ID NOs 2459-2621. In some embodiments, the RNA polynucleotide comprises a polynucleotide having at least 80% sequence identity to SEQ ID NOs 2459-2621. In some embodiments, the RNA polynucleotide comprises a polynucleotide encoding an amino acid sequence having at least 90% sequence identity to SEQ ID NO 2254. In some embodiments, the RNA polynucleotide comprises a polynucleotide encoding the amino acid sequence of SEQ ID NO 2254. In some embodiments, the RNA polynucleotide comprises a polynucleotide encoding an amino acid sequence having at least 90% sequence identity to SEQ ID NO 2259. In some embodiments, the RNA polynucleotide comprises a polynucleotide encoding the amino acid sequence of SEQ ID NO 2259. In some embodiments, the RNA polynucleotide comprises a polynucleotide encoding an amino acid sequence having at least 90% sequence identity to SEQ ID NO 2192. In some embodiments, the RNA polynucleotide comprises a polynucleotide encoding the amino acid sequence of SEQ ID NO 2192. In some embodiments, the RNA polynucleotide comprises a polynucleotide encoding an amino acid sequence having at least 90% sequence identity to SEQ ID NO 2200. In some embodiments, the RNA polynucleotide comprises a polynucleotide encoding the amino acid sequence of SEQ ID NO 2200. In some embodiments, the RNA polynucleotide comprises SEQ ID NO 2046. In some embodiments, the RNA polynucleotide comprises a polynucleotide having 80-98% sequence identity to SEQ ID NO 2046. In some embodiments, the RNA polynucleotide comprises SEQ ID NO 2119. In some embodiments, the RNA polynucleotide comprises a polynucleotide having 80-98% sequence identity to SEQ ID NO 2119. In some embodiments, the RNA polynucleotide comprises SEQ ID NO 2054. In some embodiments, the RNA polynucleotide comprises a polynucleotide having 80-98% sequence identity to SEQ ID NO 2054. In some embodiments, the RNA polynucleotide comprises SEQ ID NO 2127. In some embodiments, the RNA polynucleotide comprises a polynucleotide having 80-98% sequence identity to SEQ ID NO 2127.

[0024] The disclosure also provides the vaccine of the invention for use in a method of inducing an antigen specific immune response in a subject comprising administering the vaccines of the invention to the subject in an effective amount to produce an antigen specific immune response. In some embodiments, the antigen specific immune response comprises a T cell response. In some embodiments, the antigen specific immune response comprises a B cell response. In some embodiments, the method of producing an antigen specific immune response involves a single administration of the vaccine. In some embodiments, the method further comprises administering a booster dose of the vaccine. In some embodiments, the vaccine is administered to the subject by intradermal or intramuscular injection.

[0025] In some embodiments, the booster dose of the vaccine is administered to the subject on day twenty one. In some embodiments, a dosage of between 10 ug/kg and 400 ug/kg of the vaccine is administered to the subject. In some embodiments, a dosage of 25 micrograms of the RNA polynucleotide is included in the vaccine administered to the subject. In some embodiments, a dosage of 100 micrograms of the RNA polynucleotide is included in the vaccine administered to the subject. In some embodiments, a dosage of 400 micrograms of the RNA polynucleotide is included in the vaccine administered to the subject. In some embodiments, the RNA polynucleotide accumulates at a 100 fold higher level in the local lymph node in comparison with the distal lymph node.

[0026] The invention also provides the vaccine of the invention for use in methods of preventing or treating influenza viral infection comprising administering to a subject the vaccine of the invention. In some embodiments, the antigen specific immune response comprises a T cell response. In some embodiments, the antigen specific immune response comprises a B cell response. In some embodiments, the method of producing an antigen specific immune response involves a single administration of the vaccine. In some embodiments, the vaccine is administered to the subject by intradermal or intramuscular injection.

[0027] In some embodiments, the NAV polynucleotides may encode one or more polypeptides of an influenza strain as an antigen. Such antigens include, but are not limited to those antigens encoded by the polynucleotides listed in the Tables presented herein. In the table, the GenBank Accession Number or GI Accession Number represents either the complete or partial CDS of the encoded antigen. The NAV polynucleotides may comprise a region of any of the sequences listed in the tables or entire coding region of the mRNA listed. They may comprise hybrid or chimeric regions, or mimics or variants.

[0028] The details of various embodiments of the invention are set forth in the description below. Other features, objects, and advantages of the invention will be apparent from the description and the drawings, and from the claims.

BRIEF DESCRIPTION OF THE DRAWINGS

[0029] The foregoing and other objects, features and advantages will be apparent from the following description of particular embodiments of the invention, as illustrated in the accompanying drawings in which like reference characters refer to the same parts throughout the different views. The drawings are not necessarily to scale, emphasis instead being placed upon illustrating the principles of various embodiments of the invention.

FIG. 1 is a schematic of a polynucleotide construct. FIG. 1A is a schematic of a polynucleotide construct taught in commonly owned co-pending US Patent Application 13/791,922 filed March 9, 2013 (US2013/0259924).
FIG. 1B is a schematic of a linear polynucleotide construct.
FIG. 2 is a schematic of a series of chimeric polynucleotides of the present invention.
FIG. 3 is a schematic of a series of chimeric polynucleotides illustrating various patterns of positional modifications and showing regions analogous to those regions of an mRNA polynucleotide.
FIG. 4 is a schematic of a series of chimeric polynucleotides illustrating various patterns of positional modifications based on Formula I.
FIG. 5 is a is a schematic of a series of chimeric polynucleotides illustrating various patterns of positional modifications based on Formula I and further illustrating a blocked or structured 3' terminus.
FIGs. 6A and 6B are schematics of circular constructs of the present invention.
FIGs. 7A-7B are schematics of circular constructs of the present invention.
FIGs. 8A-8B are schematics of a circular constructs of the present invention. FIG 8A shows a circular construct comprising at least one sensor region and a spacer region. FIG. 8B shows a non-coding circular construct.
FIG. 9 is a schematic of a non-coding circular construct of the present invention.
FIG. 10 shows HA neutralization titres of a chemically modified mRNA influenza vaccine in comparison with protein and unmodified mRNA vaccines.
FIG. 11 shows hemagglutinin inhibition titers in mice following vaccination with different doses and formulations of mRNA encoding the hemagglutinin protein of the H1N1 virus.
FIG. 12A-AD shows percent survival of mice after vaccination and challenge with influenza A/PR/8/34 virus. FIG. 12A shows percent survival at 1 week post challenge. FIG. 12B shows percent survival at 2 weeks post challenge. FIG. 12C shows percent survival at 3 weeks post challenge. FIG. 12D shows percent survival at 4 weeks post challenge.
FIG. 13 shows the mean hemaglutination inhibition titers of mice after vaccination and challenge with influenza A/PR/8/34 virus
FIGs. 14A-14C shows CD4 T cell IFNy cytokine responses. FIG. 14A shows IFNγ production upon H1 protein/peptide stimulation. FIG. 14B shows IFNγ production upon H7 protein/peptide stimulation. FIG. 14C shows IFNγ production upon PMA+ionomycin stimulation.
FIGs. 15A-15D shows IgG production following H1 and H7 protein/peptide stimulation.
FIG. 16 is a graph showing hemagglutinin inhibition titers (HAI) against H10 following administration of the H10N8/N1-methyl pseudouridine /C0 formulation MC3 vaccine at the indicated dosages.
FIG. 17 is a graph showing hemagglutinin inhibition titers (HAI) against H10 following administration of the H10N8/N1-methyl pseudouridine IC1 formulation MC3 vaccine at the indicated dosages.
FIG. 18 is a graph comparing hemagglutinin inhibition titers (HAI) against H7 following administration of 10 μg/dose of the H7N9/C0 formulation compared to the H7N9/C1 formulation
FIG. 19 is a graph of the mean hemagglutinin inhibition titers (HAI) in serum samples from cynomolgus monkey at various time points prior to and after administration of the indicated formulations and dosages.
FIG. 20 is a graph showing the H7N9 viral load in ferrets challenged at day 21 after receiving a single immunization.
FIGs. 21A-21D present mouse survival and HAI titers in mice challenged with a lethal dose following administration of a single dose of mRNA NAV encoding H7N9. FIG. 21A shows survival at day 7 post challenge. FIG. 21B shows survival at day 21 post challenge. FIG. 21C shows survival at day 84 post challenge. FIG. 21D shows HAI titers.
FIG. 22 shows an alignment of amino acid sequences of hemagglutinin proteins from influenza A H7N9 strains relative to a consensus sequence.
FIG. 23 shows an alignment of amino acid sequences of hemagglutinin proteins from influenza A H10N8 strains relative to a consensus sequence.

DETAILED DESCRIPTION

[0030] It is of great interest in the fields of therapeutics, diagnostics, reagents and for biological assays to be able design, synthesize and deliver a nucleic acid, e.g., a ribonucleic acid (RNA) for example, a messenger RNA (mRNA) encoding a peptide or polypeptide of interest inside a cell, whether *in vitro, in vivo, in situ* or *ex vivo,* such as to effect

physiologic outcomes which are beneficial to the cell, tissue or organ and ultimately to an organism. One beneficial outcome is to cause intracellular translation of the nucleic acid and production of at least one encoded peptide or polypeptide of interest.

[0031] Of particular interest, is the ability to design, synthesize and deliver a nucleic acid, *e.g.*, a ribonucleic acid (RNA), for example, a messenger RNA (mRNA), which encodes an antigen, *e.g.*, an antigen derived from an infectious microorganism, for the purpose of vaccination.

[0032] The present invention is set out in the appended claims

Described herein are compositions (including pharmaceutical compositions) and methods for the selection, design, preparation, manufacture, formulation, and/or use of nucleic acid vaccines (NAVs) where at least one component of the NAV is a nucleic acid molecule, *e.g.*, a polynucleotide. In particular, described herein are compositions (including pharmaceutical compositions) and methods for the selection, design, preparation, manufacture, formulation, and/or use of nucleic acid vaccines (NAVs) where at least one component of the NAV is a polynucleotide. In particular, described herein are compositions (including pharmaceutical compositions) and methods for the selection, design, preparation, manufacture, formulation, and/or use of nucleic acid vaccines (NAVs) where at least one component of the NAV is a RNA polynucleotide, *e.g.*, an mRNA polynucleotide which encodes an antigen, *e.g.*, an antigen derived from an infectious microorganism.

[0033] Also provided are systems, processes, devices and kits for the selection, design and/or utilization of the NAVs described herein.

[0034] According to the present invention, the polynucleotides are modified in a manner as to avoid the deficiencies of or provide improvements over other polynucleotide molecules of the art.

[0035] Although attempts have been made to produce functional RNA vaccines, including mRNA vaccines and self-replicating RNA vaccines, the therapeutic efficacy of these RNA vaccines have not yet been fully established. Quite surprisingly, the inventors have discovered a class of formulations for delivering mRNA vaccines in vivo that results in significantly enhanced, and in many respects synergistic, immune responses including enhanced antigen generation and functional antibody production with neutralization capability. These results are achieved even when significantly lower doses of the mRNA are administered in comparison with mRNA doses used in other classes of lipid based formulations. The formulations of the invention have demonstrated significant unexpected in vivo immune responses sufficient to establish the efficacy of functional mRNA vaccines as prophylactic and therapeutic agents.

[0036] The invention involves, in some aspects, the surprising finding that lipid nanoparticle (LNP) formulations significantly enhance the effectiveness of mRNA vaccines, including chemically modified mRNA vaccines. The efficacy of mRNA vaccines formulated in LNP was examined in vivo using several distinct viral antigens and in a variety of different animal models. The results presented herein demonstrate the unexpected superior efficacy of the mRNA vaccines formulated in LNP over other mRNA vaccines formulated in other lipid based carriers as well as over protein antigens.

[0037] In addition to providing an enhanced immune response, the formulations of the invention generate a more rapid immune response following a single dose of antigen than other mRNA or protein based vaccines tested. A study described herein involved intravenous (IV), intramuscular (IM), or intradermal (ID) vaccination of mice, followed by challenge with a lethal dose of virus. In addition to all of the vaccinated animals surviving the lethal dose, significantly stronger early immune responses were observed (anti-viral activity via virus neutralization assay and HA inhibition (HAI)) in comparison to protein antigen and other lipid based formulations (lipoplex). The data demonstrates that as early as 1 week after-vaccination two groups of animals receiving a mRNA-LNP formulation (ID or IM) displayed HAI titers over 40, at 60 and 114, respectively. An HAI titer of greater than 40 is deemed sufficient to protect from a lethal challenge of influenza. The rapid response was unexpected, particularly when compared to the response seen with protein antigen and mRNA vaccines formulated in other lipid carriers (lipoplex), which at one week and even at three weeks following vaccination continued to show ineffective HAI titers of less than 40

[0038] At each of the later time points (3 weeks and 5 weeks), the formulations of the invention continued to provide significantly stronger therapeutic responses than did protein antigen. In fact both chemically unmodified and modified mRNA-LNP formulation administered by IV route had enhanced HAI titers with respect to the protein antigen. By week 3, all of the animals receiving an mRNA-LNP formulation by IM or ID administration displayed HAI activity over 40, as compared to protein antigen, which at one week and three weeks continued to show HAI titers of less than 40. By week 5 a mRNA-LNP formulation administered by ID route had a surprising HAI activity of greater than 10,000, in contrast to the HAI titer of around 400 for protein antigen at that time point. Mice receiving a mRNA-LNP formulation also displayed neutralizing activity of 79-250 (IM) and 250 (ID) by microneutralization assay, in comparison to protein antigen, which had undetectable neutralization activity at that time point. By week 5 following vaccination, five of the six LNP formulated groups showed high neutralizing activity between 789 and 24892. In contrast, the mice vaccinated with protein antigen displayed neutralizing activity in only 3 of 5 mice and ranging only between 79 and 250.

[0039] The mRNA-LNP formulations of the invention also produced quantitatively and qualitatively better immune responses than did mRNA vaccines formulated in a different lipid carrier (lipoplex). At week 5 the mRNA -lipoplex vaccine produced HAI titers of 197, in comparison to those achieved by the mRNA-LNP formulations of the invention (HAI titers

of 635-10,152). At all other time points and for all of the mRNA-lipoplex vaccines, none of the HAI titers reached the critical level of greater than 40. Additionally, the mRNA -lipoplex vaccines did not result in any detectable neutralizing activity in the micro neutralization activity, even as late as five weeks after vaccination.

[0040] The data described herein demonstrate that the formulations of the invention produced significant unexpected improvements over both existing protein antigen vaccines and mRNA vaccine formulations, including: 100% rescue from lethal influenza challenge with rapid onset of protective antibody titers after 1 week and high antibody titers, i.e., 50 fold over unmodified mRNA and 20 fold over the protein vaccine.

[0041] Additionally, the mRNA-LNP formulations of the invention were superior to other lipid formulations even when the dose of mRNA was significantly lower than in the other lipid formulations. For instance, the data described above was generated using 10 pg of mRNA in the mRNA-LNP formulations in contrast to 80 pg of mRNA in the mRNA-lipoplex formulation.

[0042] The formulations of the invention also showed strong efficacy in several non-rodent animal models, including non-human primates. Highly effective vaccination was observed in cynomoglus monkeys and ferrets. Cynomoglus monkeys were vaccinated with various doses of mRNA-LNP formulations (50 μg/dose, 200 μg/dose, 400 μg/dose). Quite surprisingly, the vaccine formulations of the invention at all doses measured significantly reduced viral titers in the lungs of ferrets when exposed to virus just 7 days following single vaccination. Statistically significant increases in antibody titer as measured by HAI and microneutralization were detected as early as 7 days following vaccination and through the entire length of the study (84 days). A single vaccination was able to eliminate all virus in most animals.

[0043] The LNP used in the studies described herein has been used previously to deliver siRNA various in animal models as well as in humans. In view of the observations made in association with the siRNA delivery of LNP formulations, the fact that LNP is useful in vaccines is quite surprising. It has been observed that therapeutic delivery of siRNA formulated in LNP causes an undesirable inflammatory response associated with a transient IgM response, typically leading to a reduction in antigen production and a compromised immune response. In contrast to the findings observed with siRNA, the LNP-mRNA formulations of the invention are demonstrated herein to generate enhanced IgG levels, sufficient for prophylactic and therapeutic methods rather than transient IgM responses.

1. Nucleic Acid Vaccines (NAVs)

[0044] Nucleic Acid Vaccines (NAVs) of the present disclosure comprise one or more polynucleotides, *e.g.*, polynucleotide constructs, which encode one or more wild type or engineered antigens. Exemplary polynucleotides, *e.g.*, polynucleotide constructs, include antigen-encoding RNA polynucleotides, *e.g.*, mRNAs. In the disclosure, polynucleotides of the disclosure, *e.g.*, antigen-encoding RNA polynucleotides, may include at least one chemical modification. References herein to "NAVs of the invention" relate to those as set out in the appended claims.

[0045] NAV compositions of the invention may comprise other components including, but not limited to, tolerizing agents or adjuvants.

Tolerizing agent or composition

[0046] Where auto-immunity mediated side effects occur, tolerizing mRNA and/or such as any of those taught for example in USSN 61/892,556 filed October 18, 2013, and PCT/US2014/61104 filed October 17, 2014 are co-administered with the NAV to induce antigen specific tolerance.

Adjuvants

[0047] Adjuvants or immune potentiators, may also be administered with or in combination with one or more NAVs.

[0048] In one embodiment, an adjuvant acts as a co-signal to prime T-cells and/or B-cells and/or NK cells as to the existence of an infection.

[0049] Advantages of adjuvants include the enhancement of the immunogenicity of antigens, modification of the nature of the immune response, the reduction of the antigen amount needed for a successful immunization, the reduction of the frequency of booster immunizations needed and an improved immune response in elderly and immunocompromised vaccinees. These may be co-administered by any route, e.g., intramusculary, subcutaneous, IV or intradermal injections.

[0050] Adjuvants useful in the present invention may include, but are not limited to, natural or synthetic. They may be organic or inorganic.

[0051] Aduvants may be selected from any of the classes (1) mineral salts, e.g., aluminium hydroxide and aluminium or calcium phosphate gels; (2) emulsions including: oil emulsions and surfactant based formulations, e.g., microfluidised detergent stabilised oil-in-water emulsion, purified saponin, oil-in-water emulsion, stabilised water-in-oil emulsion; (3) particulate adjuvants, e.g., virosomes (unilamellar liposomal vehicles incorporating influenza haemagglutinin), structured complex of saponins and lipids, polylactide co-glycolide (PLG); (4) microbial derivatives; (5) endogenous human immu-

nomodulators; and/or (6) inert vehicles, such as gold particles; (7) microorganism derived adjuvants; (8) tensoactive compunds; (9) carbohydrates; or combinations thereof.

[0052] Adjuvants for DNA nucleic acid vaccines (DNA) have been disclosed in, for example, Kobiyama, et al Vaccines, 2013, 1(3), 278-292. Any of the adjuvants disclosed by Kobiyama may be used in the RNAVs of the present invention.

[0053] Other adjuvants which may be utilized in the RNAVs of the present invention include any of those listed on the web-based vaccine adjuvant database, Vaxjo; http://www.violinet.org/vaxjo/ and described in for example Sayers, et al., J. Biomedicine and Biotechnology, volume 2012 (2012), Article ID 831486,13 pages

[0054] Selection of appropriate adjuvants will be evident to one of ordinary skill in the art. Specific adjuvants may include,without limitation, cationic liposome-DNA complex JVRS-100, aluminum hydroxide vaccine adjuvant, aluminum phosphate vaccine adjuvant, aluminum potassium sulfate adjuvant, alhydrogel, ISCOM(s)™, Freund's Complete Adjuvant, Freund's Incomplete Adjuvant, CpG DNA Vaccine Adjuvant, Cholera toxin, Cholera toxin B subunit, Liposomes, Saponin Vaccine Adjuvant, DDA Adjuvant, Squalene-based Adjuvants, Etx B subunit Adjuvant, IL-12 Vaccine Adjuvant, LTK63 Vaccine Mutant Adjuvant, TiterMax Gold Adjuvant, Ribi Vaccine Adjuvant, Montanide ISA 720 Adjuvant, *Corynebacterium*-derived P40 Vaccine Adjuvant, MPL™ Adjuvant, AS04, AS02, Lipopolysaccharide Vaccine Adjuvant, Muramyl Dipeptide Adjuvant, CRL1005, Killed *Corynebacterium* parvum Vaccine Adjuvant, Montanide ISA 51, *Bordetella pertussis* component Vaccine Adjuvant, Cationic Liposomal Vaccine Adjuvant, Adamantylamide Dipeptide Vaccine Adjuvant, Arlacel A, VSA-3 Adjuvant, Aluminum vaccine adjuvant, Polygen Vaccine Adjuvant, Adjumer™, Algal Glucan, Bay R1005, Theramide®, Stearyl Tyrosine, Specol, Algammulin, Avridine®, Calcium Phosphate Gel, CTA1-DD gene fusion protein, DOC/Alum Complex, Gamma Inulin, Gerbu Adjuvant, GM-CSF, GMDP, Recombinant hIFN-gamma/Interferon-g, Interleukin-1β, Interleukin-2, Interleukin-7, Sclavo peptide, Rehydragel LV, Rehydragel HPA, Loxoribine, MF59, MTP-PE Liposomes, Murametide, Murapalmitine, D-Murapalmitine, NAGO, Non-Ionic Surfactant Vesicles, PMMA, Protein Cochleates, QS-21, SPT (Antigen Formulation), nanoemulsion vaccine adjuvant, AS03, Quil-A vaccine adjuvant, RC529 vaccine adjuvant, LTR192G Vaccine Adjuvant, *E. coli* heat-labile toxin, LT, amorphous aluminum hydroxyphosphate sulfate adjuvant, Calcium phosphate vaccine adjuvant, Montanide Incomplete Seppic Adjuvant, Imiquimod, Resiquimod, AF03, Flagellin, Poly(I:C), ISCOMATRIX®, Abisco-100 vaccine adjuvant, Albumin-heparin microparticles vaccine adjuvant, AS-2 vaccine adjuvant, B7-2 vaccine adjuvant, DHEA vaccine adjuvant, Immunoliposomes Containing Antibodies to Costimulatory Molecules, SAF-1, Sendai Proteoliposomes, Sendai-containing Lipid Matrices, Threonyl muramyl dipeptide (TMDP), Ty Particles vaccine adjuvant, Bupivacaine vaccine adjuvant, DL-PGL (Polyester poly (DL-lacfide-co-glycolide)) vaccine adjuvant, IL-15 vaccine adjuvant, LTK72 vaccine adjuvant, MPL-SE vaccine adjuvant, non-toxic mutant E112K of Cholera Toxin mCT-E112K, and/or Matrix-S.

[0055] Other adjuvants which may be co-administered with the NAVs of the invention include, but are not limited to interferons, TNF-alpha, TNF-beta, chemokines such as CCL21, eotaxin, HMGB1, SA100-8alpha, GCSF, GMCSF, granulysin, lactoferrin, ovalbumin, CD-40L, CD28 agonists, PD-1, soluble PD1, L1 or L2, or interleukins such as IL-1, IL-2, IL-4, IL-6, IL-7, IL-10, IL-12, IL-13, IL-21, IL-23, IL-15, IL-17, and IL-18. These may be administered with the NAV on the same encoded polynucleotide, e.g., polycistronic, or as separate mRNA encoding the adjuvant and antigen.

Valency

[0056] NAVs of the present invention may vary in their valency. Valency refers to the number of antigenic components in the NAV or NAV polynucleotide (e.g., RNA polynucleotide) or polypeptide. In some embodiments, the NAVs are monovalent. In some embodiments, the NAVs are divalent. In some embodments the NAVs are trivalent. In some embodiments the NAVs are multi-valent. Multivalent vaccines may comprise 2, 3, 4, 5, 6, 7, 8, 9, 10, 11,12,13,14, 15, 16, 17, 18, 19, 20, or more antigens or antigenic moieties (*e.g.*, antigenic peptides, etc.). The antigenic components of the NAVs may be on a single polynucleotide or on separate polynucleotides.

Therapeutics

[0057] The NAVs of the present invention can be used as therapeutic or prophylactic agents. They are provided for use in medicine and/or for the priming of immune effector cells, e.g., stimulate/transfect PBMCs ex vivo and re-infuse the activated cells. For example, a NAV described herein can be administered to a subject, wherein the polynucleotides is translated *in vivo* to produce an antigen. Provided are compositions, methods, kits, and reagents for diagnosis, treatment or prevention of a disease or condition in humans and other mammals. The active therapeutic agents of the disclosure include NAVs, cells containing NAVs or polypeptides translated from the polynucleotides contained in said NAVs.

[0058] Provided herein are methods of inducing translation of a polypeptide (*e.g.*, antigen or immunogen) in a cell, tissue or organism using the polynucleotides of the NAVs described herein. Such translation can be *in vivo, , ex vivo, in culture,* or *in vitro.* The cell, tissue or organism is contacted with an effective amount of a composition containing a NAV which contains a polynucletotide that has at least one a translatable region encoding the polypeptide of intereste

(*e.g.*, antigen or immunogen).

[0059] An "effective amount" of the NAV composition is provided based, at least in part, on the target tissue, target cell type, means of administration, physical characteristics of the polynucleotide (e.g., size, and extent of modified nucleosides) and other components of the NAV, and other determinants. In general, an effective amount of the NAV composition provides an induced or boosted immune response as a function of antigen production in the cell, preferably more efficient than a composition containing a corresponding unmodified polynucleotide encoding the same antigen. Increased antigen production may be demonstrated by increased cell transfection (i.e., the percentage of cells transfected with the NAV), increased protein translation from the polynucleotide, decreased nucleic acid degradation (as demonstrated, e.g., by increased duration of protein translation from a modified polynucleotide), or altered innate immune response of the host cell.

[0060] Aspects of the disclosure are directed to methods of inducing *in vivo* translation of a polypeptide antigen in a mammalian subject in need thereof. Therein, an effective amount of a NAV composition containing a polynucleotide that has at least one structural or chemical modification and a translatable region encoding the polypeptide (*e.g.*, antigen or immunogen) is administered to the subject using the delivery methods described herein. The polynucleotide is provided in an amount and under other conditions such that the polynucleotide is translated in the cell. The cell in which the polynucleotide is localized, or the tissue in which the cell is present, may be targeted with one or more than one rounds of NAV administration.

[0061] In certain instances, the administered NAVs comprising polynucleotides directs production of one or more polypeptides that provide a functional immune system-related activity which is substantially absent in the cell, tissue or organism in which the polypeptide is translated. For example, the missing functional activity may be enzymatic, structural, or gene regulatory in nature. In related instances, the administered polynucleotides directs production of one or more polypeptides that increases (e.g., synergistically) a functional activity related to the immune system which is present but substantially deficient in the cell in which the polypeptide is translated.

[0062] Additionally, the polypeptide antagonizes, directly or indirectly, the activity of a biological moiety present in, on the surface of, or secreted from the cell. Examples of antagonized biological moieties include lipids (e.g., cholesterol), a lipoprotein (e.g., low density lipoprotein), a nucleic acid, a carbohydrate, a protein toxin such as shiga and tetanus toxins, or a small molecule toxin such as botulinum, cholera, and diphtheria toxins. Additionally, the antagonized biological molecule may be an endogenous protein that exhibits an undesirable activity, such as a cytotoxic or cytostatic activity.

[0063] The proteins described herein may be engineered for localization within the cell, potentially within a specific compartment such as the cytoplasms or nucleus, or are engineered for secretion from the cell or translocation to the plasma membrane of the cell.

[0064] In some instances, polynucleotides of the NAVs and their encoded polypeptides in accordance with the present disclosure may be used for treatment of any of a variety of diseases, disorders, and/or conditions, including but not limited to viral infections (*e.g.*, influenza, HIV, HCV, RSV), parasitic intentions or bacterial infections.

[0065] The subject to whom the therapeutic agent may be administered suffers from or may be at risk of developing a disease, disorder, or deleterious condition. Provided are methods of identifying, diagnosing, and classifying subjects on these bases, which may include clinical diagnosis, biomarker levels, genome-wide association studies (GWAS), and other methods known in the art.

[0066] The agents can be administered simultaneously, for example in a combined unit dose (*e.g.*, providing simultaneous delivery of both agents). The agents can also be administered at a specified time interval, such as, but not limited to, an interval of minutes, hours, days or weeks. Generally, the agents may be concurrently bioavailable, *e.g.*, detectable, in the subject. In some instances, the agents may be administered essentially simultaneously, for example two unit dosages administered at the same time, or a combined unit dosage of the two agents. In other instances, the agents may be delivered in separate unit dosages. The agents may be administered in any order, or as one or more preparations that includes two or more agents. In a preferred instance, at least one administration of one of the agents, *e.g.*, the first agent, may be made within minutes, one, two, three, or four hours, or even within one or two days of the other agent, *e.g.*, the second agent. In some instances, combinations can achieve synergistic results, *e.g.*, greater than additive results, *e.g.*, at least 25,50,75,100,200,300,400, or 500% greater than additive results.

Modulation of the Immune Response

Activation of the immune response

[0067] According to the present disclosure, the NAVs comprising the polynucleotides disclosed herein, e.g., comprising RNA polynucleotides, may act as a single composition as a vaccine. As used herein, a "vaccine" refers to a composition, for example, a substance or preparation that stimulates, induces, causes or improves immunity in an organism, *e.g.*, an animal organism, for example, a mammalian organism (*e.g.*, a human) Preferably, a vaccine provides immunity against one or more diseases or disorders in the organism, including prophylactic and/or therapeutic immunity. Exemplary

vaccines includes one or more agents that resembles an infectious agent, *e.g.*, a disease-causing microorganism, and can be made, for example, from live, attenuated, modified, weakened or killed forms of disease-causing microorganisms, or antigens derived therefrom, including combinations of antigenic components In exemplary instances, a vaccine stimulates, induces causes or improves immunity in an organism or causes or mimics infection in the organism without inducing any disease or disorder. A vaccine introduces an antigen into the tissues, extracellular space or cells of a subject and elicits an immune response, thereby protecting the subject from a particular disease or pathogen infection. The polynucleotides of the present disclosure may encode an antigen and when the polynucleotides are expressed in cells, a desired immune reponse is achieved.

[0068] NAVs may be administered prophylactically or therapeutically as part of an active immunization scheme to healthy individuals or early in infection during the incubation phase or during active infection after onset of symptoms.

[0069] The NAVs of the present disclosure may also be administered as a second line treatment after the standard first line treatments such as antibiotics and antivirals have failed to induce passive immunity. In this regard, the NAVs of the present invention are useful in settings where resistance to first line treatments has developed and disease persists and induces chronic disease.

[0070] The use of RNA in or as a vaccine overcomes the disadvantages of conventional genetic vaccination involving incorporating DNA into cells in terms of safeness, feasibility, applicability, and effectiveness to generate immune responses. RNA molecules are considered to be significantly safer than DNA vaccines, as RNAs are more easily degraded. They are cleared quickly out of the organism and cannot integrate into the genome and influence the cell's gene expression in an uncontrollable manner. It is also less likely for RNA vaccines to cause severe side effects like the generation of autoimmune disease or anti-DNA antibodies (Bringmann A. et al., Journal of Biomedicine and Biotechnology (2010), vol. 2010, article ID623687). Transfetion with RNA requires only insertion into the cell's cytoplasm, which is easier to achieve than into the nucleus. Howerver, RNA is susceptible to RNase degradation and other natural decomposition in the cytoplasm of cells.

[0071] Various attempts to increase the stability and shelf life of RNA vaccines. US 2005/0032730 to Von Der Mulbe et al. discloses improving the stability of mRNA vaccine compositions by increasing G(guanosine)/C(cytosine) content of the mRNA molecules. US 5580859 to Felgner et al. teaches incorporating polynucleotide sequences coding for regulatory proteins that binds to and regulates the stabilities of mRNA. While not wishing to be bound by theory, it is believed that the polynucleotides vaccines (NAVs) of the invention will result in improved stability and therapeutic efficacy due at least in part to the specificity, purity and selectivity of the construct designs.

[0072] Additionally, certain modified nucleosides, or combinations thereof, when introduced into the polynucleotides of the NAVs of the invention will activate the innate immune response. Such activating molecules are useful as adjuvants when combined with polypeptides and/or othervaccines. In certain instances, the activating molecules contain a translatable region which encodes for a polypeptide sequence useful as a vaccine, thus providing the ability to be a self-adjuvant.

[0073] In one instance, the polynucleotides of the NAVs of the present disclosure may be used in the prevention, treatment and diagnosis of diseases and physical disturbances caused by infectious agents. The polynucleotide of the present disclosure may encode at least one polypeptide of interest (antigen) and may be provided to an individual in order to stimulate the immune system to protect against the disease-causing agents. As a non-limiting example, the biological activity and/or effect from an antigen or infectious agent may be inhibited and/or abolished by providing one or more polynucleotides which have the ability to bind and neutralize the antigen and/or infectious agent.

[0074] As a non-limiting example, the polynucleotides encoding an immunogen may be delivered to cells to trigger multiple innate response pathways (see International Pub. No. WO2012006377 and US Patent Publication No. US20130177639). As another non-limiting example, the polynucleotides of the NAVs of the present disclosure encoding an immunogen may be delivered to a vertebrate in a dose amount large enough to be immunogenic to the vertebrate (see International Pub. No. WO2012006372 and WO2012006369 and US Publication No. US20130149375 and US20130177640).

[0075] A non-limiting list of infectious diseases that the polynucleotide vaccines may treat includes, viral infectious diseases such as AIDS (HIV), HIV resulting in mycobacterial infection, AIDS related Cacheixa, AIDS related Cytomegalovirus infection, HIV-associated nephropathy, Lipodystrophy, AID related cryptococcal meningitis, AIDS related neutropaenia, Pneumocysitis jiroveci (Pneumocystis carinii) infections, AID related toxoplasmosis, hepatitis A, B, C, D or E, herpes, herpes zoster (chicken pox), German measles (rubella virus), yellow fever, dengue fever etc. (flavi viruses), flu (influenza viruses), haemorrhagic infectious diseases (Marburg or Ebola viruses), bacterial infectious diseases such as Legionnaires' disease (Legionella), gastric ulcer (Helicobacter), cholera (Vibrio), E. coli infections, staphylococcal infections, salmonella infections or streptococcal infections, tetanus (Clostridium tetani), protozoan infectious diseases (malaria, sleeping sickness, leishmaniasis, toxoplasmosis, i.e. infections caused by plasmodium, trypanosomes, leishmania and toxoplasma), diphtheria, leprosy, measles, pertussis, rabies, tetanus, tuberculosis, typhoid, varicella, diarrheal infections such as Amoebiasis, Clostridium difficile-associated diarrhea (CDAD), Cryptosporidiosis, Giardiasis, Cyclosporiasis and Rotaviral gastroenteritis, encephalitis such as Japanese encephalitis, Wester equine encephalitis and Tick-

borne encephalitis (TBE), fungal skin diseases such as candidiasis, onychomycosis, Tinea captis/scal ringworm, Tinea corporis/body ringworm, Tinea cruris/jock itch, sporotrichosis and Tinea pedis/Athlete's foot, Meningitis such as Haemophilus influenza type b (Hib), Meningitis, viral, meningococcal infections and pneumococcal infection, neglected tropical diseases such as Argentine haemorrhagic fever, Leishmaniasis, Nematode/roundworm infections, Ross river virus infection and West Nile virus (WNV) disease, Non-HIV STDs such as Trichomoniasis, Human papillomavirus (HPV) infections, sexually transmitted chlamydial diseases, Chancroid and Syphilis, Non-septic bacterial infections such as cellulitis, lyme disease, MRSA infection, pseudomonas, staphylococcal infections, Boutonneuse fever, Leptospirosis, Rheumatic fever, Botulism, Rickettsial disease and Mastoiditis, parasitic infections such as Cysticercosis, Echinococcosis, Trematode/Fluke infections, Trichinellosis, Babesiosis, Hypodermyiasis, Diphyllobothriasis and Trypanosomiasis, respiratory infections such as adenovirus infection, aspergillosis infections, avian (H5N1) influenza, influenza, RSV infections, severe acute respiratory syndrome (SARS), sinusitis, Legionellosis, Coccidioidomycosis and swine (H1N1) influenza, sepsis such as bacteraemia, sepsis/septic shock, sepsis in premature infants, urinary tract infection such as vaginal infections (bacterial), vaginal infections (fungal) and gonococcal infection, viral skin diseases such as B19 parvovirus infections, warts, genital herpes, orofacial herpes, shingles, inner ear infections, fetal cytomegalovirus syndrome, foodborn illnesses such as brucellosis (Brucella species), Clostridium perfringens (Epsilon toxin), E. Coli 0157:H7 (Escherichia coli), Salmonellosis (Salmonella species), Shingellosis (Shingella), Vibriosis and Listeriosis, bioterrorism and potential epidemic diseases such as Ebola haemorrhagic fever, Lassa fever, Marburg haemorrhagic fever, plague, Anthrax Nipah virus disease, Hanta virus, Smallpox, Glanders (Burkholderia mallei), Melioidosis (Burkholderia pseudomallei), Psittacosis (Chlamydia psittaci), Q fever (Coxiella burnetii), Tularemia (Fancisella tularensis), rubella, mumps and polio.

[0076] NAVs of the present invention may be utilized in various settings depending on the prevalence of the infection or the degree or level of unmet medical need. As a non-limiting example, the NAVs of the present invention may be utilized to treat and/or prevent influenza infection, i.e. diseases and conditions related to influenza virus infection (seasonal and pandemic).

[0077] Symptoms of the influenza infection include dry cough, fever, chills, myalgias progressing to respiratory failure and the risk of secondary bacterial infections (e.g., MRSA). Seasonal influenza is ubiquitous and consists of three principal strains (A [H1N1], A [H3N2], and B), which are covered by the annual vaccine. Pandemic flu occurs because the viruses' unique reassortment ability allowing antigenic shift as well as transfer between avian and swine flu strains. One emerging concern in Southeast Asia is the pandemic potential of several new strains. Such pandemic outbreaks have a high mortality rate with few available treatments. Anti-virals only provide symptomatic relief and must be given in the first 48 hours.

[0078] The NAVs of the present invention have superior properties in that they produce much larger antibody titers, produce responses early than commercially available anti-virals and may be administered after the critical 48 hour period while retaining efficacy.

[0079] While not wishing to be bound by theory, the inventors hypothesize that the NAVs of the invention, as mRNA polynucleotides, are better designed to produce the appropriate protein conformation on translation as the NAVs co-opt natural cellular machinery. Unlike traditional vaccines which are manufactured ex vivo and may trigger unwanted cellular responses, the NAVs are presented to the cellular system in a more native fashion. Adding to the superior effects may also involve the formulations utilized which may neither serve to shield nor traffic the NAVs.

[0080] According to the present invention, NAVs represent a tailored active vaccine that not only can prevent infection but can limit transmission of influenza.

[0081] In some instances, the NAVs may be used to prevent pandemic influenza by reacting to emerging new strains with the very rapid NAV-based vaccine production process. In some embodiments, new NAV for treating or prophylactically preventing influenza outbreaks, including for emerging strains (e.g., H10N8), may be produced in less than six weeks, from the time of antigen identification to available vaccine.

[0082] In some instances a single injection of a single antigen encoding NAV polynucleotide may provide protection for an entire flu season.

[0083] The polynucleotides of the NAVs of the invention are not self-replicating RNA. Self-replicating RNA have been described, for instance in US Pub. No. US20110300205 and International Pub. No. WO2011005799 and WO2013055905.

[0084] In one embodiment, the polynucleotides of the NAVs of the invention may encode amphipathic and/or immunogenic amphipathic peptides.

[0085] In on embodiment, a formulation of the polynucleotides of the NAVs of the invention may further comprise an amphipathic and/or immunogenic amphipathic peptide. As a non-limiting example, the polynucleotides comprising an amphipathic and/or immunogenic amphipathic peptide may be formulated as described in US. Pub. No. US20110250237 and International Pub. Nos. WO2010009277 and WO2010009065.

[0086] In one embodiment, the polynucleotides of the NAVs of the invention may be immunostimultory. As a non-limiting example, the polynucleotides may encode all or a part of a positive-sense or a negative-sense stranded RNA virus genome (see International Pub No. WO2012092569 and US Pub No. US20120177701). In another non-limiting

example, the immunostimultory polynucleotides of the present invention may be formulated with an excipient for administration as described herein and/or known in the art (see International Pub No. WO2012068295 and US Pub No. US20120213812). The polynucleotides may further comprise a sequence region encoding a cytokine that promotes the immune response, such as a monokine, lymphokine, interleukin or chemokine, such as IL-1, IL-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-8, IL-9, IL-10, IL-12, INF-α, INF-y, GM-CFS, LT-o, or growth factors such as hGH.

[0087] In one embodiment, the response of the vaccine formulated by the methods described herein may be enhanced by the addition of various compounds to induce the therapeutic effect. As a non-limiting example, the vaccine formulation may include a MHC II binding peptide or a peptide having a similar sequence to a MHC II binding peptide (see International Pub Nos. WO2012027365, WO2011031298 and US Pub No. US20120070493, US20110110965). As another example, the vaccine formulations may comprise modified nicotinic compounds which may generate an antibody response to nicotine residue in a subject (see International Pub No. WO2012061717 and US Pub No. US20120114677).

[0088] In one embodiment, the effective amount of the polynucleotides of the NAVs of the invention provided to a cell, a tissue or a subject may be enough for immune prophylaxis.

[0089] In one instance, the polynucleotides of the NAVs of the invention may be administrated with other prophylactic or therapeutic compounds. As a non-limiting example, the prophylactic or therapeutic compound may be an adjuvant or a booster. As used herein, when referring to a prophylactic composition, such as a vaccine, the term "booster" refers to an extra administration of the prophylactic composition. A booster (or booster vaccine) may be given after an earlier administration of the prophylactic composition. The time of administration between the intial administration of the prophylactic composition and the booster may be, but is not limited to, 1 minute, 2 minutes, 3 minutes, 4 minutes, 5 minutes, 6 minutes, 7 minutes, 8 minutes, 9 minutes, 10 minutes, 15 minutes, 20 minutes 35 minutes, 40 minutes, 45 minutes, 50 minutes, 55 minutes, 1 hour, 2 hours, 3 hours, 4 hours, 5 hours, 6 hours, 7 hours, 8 hours, 9 hours, 10 hours, 11 hours, 12 hours, 13 hours, 14 hours, 15 hours, 16 hours, 17 hours, 18 hours, 19 hours, 20 hours, 21 hours, 22 hours, 23 hours, 1 day, 36 hours, 2 days, 3 days, 4 days, 5 days, 6 days, 1 week, 10 days, 2 weeks, 3 weeks, 1 month, 2 months, 3 months, 4 months, 5 months, 6 months, 7 months, 8 months, 9 months, 10 months, 11 months, 1 year, 18 months, 2years, 3years, 4years, 5years, 6years, 7years, 8 years, 9 years, 10 years, 11 years, 12 years, 13 years, 14years, 15 years, 16 years, 17 years, 18 years, 19 years, 20 years, 25 years, 30 years, 35 years, 40 years, 45 years, 50 years, 55 years, 60 years, 65 years, 70 years, 75 years, 80 years, 85 years, 90 years, 95 years or more than 99 years.

[0090] In one instance, the polynucleotides of the NAVs of the invention may be administered intranasally similar to the administration of live vaccines. In another aspect the polynucleotide may be administered intramuscularly or intradermally similarly to the administration of inactivated vaccines known in the art.

[0091] In one instance, the NAVs of the invention may be used to protect against and/or prevent the transmission of an emerging or engineered threat which may be known or unknown.

[0092] In another embodiment, the NAVs may be formulated by the methods described herein. In one aspect, the formulation may comprise a NAV or polynucleotide which can can have a therapetutic and/or prophylactic effect on more than one disease, disorder or condition.

[0093] In addition, the NAV antibodies of the present invention may be used for research in many applications, such as, but not limited to, identifying and locating intracellular and extracellular proteins, protein interaction, signal pathways and cell biology.

[0094] In another instance, the NAV may be used in to reduce the risk or inhibit the infection of influenza viruses such as, but not limited to, the highly pathogenic avian influenza virus (such as, but not limited to, H5N1 subtype) infection and human influenza virs (such as, but not limited to, H1N1 subtype and H3N2 subtype) infection. The polynucleotide described herein which may encode any of the protein sequences described in US Patent No. 8470771 may be used in the treatment or to reduce the risk of an influenza infection.

[0095] In one instance, the NAV may be used as a vaccine and may further comprise an adjuvant which may enable the vaccine to elicit a higher immune response. As a non-limiting example, the adjuvant could be a sub-micron oil-in-water emulsion which can elicit a higher immune response in human pediatric populations (see e.g., the adjuvanted vaccines described in US Patent Publication No. US20120027813 and US Patent No. US8506966).

II. Infectious Agents and Antigens

[0096] NAVs of the present invention may be used to protect, treat or cure infection arising from contact with an infectious agent. In the invention, infectious agents are viruses.

A. Managing infection

Antibiotic Combinations

[0097] In one instance, the NAVs of the present invention, e.g., NAVs comprising one or more antigen-encoding

polynucleotides of the present disclosure, may be administered in conjunction with one or more antibiotics.

Antibacterial agents

[0098] Anti-bacterial agents include, but are not limited to, aminoglycosides (*e.g.*, amikacin (AMIKIN®), gentamicin (GARAMYCIN®), kanamycin (KANTREX®), neomycin (MYCIFRADIN®), netilmicin (NETROMYCIN®), tobramycin (NEBCIN®), Paromomycin (HUMATIN®)), ansamycins (*e.g.*, geldanamycin, herbimycin), carbacephem (*e.g.*, loracarbef (LORABID®), Carbapenems (*e.g.*, ertapenem (INVANZ®), doripenem (DORIBAX®), imipenem/cilastatin (PRIMAXIN®), meropenem (MERREM®), cephalosporins (first generation) (*e.g.*, cefadroxil (DURICEF®), cefazolin (ANCEF®), cefalotin or cefalothin (KEFLIN®), cefalexin (KEFLEX®), cephalosporins (second generation) (*e.g.*, cefaclor (CECLOR®), cefamandole (MANDOL®), cefoxitin (MEFOXIN®), cefprozil (CEFZIL®), cefuroxime (CEFTIN®, ZINNAT®)), cephalosporins (third generation) (*e.g.*, cefixime (SUPRAX®), cefdinir (OMNICEF®, CEFDIEL®), cefditoren (SPECTRACEF®), cefoperazone (CEFOBID®), cefotaxime (CLAFORAN®), cefpodoxime (VANTIN®), ceftazidime (FORTAZ®), ceftibuten (CEDAX®), ceffizoxime (CEFIZOX®), ceftriaxone (ROCEPHIN®)), cephalosporins (fourth generation) (*e.g.*, cefepime (MAXIPIME®)), cephalosporins (fifth generation) (*e.g.*, ceffobiprole (ZEFTERA®)), glycopeptides (*e.g.*, teicoplanin (TARGOCID®), vancomycin (VANCOCIN®), telavancin (VIBATIV®)), lincosamides (*e.g.*, clindamycin (CLEOCIN®), lincomycin (LINCOCIN®)), lipopeptide (*e.g.*, daptomycin (CUBICIN®)), macrolides (*e.g.*, azithromycin (ZITHROMAX®, SUMAMED®, ZITROCIN®), clarithromycin (BIAXIN®), dirithromycin (DYNABAC®), erythromycin (ERYTHOCIN®, ERYTHROPED®), roxithromycin, troleandomycin (TAO®), telithromycin (KETEK®), spectinomycin (TROBICIN®)), monobactams (*e.g.*, aztreonam (AZACTAM®)), nitrofurans (*e.g.*, furazolidone (FUROXONE®), nitrofurantoin (MACRODANTIN®, MACROBID®)), penicillins (*e.g.*, amoxicillin (NOVAMOX®, AMOXIL®), ampicillin (PRINCIPEN®), azlocillin, carbenicillin (GEOCILLIN®), cloxacillin (TEGOPEN®), dicloxacillin (DYNAPEN®), flucloxacillin (FLOXAPEN®), mezlocillin (MEZLIN®), methicillin (STAPHCILLIN®), nafcillin (UNIPEN®), oxacillin (PROSTAPHLIN®), penicillin G (PENTIDS®), penicillin V (PENVEE-K®), piperacillin (PIPRACIL®), temocillin (NEGABAN®), ticarcillin (TICAR®)), penicillin combinations (*e.g.*, amoxicillin/clavulanate (AUGMENTIN®), ampicillin/sulbactam (UNASYN®), piperacillin/tazobactam (ZOSYN®), ticarcillin/clavulanate (TIMENTIN®)), polypeptides (*e.g.*, bacitracin, colistin (COLY-MYCIN-S®), polymyxin B, quinolones (*e.g.*, ciprofloxacin (CIPRO®, CIPROXIN®, CIPROBAY®), enoxacin (PENETREX®), gatifloxacin (TEQUIN®), levofloxacin (LEVAQUIN®), lomefloxacin (MAXAQUIN®), moxifloxacin (AVELOX®), nalidixic acid (NEGGRAM®), norfloxacin (NOROXIN®), ofloxacin (FLOXIN®, OCUFLOX®), trovafloxacin (TROVAN®), grepafloxacin (RAXAR®), sparfloxacin (ZAGAM®), femafloxacin (OMNIFLOX®)), sulfonamides (*e.g.*, mafenide (SULFAMYLON®), sulfonamidochrysoidine (PRONTOSIL®), sulfacetamide (SULAMYD®, BLEPH-10®), sulfadiazine (MICRO-SULFON®), silver sulfadiazine (SILVADENE®), sulfamethizole (THIOSULFIL FORTE®), sulfamethoxazole (GANTANOL®), sulfanilimide, sulfasalazine (AZULFIDINE®), sulfisoxazole (GANTRISIN®), trimethoprim (PROLOPRIM®), TRIMPEX®), trimethoprim-sulfamethoxazole (co-trimoxazole) (TMP-SMX) (BACTRIM®, SEPTRA®)), tetracyclines (*e.g.*, demeclocycline (DECLOMYCIN®), doxycycline (VIBRAMYCIN®), minocycline (MINOCIN®), oxytetracycline (TERRAMYCIN®), tetracycline (SUMYCIN®, ACHROMYCIN® V, STECLIN®)), drugs against mycobacteria (*e.g.*, clofazimine (LAMPRENE®), dapsone (AVLOSULFON®), capreomycin (CAPASTAT®), cycloserine (SEROMYCIN®), ethambutol (MYAMBUTOL®), ethionamide (TRECATOR®), isoniazid (I.N.H.®), pyrazinamide (ALDINAMIDE®), rifampin (RIFADIN®, RIMACTANE®), rifabutin (MYCOBUTIN®), rifapentine (PRIFTIN®), streptomycin), and others (*e.g.*, arsphenamine (SALVARSAN®), chloramphenicol (CHLOROMYCETIN®), fosfomycin (MONUROL®), fusidic acid (FUCIDIN®), linezolid (ZYVOX®), metronidazole (FLAGYL®), mupirocin (BACTROBAN®), platensimycin, quinupristin/dalfopristin (SYNERGID®), rifaximin (XIFAXAN®), thiamphenicol, tigecycline (TIGACYL®), tinidazole (TINDAMAX®, FASIGYN®)).

Conditions associated with viral infection

[0099] In another instance, provided are methods for treating or preventing a viral infection and/or a disease, disorder, or condition associated with a viral infection, or a symptom thereof, in a subject, by administering aRNAV comprising one or more polynucleotides encoding an anti-viral polypeptide, *e.g.*, an anti-viral polypeptide described herein in combination with an anti-viral agent, *e.g.*, an anti-viral polypeptide or a small molecule anti-viral agent described herein.

[0100] Diseases, disorders, or conditions associated with viral infections which may be treated using the NAVs of the invention include, but are not limited to, acute febrile pharyngitis, pharyngoconjunctival fever, epidemic keratoconjunctivitis, infantile gastroenteritis, Coxsackie infections, infectious mononucleosis, Burkitt lymphoma, acute hepatitis, chronic hepatitis, hepatic cirrhosis, hepatocellular carcinoma, primary HSV-1 infection (*e.g.*, gingivostomatitis in children, tonsillitis and pharyngitis in adults, keratoconjunctivitis), latent HSV-1 infection (*e.g.*, herpes labialis and cold sores), primary HSV-2 infection, latent HSV-2 infection, aseptic meningitis, infectious mononucleosis, Cytomegalic inclusion disease, Kaposi sarcoma, multicentric Castleman disease, primary effusion lymphoma, AIDS, influenza, Reye syndrome, measles, postinfectious encephalomyelitis, Mumps, hyperplastic epithelial lesions (*e.g.*, common, flat, plantar and anogenital warts, laryngeal papillomas, epidermodysplasia verruciformis), cervical carcinoma, squamous cell carcinomas, croup, pneu-

monia, bronchiolitis, common cold, Poliomyelitis, Rabies, bronchiolitis, pneumonia, influenza-like syndrome, severe bronchiolitis with pneumonia, German measles, congenital rubella, Varicella, and herpes zoster.

Viral pathogens

[0101] Examples of viral infectious agents include, but are not limited to, adenovirus; Herpes simplex, type 1; Herpes simplex, type 2; encephalitis virus, papillomavirus, Varicella-zoster virus; Epstein-barr virus; Human cytomegalovirus; Human herpesvirus, type 8; Human papillomavirus; BK virus; JC virus; Smallpox; polio virus, Hepatitis B virus; Human bocavirus; Parvovirus B19; Human astrovirus; Norwalk virus; coxsackievirus; hepatitis A virus; poliovirus; rhinovirus; Severe acute respiratory syndrome virus; Hepatitis C virus; yellow fever virus; dengue virus; West Nile virus; Rubella virus; Hepatitis E virus; Human immunodeficiency virus (HIV); Influenza virus, type A or B; Guanarito virus; Junin virus; Lassa virus; Machupo virus; Sabia virus; Crimean-Congo hemorrhagic fevervirus; Ebola virus; Marburg virus; Measles virus; Mumps virus; Parainfluenza virus; Respiratory syncytial virus; Human metapneumovirus; Hendra virus; Nipah virus; Rabies virus; Hepatitis D; Rotavirus; Orbivirus; Coltivirus; Hantavirus, Middle East Respiratory Coronavirus; Chikungunya virus or Banna virus.

[0102] Viral pathogens may also include viruses that cause resistant viral infections.

Antiviral agents

[0103] Exemplary anti-viral agents include, but are not limited to, abacavir (ZIAGEN®), abacavir/lamivudine/zidovudine (trizivir®), aciclovir or acyclovir (CYCLOVIR®, HERPEX®, ACIVIR®, ACIVIRAX®, ZOVIRAX®, ZOVIR®), adefovir (Preveon®, Hepsera®), amantadine (SYMMETREL®), amprenavir (AGENERASE®), ampligen, arbidol, atazanavir (REYATAZ®), boceprevir, cidofovir, darunavir (PREZISTA®), delavirdine (RESCRIPTOR®), didanosine (VIDEX®), docosanol (ABREVA®), edoxudine, efavirenz (SUSTIVA®, STOCRIN®), emtricitabine (EMTRIVA®), emtricitabine/tenofovir/efavirenz (ATRIPLA®), enfuvirtide (FUZEON®), entecavir (BARACLUDE®, ENNAVIR®), famciclovir (FAMVIR®), fomivirsen (VITRAVENE®), fosamprenavir (LEXIVA®, TELZIR®), foscarnet (FOSCAVIR®), fosfonet, ganciclovir (CYTOVENE®, CYMEVENE®, VITRASERT®), GS 9137 (ELVITEGRAVIR®), imiquimod (ALDARA®, ZYCLARA®, BESELNA®), indinavir (CRIXIVAN®), inosine, inosine pranobex (IMUNOVIR®), interferon type I, interferon type II, interferon type III, kutapressin (NEXAVIR®), lamivudine (ZEFFIX®, HEPTOVIR®, EPIVIR®), lamivudine/zidovudine (COMBI VIR®), lopinavir, loviride, maraviroc (SELZENTRY®, CELSENTRI®), methisazone, MK-2048, moroxydine, nelfinavir (VIRACEPT®), nevirapine (VIRAMUNE®), oseltamivir (TAMIFLU®), peginterferon alfa-2a (PEGASYS®), penciclovir (DENAVIR®), peramivir, pleconaril, podophyllotoxin (CONDYLOX®), raltegravir (ISENTRESS®), ribavirin (COPEGUs®, REBETOL®, RIBASPHERE®, VILONA® AND VIRAZOLE®), rimantadine (FLUMADINE®), ritonavir (NORVIR®), pyramidine, saquinavir (INVIRASE®, FORTOVASE®), stavudine, tea free oil (melaleuca oil), tenofovir (VIREAD®), tenofovir/emtricitabine (TRUVADA®), tipranavir (APTIVUS®), trifluridine (VIROPTIC®), tromantadine (VIRU-MERZ®), valaciclovir (VALTREX®), valganciclovir (VALCYTE®), vicriviroc, vidarabine, viramidine, zalcitabine, zanamivir (RELENZA®), and zidovudine (azidothymidine (AZT), RETROVIR®, RETROVIS®).

B. Therapeutic Settings and/or Situations

[0104] NAVs of the present invention may be utilized in various settings depending on the prevalence of the infection or the degree or level of unmet medical need. Some applications of the NAVs of the disclosure are outlined in Table 1.

Table 1. Infectious Agents by prevalence and Medical Need

	Unmet Need (Infectious Agent Target)		
Prevalence	**Short term benign Long term sequelae**	**Short term morbidity but some treatment**	**Short term mortality; no treatment available**
Ubiquitous	HPV, HCV, UTRIs	Dengue, Chikungunya, ETEC and GI bacteria and S. Pneumo PNA	seasonal and pandemic influenza, MRSA and TB
At risk populations	VZV, Lyme and Chlamydia, N. gonorrhea and HSV	Noroviruses, HEV, CMV, HIV and N. meningitis	Klebsiella, Pseudomonas, Rabies and C. difficile
Rare disease	--	VEV	toxin-mediated diseases, hantavirus, arborviruses such as JE, WNV and EEE

Certain abbreviations include: HPV – Human Papillomavirus; HCV – Hepatitis C Virus; HEV – Human Enterovirus; MERS-CoV: Middle East Respiratory Syndrom Corona Virus; VZV – Varicella-zoster Virus; MRSA – Methicillin-resistant Staph areus; TB—tuberculosis; WNV—West Nile Virus; VEV--vesicular exanthema virus; EEE—Eastern equine encephalitis, JE—Japanese encephalitis, ETEC—Enterotoxigenic E. coli.

Influenza (seasonal and pandemic)

[0105] Symptoms of the flu include dry cough, fever, chills, myalgias progressing to respiratory failure and the risk of secondary bacterial infections (e.g., MRSA). Seasonal influenza is ubiquitous and consists of three principal strains (A [H1N1], A [H3N2], and B), which are covered by the annual vaccine. Pandemic flu occurs because the viruses' unique reassortment ability allowing antigenic shift as well as transfer between avian and swine flu strains. One emerging concern in Southeast Asia is the pandemic potential of several new strains. Such pandemic outbreaks have a high mortality rate with few available treatments. Anti-virals only provide symptomatic relief and must be given in the first 48 hours.

[0106] The NAVs of the present invention have superior properties in that they produce much larger antibody titers, produce responses early than commercially available anti-virals and may be administered after the critical 48 hour period while retaining efficacy.

[0107] While not wishing to be bound by theory, the inventors hypothesize that the NAVs of the invention, as mRNA polynucleotides, are better designed to produce the appropriate protein conformation on translation as the NAVs co-opt natural cellular machinery. Unlike tranditional vaccines which are manufactured ex vivo and may trigger unwanted cellular responses, the NAVs are presented to the cellular system in amore native fashion. Adding to the superior effects may also involve the formulations utilized which may enither serve to shield or traffic the NAVs.

[0108] According to the present invention, NAVs represent a tailored active vaccine that not only can prevent infection but can limit transmission of influenza.

[0109] In some embodiments, the NAVs may be used to prevent pandemic influenza by reacting to emerging new strains with the very rapid NAV-based vaccine production process. In some embodiments, new NAV for treating or prophylactically preventing influenza outbreaks, including for emerging strains (e.g., H7N9 and H10N8), may be produced in less than six weeks, from the time of antigen identification to available vaccine.

[0110] In some instances a single injection of a single antigen encoding NAV polynucleotide may provide protection for an entire flu season.

Influenza: Maintenance of Antigenic memory

[0111] The NAV compositions of the present invention may also be used to maintain or restore antigenic memory in a subject or population as part of a vaccination plan.

[0112] With the speed and versatility of the NAV technology of the present invention, it is now possible to create a vaccination plan that spans both temporal and viral strain space.

[0113] In one embodiment, NAV compositions may be created which include polynucleotides that encode one or more flu year antigens. As used herein a flu year antigen is an antigen which is selected from a strain of influenza used as a component of a flu vaccine from a particularyear. For example, the influenza A strain, A/Port Chalmers/1/1973(H3N2)-like virus, represents one strain component of the Northern Hemisphere vaccine from 1974-1975.

[0114] According to the present disclosure, a vaccination scheme or plan is developed which allows for not only ongoing vaccination in the current year but antigenic memory booster vaccinations across years, strains, or groups thereof to establish and maintain antigenic memory in a population In this manner, a population is less likely to succumb to any pandemic or outbreak involving recurrence of older strains or the appearance of antigens from older strains.

[0115] Any combination of prior vaccine component strains utilized to create or design an antigenic memory booster vaccine is referred to here as a reference set.

[0116] In one instance, NAVs which are antigenic memory booster vaccines are administered to boost antigenic

memory across a time period of 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 35, 40, 45, 50or more years.

[0117] In one instance, NAVs which are antigenic memory booster vaccines are administered to boost antigenic memory for alternating historic years including every other year from the past vaccine component strains relative to a current year. In some instancesthe selection of the vaccine components can be from every 3rd, 4th, 5 h, 6th, 7th, 8 h, 9th, 10th or more years.

[0118] In one instance, NAVs which are antigenic memory booster vaccines are administered to boost antigenic memory over ten year periods.

[0119] In some instancesNAVs which are antigenic memory booster vaccines are administered to boost antigenic memory and are selected from a number of influenza type A strains as a first selection combined with a selection from a number of influenza type B strains or other strains listed herein. The number of selections of type A or type B may be independently, 1, 2, 1, 4, 5, 6, 7, 8, 9, 10 or more.

[0120] In all cases, the antigenic memory booster vaccine strains for antigen encoding in the NAVs may be selected from either the Northern or Southern hemisphere vaccine components independently

[0121] In some instances, the NAV booster vaccine may be used in a population either once or periodically to create herd immunity. Such immunity is present when greater than 30% of a population is protected.

[0122] The components or strains of influenza which may be utilized in the antigenic memory booster vaccines include, but are not limited to, those in Tables 2-5.

Table 2. Influenza vaccine components by year

Northern hemisphere	H1N1	H3N2	B-strain	additional B-strain for QIV
1974-1975	N/A	A/Port Chalmers/1/1973(H3N2)-like virus	B/HongKong/05/1972-like virus	N/A
1975-1976	A/Scotland/840/74-like virus H1N1	A/Port Chalmers/1/1973(H3N2)-like virus	B/HongKong/05/1972-like virus	N/A
1976-1977	N/A	A/Victoria/3/75(H3N2)-like virus	B/HongKong/05/1972-like virus	N/A
1977-1978	N/A	A/Victoria/3/75(H3N2)-like virus	B/HongKong/05/1972-like virus	N/A
1978-1979	A/USSR/90/77(H1N1)-like virus	A/Texas/1/77(H3N2)-like virus	B/HongKong/05/1972-like virus	N/A
1979-1980	A/USSR/90/77(H1N1)-like virus	A/Texas/1/77(H3N2)-like virus	N/A	N/A
1980-1981	A/Brazil/11/78(H1N1)-like virus	A/Bangkok/01/1979(H3N2)-like virus	B/Singapore/222/79-like virus	N/A
1981-1982	A/Brazil/11/78(H1N1)-like virus	A/Bangkok/01/1979(H3N2)-like virus	B/Singapore/222/79-like virus	N/A
1982-1983	A/Brazil/11/78(H1N1)-like virus	A/Bangkok/01/1979(H3N2)-like virus	B/Singapore/222/79-like virus	N/A
1983-1984	A/Brazil/11/78(H1N1)-like virus	A/Philippines/2/82(H3N2)-like virus	B/Singapore/222/79-like virus	N/A

1984-1985	A/Chile/1/83(H1N1)-like virus	A/Philippines/2/82(H3N2)-like virus	B/USSR/100/83-like virus	N/A
1985-1986	A/Chile/1/83(H1N1)-like virus	A/Philippines/2/82(H3N2)-like virus	B/USSR/100/83-like virus	N/A
1986-1987	A/Chile/1/83(H1N1)-like virus	A/Christchurch/4/1985(H3N2)-like virus and A/Mississippi/1/85(H3N2)-like virus	B/Ann Arbor/1/86-like virus	N/A
1987-1988	A/Singapore/6/1986(H1N1)-like virus	A/Leningrad/360/1986(H3N2)-like strain	N/A	N/A
1988-1989	A/Singapore/6/1986(H1N1)-like virus	A/Sichuan/02/87(H3N2)-like virus	B/Beijing/1/87-like rivus	N/A
1989-1990	A/Singapore/6/1986(H1N1)-like virus	A/Shanghai/11/87(H3N2)-like virus	B/Yamagata/16/88-like virus	N/A
1990-1991	A/Singapore/6/1986(H1N1)-like virus	A/Guizhou/54/89(H3N2)-like virus	B/Yamagata/16/88-like virus	N/A
1991-1992	A/Singapore/6/1986(H1N1)-like virus	A/Beijing/353/89(H3N2)-like virus	B/Yamagata/16/88-like virus	N/A
1992-1993	N/A	A/Beijing/353/89(H3N2)-like virus	B/Yamagata/16/88-like virus	N/A
1993-1994	A/Singapore/6/1986(H1N1)-like virus	A/Beijing/32/92(H3N2)-like virus	B/Panama/45/90-like virus	N/A
1994-1995	A/Singapore/6/1986(H1N1)-like virus	A/Shangdong/9/93(H3N2)-like virus	B/Panama/45/90-like virus	N/A
1995-1996	A/Singapore/6/1986(H1N1)-like virus	A/Johannesburg/33/94(H3N2)-like virus	B/Beijing/184/93-like virus	N/A
1996-1997	A/Singapore/6/1986(H1N1)-like virus	A/Wuhan/359/95(H3N2)-like virus	B/Beijing/184/93-like virus	N/A
1997-1998	A/Bayern/7/95(H1N1)-like virus	A/Wuhan/359/95 (H3N2)-like virus	B/Beijing/184/93-like virus	N/A

Table 3. Influenza vaccine components by year-Southern Hemisphere

Southern Hemisphere	H1N1	H3N2	B-strain	additional B-strain for QIV
1975	N/A	A/Port Chalmers/1/1973(H3N2)-like virus	B/HongKong/05/1972-like virus	N/A
1976	A/Scotland/840/74-like virus (H1N1)	A/Port Chalmers/1/1973(H3N2)-like virus	B/HongKong/05/1972-like virus	N/A
1977	N/A	A/Victoria/3/75(H3N2)-like virus	B/HongKong/05/1972-like virus	N/A
1978	N/A	A/Victoria/3/75(H3N2)-like virus	B/HongKong/05/1972-like virus	N/A
1979	A/USSR/90/77(H1N1)-like virus	A/Texas/1/77(H3N2)-like virus	B/HongKong/05/1972-like virus	N/A
1980	A/USSR/90/77(H1N1)-like virus	A/Texas/1/77(H3N2)-like virus	N/A	N/A
1981	A/Brazil/11/78(H1N1)-like virus	A/Bangkok/01/1979(H3N2) -like virus	B/Singapore/222/79-like virus	N/A
1982	A/Brazil/11/78(H1N1)-like virus	A/Bangkok/01/1979(H3N2) -like virus	B/Singapore/222/79-like virus	N/A
1983	A/Brazil/11/78(H1N1)-like virus	A/Bangkok/01/1979(H3N2) -like virus	B/Singapore/222/79-like virus	N/A
1984	A/Brazil/11/78(H1N1)-like	A/Philippines/2/82(H3N2)-	B/Singapore/222/79-like	N/A

	virus	like virus	virus	
1985	A/Chile/1/83(H1N1)-like virus	A/Philippines/2/82(H3N2)-like virus	B/USSR/100/83-like virus	N/A
1986	A/Chile/1/83(H1N1)-like virus	A/Philippines/2/82(H3N2)-like virus	B/USSR/100/83-like virus	N/A
1987	A/Chile/1/83(H1N1)-like virus	A/Christchurch/4/1985(H3N2)-like virus and A/Mississippi/1/85(H3N2)-like virus	B/Ann Arbor/1/86-like virus	N/A
1988	A/Singapore/6/1986(H1N1)-like virus	A/Leningrad/360/1986(H3N2)-like virus	N/A	N/A
1989	A/Singapore/6/1986(H1N1)-like virus	A/Sichuan/02/87(H3N2)-like virus	B/Beijing/1/87-like virus	N/A
1990	A/Singapore/6/1986(H1N1)-like virus	A/Shanghai/11/87(H3N2)-like virus	B/Yamagata/16/88-like virus	N/A
1991	A/Singapore/6/1986(H1N1)-like virus	A/Guizhou/54/89(H3N2)-like virus	B/Yamagata/16/88-like virus	N/A
1992	A/Singapore/6/1986(H1N1)-like virus	A/Beijing/353/89(H3N2)-like virus	B/Yamagata/16/88-like virus	N/A
1993	A/Singapore/6/1986(H1N1)-like virus	A/Beijing/353/89(H3N2)-like virus	B/Yamagata/16/88-like virus	N/A
1994	A/Singapore/6/1986(H1N1)-like virus	A/Beijing/32/92(H3N2)-like virus	B/Panama/45/90-like virus	N/A
1995	A/Singapore/6/1986(H1N1)-like virus	A/Shangdong/9/93(H3N2)-like virus	B/Panama/45/90-like virus	N/A
1996	A/Singapore/6/1986(H1N1)-like virus	A/Johannesburg/33/94(H3N2)-like virus	B/Beijing/184/93-like virus	N/A
1997	A/Singapore/6/1986(H1N1)-like virus	A/Wuhan/359/95(H3N2)-like virus	B/Beijing/184/93-like strain	N/A
1998	A/Bayern/7/95(H1N1)-like virus	A/Wuhan/359/95(H3N2)-like virus	B/Beijing/184/93-like virus	N/A
1999	A/Beijing/262/95(H1N1)-like virus	A/Sydney/5/97(H3N2)-like virus	B/Beijing/184/93-like virus	N/A

Table 4. Influenza Vaccine components by year-Northern Hemisphere

Northern hemisphere	H1N1	H3N2	B-strain	additional B-strain for QIV
November 1998–April 1999	A/Beijing/262/95(H1N1)-like virus	A/Sydney/5/97(H3N2)-like virus	B/Beijing/184/93-like virus	N/A
November 1999–April 2000	A/Beijing/262/95 (H1N1)-like virus	A/Sydney/5/97 (H3N2)-like virus	B/Beijing/184/93-like virus or B/Shangdong/7/97-like virus	N/A
2000–2001	A/New Caledonia/20/99 (H1N1)-like virus	A/Moscow/10/99 (H3N2)-like virus	B/Beijing/184/93-like virus	N/A
2001–2002	A/New Caledonia/20/99(H1N1)-like virus	A/Moscow/10/99(H3N2)-like virus	B/Sichuan/379/99-like virus	N/A
2002–2003	A/New Caledonia/20/99(H1N1)-like virus	A/Moscow/10/99(H3N2)-like virus	B/Hong Kong/330/2001-like virus	N/A
2003–2004	A/New Caledonia/20/99(H1N1)-like virus	A/Moscow/10/99(H3N2)-like virus	B/Hong Kong/330/2001-like virus	N/A
2004-2005	A/New Caledonia/20/99(H1N1)-like virus	A/Fujian/411/2002(H3N2)-like virus	B/Shanghai/361/2002-like virus	N/A
2005-2006	A/New Caledonia/20/99(H1N1)-like virus	A/California/7/2004(H3N2)-like virus	B/Shanghai/361/2002-like virus	N/A
2006-2007	A/New Caledonia/20/99(H1N1)-like	A/Wisconsin/67/2005 (H3N2)-like virus	B/Malaysia/2506/2004-like virus	N/A
2007-2008	A/Solomon Islands/3/2006 (H1N1)-like virus	A/Wisconsin/67/2005 (H3N2)-like virus	B/Malaysia/2506/2004-like virus	N/A
2008-2009	A/Brisbane/59/2007 (H1N1)-like virus	A/Brisbane/10/2007 (H3N2)-like virus	B/Florida/4/2006-like virus	N/A
2009-2010	A/Brisbane/59/2007 (H1N1)-like virus	A/Brisbane/10/2007 (H3N2)-like virus	B/Brisbane/60/2008-like virus	N/A
2010-2011	A/California/7/2009 (H1N1)-like virus	A/Perth/16/2009 (H3N2)-like virus	B/Brisbane/60/2008-like virus	N/A
2011-2012	A/California/7/2009 (H1N1)-like virus	A/Perth/16/2009 (H3N2)-like virus	B/Brisbane/60/2008-like virus	N/A
2012-2013	A/California/7/2009 (H1N1)pdm09-like virus	A/Victoria/361/2011 (H3N2)-like virus	B/Wisconsin/1/2010-like virus	B/Brisbane/60/2008-like virus
2013-2014	A/California/7/2009 (H1N1)pdm09-like virus	A(H3N2) virus antigenically like the cell-propagated prototype virus A/Victoria/361/2011	B/Massachusetts/2/2012-like virus	B/Brisbane/60/2008-like virus
2014-2015	A/California/7/2009 (H1N1)pdm09-like virus	A/Texas/50/2012 (H3N2)-like virus	B/Massachusetts/2/2012-like virus	B/Brisbane/60/2008-like virus

Table 5. Influenza Vaccine components by year-Southern Hemisphere

Southern hemisphere	H1N1	H3N2	B-strain	additional B-strain for QIV
1999	A/Beijing/262/95(H1N1)-like virus	A/Sydney/5/97(H3N2)-like virus	B/Beijing/184/93-like virus	N/A
May–October 2000	A/New Caledonia/20/99 (H1N1)-like virus	A/Moscow/10/99 (H3N2)-like virus	B/Beijing/184/93-like virus or B/Shangdong/7/97-like virus	N/A
May–October 2001	A/New Caledonia/20/99 (H1N1)-like virus	A/Moscow/10/99 (H3N2)-like virus	B/Sichuan/379/99-like virus	N/A
2002	A/New Caledonia/20/99(H1N1)-like virus	A/Moscow/10/99(H3N2)-like virus	B/Sichuan/379/99-like virus	N/A
2003	A/New Caledonia/20/99(H1N1)-like virus	A/Moscow/10/99(H3N2)-like virus	B/Hong Kong/330/2001-like virus	N/A
2004	A/New Caledonia/20/99(H1N1)-like virus	A/Fujian/411/2002(H3N2) -like virus	B/Hong Kong/330/2001-like virus	N/A
2005	A/New Caledonia/20/99(H1N1)-like virus	A/Wellington/1/2004(H3N2)-like virus	B/Shanghai/361/2002-like virus	N/A
2006	A/New Caledonia/20/99(H1N1)-like virus	A/California/7/2004(H3N2)-like virus	B/Malaysia/2506/2004-like virus	N/A
2007	A/New Caledonia/20/99(H1N1)-like	A/Wisconsin/67/2005 (H3N2)-like virus	B/Malaysia/2506/2004-like virus	N/A
2008	A/Solomon Islands/3/2006 (H1N1)-like virus	A/Brisbane/10/2007 (H3N2)-like virus	B/Florida/4/2006-like virus	N/A
2009	A/Brisbane/59/2007 (H1N1)-like virus	A/Brisbane/10/2007 (H3N2)-like virus	B/Florida/4/2006-like virus	N/A
2010	A/California/7/2009 (H1N1)-like virus	A/Perth/16/2009 (H3N2)-like virus	B/Brisbane/60/2008-like virus	N/A
2011	A/California/7/2009 (H1N1)-like virus	A/Perth/16/2009 (H3N2)-like virus	B/Brisbane/60/2008-like virus	N/A
2013	A/California/7/2009 (H1N1)pdm09 like virus	A/Perth/16/2009 (H3N2)-like virus	B/Brisbane/60/2008-like virus	N/A
2013	A/California/7/2009 (H1N1)pdm09-like virus	A/Victoria/361/2011 (H3N2)-like virus	B/Wisconsin/1/2010-like virus	B/Brisbane/60/2008-like virus
2014	A/California/7/2009 (H1N1)pdm09-like virus	A/Texas/50/2012 (H3N2)-like virus	B/Massachusetts/2/2012-like virus	B/Brisbane/60/2008-like virus

Influenza antigens

[0123] In some embodiments, the NAV polnynucleotides may encode one or more polypeptides of an influenza strain as an antigen. Such antigens include, but are not limited to those antigens encoded by the polynucleotides listed in Tables 6-18. In the table, the GenBank Accession Number represents either the complete or partial CDS of the encoded antigen. The NAV polynucleotides may comprise a region of any of the sequences listed in the tables or entire coding region of the mRNA listed They may comprise hybrid or chimeric regions, or mimics orvariants.

[0124] Any of the strains referred to in Tables 6-14 may also be used in an antigenic memory booster vaccine as described herein.

Table 6. Influenza H1N1 Antigens

Strain/Protein	Length	GenBank / GI Accession No.
Influenza A virus (A/Bayern/7/95(H1N1)) NA gene for neuraminidase, genomic RNA	1,459 bp linear mRNA	AJ518104.1 GI:31096418
Influenza A virus (A/Brazil/11/1978(X-71)(H1N1)) mRNA for hemagglutinin HA1, escape variant 1	1,072 bp linear mRNA	X86654.1 GI:995549
Influenza A virus (A/Brazil/11/1978(X-71)(H1N1)) mRNA for hemagglutinin HA1, escape variant 2	1,072 bp linear mRNA	X86655.1 GI:995550
Influenza A virus (A/Brazil/11/1978(X-71)(H1N1)) mRNA for hemagglutinin HA1, escape variant 3	1,072 bp linear mRNA	X86656.1 GI:995551
Influenza A virus (A/Brazil/11/1978(X-71)(H1N1)) mRNA for hemagglutinin HA1, escape variant 4	1,072 bp linear mRNA	X86657.1 GI:995552
Influenza A virus (A/Brevig_Mission/1/18(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,220 bp linear mRNA	AF116575.1 GI:4325017
Influenza A virus (A/Brevig_Mission/1/18(H1N1)) neuraminidase (NA) gene, complete cds	1,410 bp linear mRNA	AF250356.2 GI:13260556
Influenza A virus (A/Brevig Mission/1/1918(H1N1)) nucleoprotein (np) mRNA, complete cds	1,497 bp linear mRNA	AY744935.1 GI:55273940
Influenza A virus (A/Brevig Mission/1/1918(H1N1)) polymerase PB2 (PB2) mRNA, complete cds	2,280 bp linear mRNA	DQ208309.1 GI:76786704

Influenza A virus (A/Brevig Mission/1/1918(H1N1)) polymerase PB1 (PB1) mRNA, complete cds	2,274 bp linear mRNA	DQ208310.1 GI:76786706
Influenza A virus (A/Brevig Mission/1/1918(H1N1)) polymerase PA (PA) mRNA, complete cds	2,151 bp linear mRNA	DQ208311.1 GI:76786708
Influenza A virus (A/camel/Mongolia/1982(H1N1)) hemagglutinin mRNA, partial cds	366 bp linear mRNA	M73975.1 GI:324242
Influenza A virus (A/camel/Mongolia/1982(H1N1)) matrix protein mRNA, partial cds	460 bp linear mRNA	M73978.1 GI:324402
Influenza A virus (A/camel/Mongolia/1982(H1N1)) neuraminidase (NA) mRNA, partial cds	310 bp linear mRNA	M73976.1 GI:324579
Influenza A Virus A/camel/Mongolia/82 NS1 protein mRNA, partial cds	273 bp linear mRNA	M73977.1 GI:324768
Influenza A virus (A/camel/Mongolia/1982(H1N1)) PA polymerase mRNA, partial cds	227 bp linear mRNA	M73974.1 GI:324931
Influenza A virus (A/camel/Mongolia/1982(H1N1)) PB1 protein mRNA, partial cds	531 bp linear mRNA	M73973.1 GI:324971
Influenza A Virus (A/camel/Mongolia/82(H1N1)) polymerase 2 (P2) mRNA, partial cds	379 bp linear mRNA	M73972.1 GI:324993
Influenza A virus (A/chicken/Hong Kong/14/1976(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	1,169 bp linear mRNA	U46782.1 GI:1912328
Influenza A virus (A/Chonnam/07/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,452 bp linear mRNA	AY297141.1 GI:31871990
Influenza A virus (A/Chonnam/07/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,137 bp linear mRNA	AY297154.1 GI:32140347
Influenza A virus (A/Chonnam/18/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	AY297143.1 GI:31871994
Influenza A virus (A/Chonnam/18/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AY297156.1 GI:32140355
Influenza A virus (A/Chonnam/19/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	AY310410.1 GI:31872389
Influenza A virus (A/Chonnam/19/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,167 bp linear mRNA	AY299502.1 GI:32140392
Influenza A virus (A/Chonnam/51/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,443 bp linear mRNA	AY310412.1 GI:31873090
Influenza A virus (A/Chonnam/51/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,161 bp linear mRNA	AY299498.1 GI:32140384
Influenza A virus (A/Chungbuk/50/2002(H1N1)) neuraminidase (NA) mRNA, partial cds	1,425 bp linear mRNA	AY297150.1 GI:31872010

Influenza A virus (A/Chungbuk/50/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,161 bp linear mRNA	AY299506.1 GI:32140400
Influenza A virus (A/Denmark/40/2000(H1N1)) NA gene for neuraminidase, genomic RNA	1,458 bp linear mRNA	AJ518095.1 GI:31096400
Influenza A virus (A/Denver/1/57(H1N1)) neuraminidase mRNA, partial cds	379 bp linear mRNA	AF305216.1 GI:10732818
Influenza A virus (A/Denver/1/57(H1N1)) matrix protein gene, partial cds	442 bp linear mRNA	AF305217.1 GI:10732820
Influenza A virus (A/Denver/1/57(H1N1)) hemagglutinin gene, partial cds	215 bp linear mRNA	AF305218.1 GI:10732822
Influenza A virus (A/duck/Australia/749/80(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	981 bp linear mRNA	U47309.1 GI:1912348
Influenza A virus (A/duck/Australia/749/80(H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,777 bp linear mRNA	AF091312.1 GI:4585166
Influenza A virus (A/duck/Bavaria/1/77 (H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,777 bp linear mRNA	AF091313.1 GI:4585168
Influenza A virus (A/duck/Bavaria/2/77(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	981 bp linear mRNA	U47308.1 GI:1912346
Influenza A virus (A/duck/Eastern China/103/2003(H1N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	EU429749.1 GI:167859463
Influenza A virus (A/duck/Eastern China/152/2003(H1N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,461 bp linear mRNA	EU429751.1 GI:167859467
Influenza A virus (A/Duck/Ohio/118C/93 (H1N1)) neuraminidase (NA) gene, complete cds	1,410 bp linear mRNA	AF250361.2 GI:13260576
Influenza A virus (A/Duck/Ohio/175/86 (H1N1)) neuraminidase (NA) gene, complete cds	1,410 bp linear mRNA	AF250358.2 GI:13260565
Influenza A virus (A/Duck/Ohio/194/86 (H1N1)) neuraminidase (NA) gene, complete cds	1,410 bp linear mRNA	AF250360.2 GI:13260573
Influenza A virus (A/Duck/Ohio/30/86 (H1N1)) neuraminidase (NA) gene, complete cds	1,410 bp linear mRNA	AF250359.2 GI:13260570
Influenza A virus strain A/Fiji/15899/83(H1N1) mRNA for neuraminidase	1,460 bp linear mRNA	AJ006954.1 GI:4210707
Influenza A Virus (A/Fiji/15899/83(H1N1)) mRNA for PB2 protein	2,341 bp linear mRNA	AJ564805.1 GI:31442134
Influenza A Virus (A/Fiji/15899/83(H1N1)) partial mRNA for PB1 protein	2,113 bp linear mRNA	AJ564807.1 GI:31442138
Influenza A virus (A/FM/1/47 (H1N1)) neuraminidase (NA) gene, complete cds	1,395 bp linear mRNA	AF250357.2 GI:13260561

Influenza A virus (A/goose/Hong Kong/8/1976(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	1,091 bp linear mRNA	U46021.1 GI:1912326
Influenza A virus (A/goose/Hong Kong/8/1976(H1N1)) polymerase (PB1) mRNA, partial cds	261 bp linear mRNA	U48284.1 GI:1912372
Influenza A virus (A/goose/Hong Kong/8/1976(H1N1)) nucleoprotein (NP) mRNA, partial cds	1,395 bp linear mRNA	U49093.1 GI:1912384
Influenza A virus (A/Guangzhou/1561/2006(H1N1)) segment 4 hemagglutinin (HA) mRNA, complete cds	1,775 bp linear mRNA	EU382986.1 GI:170762603
Influenza A virus (A/Guangzhou/1561/2006(H1N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,462 bp linear mRNA	EU382993.1 GI:170762617
Influenza A virus (A/Guangzhou/1684/2006(H1N1)) segment 4 hemagglutinin (HA) mRNA, complete cds	1,775 bp linear mRNA	EU382987.1 GI:170762605
Influenza A virus (A/Guangzhou/1684/2006(H1N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,462 bp linear mRNA	EU382994.1 GI:170762619
Influenza A virus (A/Guangzhou/483/2006(H1N1)) segment 4 hemagglutinin (HA) mRNA, complete cds	1,775 bp linear mRNA	EU382981.1 GI:170762593
Influenza A virus (A/Guangzhou/483/2006(H1N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,462 bp linear mRNA	EU382988.1 GI:170762607
Influenza A virus (A/Guangzhou/506/2006(H1N1)) segment 4 hemagglutinin (HA) mRNA, complete cds	1,775 bp linear mRNA	EU382982.1 GI:170762595
Influenza A virus (A/Guangzhou/506/2006(H1N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,461 bp linear mRNA	EU382989.1 GI:170762609
Influenza A virus (A/Guangzhou/555/2006(H1N1)) segment 4 hemagglutinin (HA) mRNA, complete cds	1,775 bp linear mRNA	EU382983.1 GI:170762597
Influenza A virus (A/Guangzhou/555/2006(H1N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,462 bp linear mRNA	EU382990.1 GI:170762611
Influenza A virus (A/Guangzhou/657/2006(H1N1)) segment 4 hemagglutinin (HA) mRNA, complete cds	1,775 bp linear mRNA	EU382984.1 GI:170762599
Influenza A virus (A/Guangzhou/657/2006(H1N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,462 bp linear mRNA	EU382991.1 GI:170762613
Influenza A virus (A/Guangzhou/665/2006(H1N1)) segment 4 hemagglutinin (HA) mRNA, complete cds	1,775 bp linear mRNA	EU382985.1 GI:170762601
Influenza A virus (A/Guangzhou/665/2006(H1N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,462 bp linear mRNA	EU382992.1 GI:170762615
Influenza A virus (A/Gwangju/55/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,431 bp linear mRNA	AY297151.1 GI:31872012
Influenza A virus (A/Gwangju/55/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,179 bp linear mRNA	AY299507.1 GI:32140402

Influenza A virus (A/Gwangju/57/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,446 bp linear mRNA	AY297152.1 GI:31872014
Influenza A virus (A/Gwangju/57/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,167 bp linear mRNA	AY299508.1 GI:32140404
Influenza A virus (A/Gwangju/58/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,434 bp linear mRNA	AY297153.1 GI:31872016
Influenza A virus (A/Gwangju/58/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AY299509.1 GI:32140406
Influenza A virus (A/Gwangju/90/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,446 bp linear mRNA	AY297147.1 GI:31872002
Influenza A virus (A/Gwangju/90/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,164 bp linear mRNA	AY299499.1 GI:32140386
Influenza A virus (A/Hong Kong/437/2002(H1N1)) partial NA gene for neuraminidase, genomic RNA	1,403 bp linear mRNA	AJ518101.1 GI:31096412
Influenza A virus (A/Hong Kong/747/2001(H1N1)) partial NA gene for neuraminidase, genomic RNA	1,352 bp linear mRNA	AJ518102.1 GI:31096414
Influenza A virus (A/London/1/1918(H1N1)) hemagglutinin (HA) mRNA, partial cds	563 bp linear mRNA	AY184805.1 GI:32395285
Influenza A virus (A/London/1/1919(H1N1)) hemagglutinin (HA) mRNA, partial cds	563 bp linear mRNA	AY184806.1 GI:32395287
Influenza A virus (A/Loygang/4/1957(H1N1)) nucleoprotein mRNA, complete cds	1,565 bp linear mRNA	M76604.1 GI:324255
Influenza A virus (A/Lyon/651/2001(H1N1)) partial NA gene for neuraminidase, genomic RNA	1,318 bp linear mRNA	AJ518103.1 GI:31096416
Influenza A virus (A/mallard/Alberta/119/98 (H1N1)) nonfunctional matrix protein mRNA, partial sequence	947 bp linear mRNA	AY664487.1 GI:51011891
Influenza A virus (A/duck/Alberta/35/76(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	981 bp linear mRNA	U47310.1 GI:1912350
Influenza A virus (A/duck/Alberta/35/76(H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,777 bp linear mRNA	AF091309.1 GI:4585160
Influenza A virus (A/duck/Alberta/35/76(H1N1)) neuraminidase (NA) gene, complete cds	1,410 bp linear mRNA	AF250362.2 GI:13260579
Influenza A virus (A/mallard/Tennessee/11464/85 (H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	981 bp linear mRNA	U47307.1 GI:1912344
Influenza A virus (A/mallard/Tennessee/11464/85 (H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,777 bp linear mRNA	AF091311.1 GI:4585164
Influenza A virus (A/New Caledonia/20/1999(H1N1)) segment 7 matrix protein 2 (M2) mRNA, complete cds	294 bp linear mRNA	HQ008884.1 GI:302566794

Influenza A virus (A/New Jersey/4/1976(H1N1)) nucleoprotein mRNA, complete cds	1,565 bp linear mRNA	M76605.1 GI:324581
Influenza A virus (A/New Jersey/8/1976(H1N1)) nucleoprotein mRNA, complete cds	1,565 bp linear mRNA	M76606.1 GI:324583
Influenza A virus (A/New_York/1/18(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,220 bp linear mRNA	AF116576.1 GI:4325019
Influenza A virus (A/Ohio/3523/1988(H1N1)) nucleoprotein mRNA, complete cds	1,565 bp linear mRNA	M76602.1 GI:324889
Influenza A virus (A/Pusan/22/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,455 bp linear mRNA	AY310411.1 GI:31872391
Influenza A virus (A/Pusan/22/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,149 bp linear mRNA	AY299503.1 GI:32140394
Influenza A virus (A/Pusan/23/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,440 bp linear mRNA	AY297144.1 GI:31871996
Influenza A virus (A/Pusan/23/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,158 bp linear mRNA	AY297157.1 GI:32140357
Influenza A virus (A/Pusan/24/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,449 bp linear mRNA	AY297145.1 GI:31871998
Influenza A virus (A/Pusan/24/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,128 bp linear mRNA	AY299494.1 GI:32140376
Influenza A virus (A/Pusan/44/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,431 bp linear mRNA	AY297148.1 GI:31872004
Influenza A virus (A/Pusan/44/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,167 bp linear mRNA	AY299504.1 GI:32140396
Influenza A virus (A/Pusan/45/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,434 bp linear mRNA	AY297146.1 GI:31872000
Influenza A virus (A/Pusan/45/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,167 bp linear mRNA	AY299496.1 GI:32140380
Influenza A virus (A/Pusan/46/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,422 bp linear mRNA	AY310408.1 GI:31872385
Influenza A virus (A/Pusan/46/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AY299497.1 GI:32140382
Influenza A virus (A/Pusan/47/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,437 bp linear mRNA	AY297149.1 GI:31872008
Influenza A virus (A/Pusan/47/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,170 bp linear mRNA	AY299505.1 GI:32140398
Influenza A virus (A/Saudi Arabia/7971/2000(H1N1)) partial NS1 gene for non structural protein 1 and partial NS2 gene for non structural protein 2, genomic RNA	789 bp linear mRNA	AJ519463.1 GI:31096450

Influenza A virus (A/Seoul/11/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,452 bp linear mRNA	AY297142.1 GI:31871992
Influenza A virus (A/Seoul/11/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AY297155.1 GI:32140349
Influenza A virus (A/Seoul/13/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,452 bp linear mRNA	AY310409.1 GI:31872387
Influenza A virus (A/Seoul/13/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,167 bp linear mRNA	AY299500.1 GI:32140388
Influenza A virus (A/Seoul/15/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,449 bp linear mRNA	AY297140.1 GI:31871988
Influenza A virus (A/Seoul/15/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,149 bp linear mRNA	AY299501.1 GI:32140390
Influenza A virus (A/Seoul/33/2002(H1N1)) neuraminidase (NA) mRNA, complete cds	1,437 bp linear mRNA	AY310407.1 GI:31872383
Influenza A virus (A/Seoul/33/2002(H1N1)) hemagglutinin (HA) mRNA, partial cds	1,167 bp linear mRNA	AY299495.1 GI:32140378
Influenza A virus (A/swine/Arnsberg/6554/1979(H1N1)) mRNA for hemagglutinin HA1	1,050 bp linear mRNA	Z46437.1 GI:565609
Influenza A virus (A/swine/Beijing/47/1991(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	1,595 bp linear mRNA	U46783.1 GI:1912330
Influenza A virus (A/swine/Beijing/94/1991(H1N1)) nucleoprotein (NP) mRNA, complete cds	1,565 bp linear mRNA	U49091.1 GI:1912380
Influenza A virus (A/swine/Belgium/1/83(H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,778 bp linear mRNA	AF091316.1 GI:4585174
Influenza A virus (A/swine/Cotes d'Armor/0118/2006(H1N1)) partial mRNA for haemagglutinin precursor (HA1 gene)	1,116 bp linear mRNA	AM490219.1 GI:222062898
Influenza A virus (A/swine/Cotes d'Armor/0136_18/2006(H1N1)) partial mRNA for haemagglutinin precursor (HA1 gene)	1,043 bp linear mRNA	AM490223.1 GI:222062906
Influenza A virus (A/swine/Cotes d'Armor/0184/2006(H1N1)) partial mRNA for haemagglutinin precursor (HA1 gene)	1,089 bp linear mRNA	AM490220.1 GI:222062900
Influenza A virus (A/swine/Cotes d'Armor/0227/2005(H1N1)) partial mRNA for haemagglutinin precursor (HA1 gene)	1,068 bp linear mRNA	AM490221.1 GI:222062902
Influenza A virus (A/swine/Cotes d'Armor/0250/2006(H1N1)) partial mRNA for haemagglutinin precursor (HA1 gene)	1,024 bp linear mRNA	AM490222.1 GI:222062904
Influenza A virus (A/swine/Cotes d'Armor/736/2001(H1N1)) partial HA gene for Haemagglutinin, genomic RNA	1,011 bp linear mRNA	AJ517820.1 GI:38422533
Influenza A virus (A/Swine/England/195852/92 (H1N1)) neuraminidase (NA) gene, complete cds	1,410 bp linear mRNA	AF250366.2 GI:13260593

Influenza A virus PB2 gene for Polymerase 2 protein, genomic RNA, strain A/Swine/Finistere/2899/82	2,268 bp linear mRNA	AJ311457.1 GI:13661037
Influenza A virus PB1 gene for Polymerase 1 protein, genomic RNA, strain A/Swine/Finistere/2899/82	2,341 bp linear mRNA	AJ311462.1 GI:13661047
Influenza A virus PA gene for Polymerase A protein, genomic RNA, strain A/Swine/Finistere/2899/82	2,233 bp linear mRNA	AJ311463.1 GI:13661049
Influenza A virus (A/swine/Finistere/2899/82(H1N1) M1 gene for matrix protein 1 and M2 gene for matrix protein 2, genomic RNA	1,002 bp linear mRNA	AJ316059.1 GI:20068128
Influenza A virus (A/swine/Finistere/2899/82(H1N1)) NS1 gene for non structural protein 1 and NS2 gene for non structural protein 2, genomic RNA	864 bp linear mRNA	AJ344037.1 GI:20068185
Influenza A virus (A/swine/Germany/2/1981(H1N1)) mRNA for PA polymerase	838 bp linear mRNA	X75786.1 GI:438106
Influenza A virus (A/swine/Germany/2/1981(H1N1)) mRNA for neuraminidase (partial)	305 bp linear mRNA	Z30277.1 GI:530399
Influenza A virus (A/swine/Germany/2/1981(H1N1)) mRNA for hemagglutinin	1,730 bp linear mRNA	Z30276.1 GI:563490
165. Influenza A virus (A/swine/Germany/8533/1991(H1N1)) mRNA for hemagglutinin precursor	1,730 bp linear mRNA	Z46434.1 GI:565611
Influenza A virus (A/swine/Guangdong/711/2001(H1N1)) nonfunctional hemagglutinin (HA) mRNA, partial sequence	1,690 bp linear mRNA	AY852271.1 GI:60327789
Influenza A virus (A/swine/Haseluenne/IDT2617/03(H1N1)) hemagglutinin mRNA, complete cds	1,809 bp linear mRNA	EU163946.1 GI:157679548
Influenza A virus (A/swine/Hokkaido/2/81 (H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	981 bp linear mRNA	U47306.1 GI:1912342
Influenza A virus (A/swine/Hokkaido/2/81 (H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,778 bp linear mRNA	AF091306.1 GI:4585154
Influenza A virus (A/swine/Hong Kong/168/1993(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	1,113 bp linear mRNA	U44482.1 GI:1912318
Influenza A virus (A/swine/Hong Kong/168/1993(H1N1)) neuraminidase (NA) mRNA, partial cds	416 bp linear mRNA	U47817.1 GI:1912354
Influenza A virus (A/swine/Hong Kong/168/1993(H1N1)) polymerase (PB2) mRNA, partial cds	286 bp linear mRNA	U48286.1 GI:1912358
Influenza A virus (A/swine/Hong Kong/168/1993(H1N1)) polymerase (PB1) mRNA, partial cds	379 bp linear mRNA	U48283.1 GI:1912370
Influenza A virus (A/swine/Hong Kong/168/1993(H1N1)) polymerase (PA) mRNA, partial cds	308 bp linear mRNA	U48850.1 GI:1912376
Influenza A virus (A/swine/Hong Kong/168/1993(H1N1)) nucleoprotein (NP) mRNA, partial cds	1,397 bp linear mRNA	U49096.1 GI:1912390

Influenza A virus (A/swine/Hong Kong/172/1993(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	1,315 bp linear mRNA	U46020.1 GI:1912324
Influenza A virus (A/swine/Hong Kong/176/1993(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	1,113 bp linear mRNA	U45451.1 GI:1912320
Influenza A virus (A/swine/Hong Kong/273/1994(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	1,330 bp linear mRNA	U45452.1 GI:1912322
Influenza A virus (A/swine/Hong Kong/273/1994(H1N1)) neuraminidase (NA) mRNA, partial cds	241 bp linear mRNA	U47818.1 GI:1912356
Influenza A virus (A/swine/Hong Kong/273/1994(H1N1)) polymerase (PB2) mRNA, partial cds	328 bp linear mRNA	U48287.1 GI:1912360
Influenza A virus (A/swine/Hong Kong/273/1994(H1N1)) polymerase (PB1) mRNA, partial cds	240 bp linear mRNA	U48282.1 GI:1912368
Influenza A virus (A/swine/Hong Kong/273/1994(H1N1)) polymerase (PA) mRNA, partial cds	336 bp linear mRNA	U48851.1 GI:1912378
Influenza A virus (A/swine/Hong Kong/273/1994(H1N1)) nucleoprotein (NP) mRNA, partial cds	1,422 bp linear mRNA	U49092.1 GI:1912382
Influenza A virus (A/swine/IDT/Re230/92hp(H1N1)) hemagglutinin mRNA, complete cds	1,761 bp linear mRNA	EU163947.1 GI:157679550
Influenza A virus (A/swine/IN/1726/1988(H1N1)) nucleoprotein (segment 5) mRNA, complete cds	1,550 bp linear mRNA	L46849.1 GI:954755
Influenza A virus (A/swine/Iowa/15/30(H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	981 bp linear mRNA	U47305.1 GI:1912340
Influenza A virus (A/swine/Iowa/15/30 (H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,778 bp linear mRNA	AF091308.1 GI:4585158
Influenza A virus (A/Swine/Iowa/30 (H1N1)) neuraminidase (NA) gene, complete cds	1,410 bp linear mRNA	AF250364.2 GI:13260586
Influenza A virus (A/swine/Iowa/17672/88 (H1N1)) hemagglutinin precursor (HA) mRNA, partial cds	981 bp linear mRNA	U47304.1 GI:1912338
Influenza A virus (A/swine/Italy/3364/00(H1N1)) partial NS1 gene for non structural protein 1 and partial NS2 gene for non structural protein 2, genomic RNA	864 bp linear mRNA	AJ519462.1 GI:31096447
Influenza A virus (A/swine/Italy-Virus/671/87(H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,777 bp linear mRNA	AF091315.1 GI:4585172
Influenza A Virus (A/swine/Italy/v.147/1981(H1N1)) mRNA for hemagglutinin HA1	1,028 bp linear mRNA	Z46436.1 GI:854214
Influenza A virus (A/swine/Morbihan/0070/2005(H1N1)) partial mRNA for haemagglutinin precursor (HA1 gene)	1,118 bp linear mRNA	AM490218.1 GI:222062896
Influenza A virus (A/swine/Nebraska/1/92(H1N1)) HA protein mRNA, complete cds	1,770 bp linear mRNA	L09063.1 GI:290722

Influenza A virus (A/swine/Nebraska/1/1992(H1N1)) segment 5 nucleoprotein (NP) mRNA, complete cds	1,550 bp linear mRNA	L11164.1 GI:290724
Influenza A virus (A/swine/Netherlands/12/1985(H1N1)) hemagglutinin (HA) mRNA, partial cds	981 bp linear mRNA	U46943.1 GI:1912336
Influenza A virus (A/swine/Netherlands/12/85(H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,776 bp linear mRNA	AF091317.1 GI:4585176
Influenza A virus (A/swine/Netherlands/25/1980(H1N1)) mRNA for nucleoprotein	539 bp linear mRNA	X75791.1 GI:438105
Influenza A virus (A/swine/Netherlands/3/1980(H1N1)) hemagglutinin (HA) mRNA, partial cds	981 bp linear mRNA	U46942.1 GI:1912334
Influenza A virus (A/swine/Netherlands/3/80(H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,778 bp linear mRNA	AF091314.1 GI:4585170
Influenza A virus (A/NJ/11/76 (H1N1)) neuraminidase (NA) gene, complete cds	1,410 bp linear mRNA	AF250363.2 GI:13260583
Influenza A virus (A/Swine/Quebec/192/81 (SwQc81)) neuraminidase mRNA, complete cds	1,438 bp linear mRNA	U86144.1 GI:4099318
Influenza A virus (A/Swine/Quebec/5393/91 (SwQc91)) neuraminidase mRNA, complete cds	1,438 bp linear mRNA	U86145.1 GI:4099320
Influenza A virus (A/swine/Schleswig-Holstein/1/1992(H1N1)) mRNA for hemagglutinin precursor	1,730 bp linear mRNA	Z46435.1 GI:854216
Influenza A Virus (A/swine/Schleswig-Holstein/1/1993(H1N1)) mRNA for nucleoprotein	1,554 bp linear mRNA	Z46438.1 GI:854222
Influenza A virus (A/swine/Wisconsin/1/61(H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,778 bp linear mRNA	AF091307.1 GI:4585156
212. Influenza A virus (A/swine/Wisconsin/1/1967(H1N1)) nucleoprotein mRNA, complete cds	1,565 bp linear mRNA	M76607.1 GI:325086
Influenza A virus (A/swine/Wisconsin/1915/1988(H1N1)) nucleoprotein mRNA, complete cds	1,565 bp linear mRNA	M76608.1 GI:325088
Influenza A virus (A/swine/WI/1915/1988(H1N1)) nucleoprotein (segment 5) mRNA, complete cds	1,550 bp linear mRNA	L46850.1 GI:954757
Influenza A virus (A/Switzerland/8808/2002(H1N1)) partial m1 gene for matrix protein 1 and partial m2 gene for matrix protein 2, genomic RNA	729 bp linear mRNA	AJ532568.1 GI:31096461
Influenza A virus (A/human/Taiwan/0012/00(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362803.1 GI:14571975
Influenza A virus (A/human/Taiwan/0016/00(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362779.1 GI:14571927
Influenza A virus (A/Taiwan/0016/2000 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303752.1 GI:32330993

Influenza A virus (A/human/Taiwan/0030/00(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362780.1 GI:14571929
Influenza A virus (A/Taiwan/0030/2000 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303704.1 GI:32330897
Influenza A virus (A/Taiwan/0032/2002(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604804.1 GI:50727488
Influenza A virus (A/Taiwan/0061/2002(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604795.1 GI:50727470
Influenza A virus (A/Taiwan/0069/2002(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604803.1 GI:50727486
Influenza A virus (A/Taiwan/0078/2002(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604805.1 GI:50727490
Influenza A virus (A/Taiwan/0094/2002(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604797.1 GI:50727474
Influenza A virus (A/Taiwan/0116/2002(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604796.1 GI:50727472
Influenza A virus (A/human/Taiwan/0130/96(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362781.1 GI:14571931
Influenza A virus (A/Taiwan/0130/96 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303707.1 GI:32330903
Influenza A virus (A/human/Taiwan/0132/96(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362782.1 GI:14571933
Influenza A virus (A/Taiwan/0132/96 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303708.1 GI:32330905
Influenza A virus (A/human/Taiwan/0211/96(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362783.1 GI:14571935
Influenza A virus (A/Taiwan/0211/96 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303709.1 GI:32330907
Influenza A virus (A/human/Taiwan/0235/96(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362784.1 GI:14571937
Influenza A virus (A/Taiwan/0235/96 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303710.1 GI:32330909
Influenza A virus (A/human/Taiwan/0255/96(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362785.1 GI:14571939
Influenza A virus (A/Taiwan/0255/96 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303711.1 GI:32330911
Influenza A virus (A/human/Taiwan/0337/96(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362786.1 GI:14571941

Influenza A virus (A/human/Taiwan/0342/96(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362787.1 GI:14571943
Influenza A virus (A/Taiwan/0342/96 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303714.1 GI:32330917
Influenza A virus (A/human/Taiwan/0464/99(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362788.1 GI:14571945
Influenza A virus (A/human/Taiwan/0562/95(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362789.1 GI:14571947
Influenza A virus (A/Taiwan/0562/95 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303720.1 GI:32330929
Influenza A virus (A/human/Taiwan/0563/95(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362790.1 GI:14571949
Influenza A virus (A/Taiwan/0563/95 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303721.1 GI:32330931
Influenza A virus (A/human/Taiwan/0657/95(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362791.1 GI:14571951
Influenza A virus (A/Taiwan/0657/95 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303724.1 GI:32330937
Influenza A virus (A/Taiwan/0859/2002(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604801.1 GI:50727482
Influenza A virus (A/human/Taiwan/0892/99(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362792.1 GI:14571953
Influenza A virus (A/Taiwan/0983/2002(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604800.1 GI:50727480
Influenza A virus (A/Taiwan/1007/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068163.1 GI:158452199
Influenza A virus (A/Taiwan/1015/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068171.1 GI:158452215
Influenza A virus (A/Taiwan/112/1996-1(H1N1)) haemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AF026153.1 GI:2554950
Influenza A virus (A/Taiwan/112/1996-2(H1N1)) haemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AF026154.1 GI:2554952
Influenza A virus (A/Taiwan/117/1996-1(H1N1)) haemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AF026155.1 GI:2554954
Influenza A virus (A/Taiwan/117/1996-2(H1N1)) haemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AF026156.1 GI:2554956
Influenza A virus (A/Taiwan/117/1996-3(H1N1)) haemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AF026157.1 GI:2554958

Influenza A virus (A/Taiwan/118/1996-1(H1N1)) haemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AF026158.1 GI:2554960
Influenza A virus (A/Taiwan/118/1996-2(H1N1)) haemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AF026159.1 GI:2554962
Influenza A virus (A/Taiwan/118/1996-3(H1N1)) haemagglutinin (HA) mRNA, partial cds	1,176 bp linear mRNA	AF026160.1 GI:2554964
Influenza A virus (A/human/Taiwan/1184/99(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362793.1 GI:14571955
Influenza A virus (A/Taiwan/1184/99 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303726.1 GI:32330941
Influenza A virus (A/human/Taiwan/1190/95(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362794.1 GI:14571957
Influenza A virus (A/Taiwan/1190/95 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303727.1 GI:32330943
Influenza A virus (A/Taiwan/1523/2003(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604808.1 GI:50727496
Influenza A virus (A/Taiwan/1566/2003(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604806.1 GI:50727492
Influenza A virus (A/Taiwan/1769/96(H1N1)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138710.2 GI:4996871
Influenza A virus (A/Taiwan/1906/2002(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604799.1 GI:50727478
Influenza A virus (A/Taiwan/1922/2002(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604802.1 GI:50727484
Influenza A virus (A/Taiwan/2069/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068168.1 GI:158452209
Influenza A virus (A/Taiwan/2157/2001 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303733.1 GI:32330955
Influenza A virus (A/Taiwan/2175/2001 (H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AY303734.1 GI:32330957
Influenza A virus (A/human/Taiwan/2200/95(H1N1)) hemagglutinin (HA) mRNA, partial cds	564 bp linear mRNA	AF362795.1 GI:14571959
Influenza A virus (A/Taiwan/2200/95 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303737.1 GI:32330963
Influenza A virus (A/Taiwan/2966/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068170.1 GI:158452213
Influenza A virus (A/Taiwan/3168/2005(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068174.1 GI:158452221

Influenza A virus (A/human/Taiwan/3355/97(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362796.1 GI:14571961
Influenza A virus (A/Taiwan/3355/97 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303739.1 GI:32330967
Influenza A virus (A/Taiwan/3361/2001 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303740.1 GI:32330969
Influenza A virus (A/Taiwan/3361/2001 (H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AY303741.1 GI:32330971
Influenza A virus (A/Taiwan/3518/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068169.1 GI:158452211
Influenza A virus (A/human/Taiwan/3825/00(H1N1)) hemagglutinin (HA) mRNA, partial cds	581 bp linear mRNA	AF362797.1 GI:14571963
Influenza A virus (A/Taiwan/3896/2001 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303746.1 GI:32330981
Influenza A virus (A/Taiwan/3896/2001 (H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AY303747.1 GI:32330983
Influenza A virus (A/Taiwan/4050/2003(H1N1)) hemagglutinin mRNA, partial cds	494 bp linear mRNA	AY604807.1 GI:50727494
Influenza A virus (A/Taiwan/4054/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068160.1 GI:158452193
Influenza A virus (A/human/Taiwan/4360/99(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362798.1 GI:14571965
Influenza A virus (A/Taiwan/4360/99 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303748.1 GI:32330985
Influenza A virus (A/human/Taiwan/4415/99(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362799.1 GI:14571967
Influenza A virus (A/Taiwan/4415/99 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303749.1 GI:32330987
Influenza A virus (A/Taiwan/4509/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068165.1 GI:158452203
Influenza A virus (A/human/Taiwan/4845/99(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362800.1 GI:14571969
Influenza A virus (A/Taiwan/4845/99 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303750.1 GI:32330989
Influenza A virus (A/human/Taiwan/4943/99(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362801.1 GI:14571971
Influenza A virus (A/Taiwan/5010/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068167.1 GI:158452207

Influenza A virus (A/human/Taiwan/5063/99(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362802.1 GI:14571973
Influenza A virus (A/Taiwan/5063/99 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303751.1 GI:32330991
Influenza A virus (A/Taiwan/5084/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068166.1 GI:158452205
Influenza A virus (A/Taiwan/511/96(H1N1)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138708.2 GI:4996867
Influenza A virus (A/Taiwan/557/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068156.1 GI:158452185
Influenza A virus (A/Taiwan/562/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068159.1 GI:158452191
Influenza A virus (A/human/Taiwan/5779/98(H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AF362778.1 GI:14571925
Influenza A virus (A/Taiwan/5779/98 (H1N1)) polymerase basic protein 1 (PB1) mRNA, partial cds	303 bp linear mRNA	AY303702.1 GI:32330893
Influenza A virus (A/Taiwan/6025/2005(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068172.1 GI:158452217
Influenza A virus (A/Taiwan/607/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068157.1 GI:158452187
Influenza A virus (A/Taiwan/615/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068162.1 GI:158452197
Influenza A virus (A/Taiwan/645/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068164.1 GI:158452201
Influenza A virus (A/Taiwan/680/2005(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068173.1 GI:158452219
Influenza A virus (A/Taiwan/719/2006(H1N1)) hemagglutinin (HA) mRNA, partial cds	507 bp linear mRNA	EU068158.1 GI:158452189
Influenza A virus (A/Thailand/CU124/2006(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	EU021285.1 GI:154224724
Influenza A virus (A/Thailand/CU32/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021265.1 GI:154224704
Influenza A virus (A/Thailand/CU32/2006(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,698 bp linear mRNA	EU021264.1 GI:154224775
Influenza A virus (A/Thailand/CU41/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021247.1 GI:154224686
Influenza A virus (A/Thailand/CU41/2006(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,698 bp linear mRNA	EU021246.1 GI:154224757

Influenza A virus (A/Thailand/CU44/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021259.1 GI:154224698
Influenza A virus (A/Thailand/CU44/2006(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,698 bp linear mRNA	EU021258.1 GI:154224769
Influenza A virus (A/Thailand/CU51/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021255.1 GI:154224694
Influenza A virus (A/Thailand/CU51/2006(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,698 bp linear mRNA	EU021254.1 GI:154224765
Influenza A virus (A/Thailand/CU53/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021249.1 GI:154224688
Influenza A virus (A/Thailand/CU53/2006(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,698 bp linear mRNA	EU021248.1 GI:154224759
Influenza A virus (A/Thailand/CU57/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021257.1 GI:154224696
Influenza A virus (A/Thailand/CU57/2006(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,698 bp linear mRNA	EU021256.1 GI:154224767
Influenza A virus (A/Thailand/CU67/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021251.1 GI:154224690
Influenza A virus (A/Thailand/CU67/2006(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,698 bp linear mRNA	EU021250.1 GI:154224761
Influenza A virus (A/Thailand/CU68/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021261.1 GI:154224700
Influenza A virus (A/Thailand/CU68/2006(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,698 bp linear mRNA	EU021260.1 GI:154224771
Influenza A virus (A/Thailand/CU75/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021263.1 GI:154224702
Influenza A virus (A/Thailand/CU75/2006(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,698 bp linear mRNA	EU021262.1 GI:154224773
Influenza A virus (A/Thailand/CU88/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021253.1 GI:154224692
Influenza A virus (A/Thailand/CU88/2006(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,698 bp linear mRNA	EU021252.1 GI:154224763
Influenza A virus (A/turkey/England/647/1977(H1N1)) nucleoprotein mRNA, complete cds	1,565 bp linear mRNA	M76603.1 GI:325094
Influenza A virus (A/turkey/France/87075/87(H1N1)) N1 gene for neuraminidase, genomic RNA	1,445 bp linear mRNA	AJ416626.1 GI:39840719

Influenza A virus (A/turkey/Germany/3/91(H1N1)) mRNA for PB2 polymerase (partial)	394 bp linear mRNA	Z30272.1 GI:456652
Influenza A virus (A/turkey/Germany/3/91(H1N1)) mRNA for neuraminidase (UTR)	97 bp linear mRNA	Z30275.1 GI:530398
Influenza A virus (A/turkey/Germany/3/91(H1N1)) mRNA for PA polymerase	264 bp linear mRNA	Z30274.1 GI:530401
Influenza A virus (A/turkey/Germany/3/91(H1N1)) mRNA for PBI polymerase (partial)	247 bp linear mRNA	Z30273.1 GI:530403
Influenza A virus (A/turkey/Germany/3/91(H1N1)) mRNA for hemagglutinin HA1	1,038 bp linear mRNA	Z46441.1 GI:854218
Influenza A virus (A/turkey/Minnesota/1661/1981(H1N1)) hemagglutinin (HA) mRNA, partial cds	981 bp linear mRNA	U46941.1 GI:1912332
Influenza A virus (A/turkey/Minnesota/1661/81(H1N1)) segment 4 hemagglutinin precursor (HA) mRNA, complete cds	1,777 bp linear mRNA	AF091310.1 GI:4585162
Influenza A virus (A/turkey/North Carolina/1790/1988(H1N1)) nucleoprotein mRNA, complete cds	1,565 bp linear mRNA	M76609.1 GI:325096
Influenza A virus (A/Weiss/43 (H1N1)) neuraminidase (NA) gene, complete cds	1,410 bp linear mRNA	AF250365.2 GI:13260589
Influenza A virus (A/Wilson-Smith/1933(H1N1)) nucleocapsid protein (NP) mRNA, complete cds	1,497 bp linear mRNA	EU330203.1 GI:167989512
Influenza A virus (A/Wisconsin/3523/1988(H1N1)) neuraminidase (NA) mRNA, partial cds	241 bp linear mRNA	U47816.1 GI:1912352
Influenza A virus (A/Wisconsin/3623/1988(H1N1)) nucleoprotein mRNA, complete cds	1,565 bp linear mRNA	M76610.1 GI:325103
Influenza A virus (A/WI/4754/1994(H1N1)) PB1 (PB1) mRNA, partial cds	235 bp linear mRNA	U53156.1 GI:1399590
Influenza A virus (A/WI/4754/1994(H1N1)) PB2 (PB2) mRNA, partial cds	168 bp linear mRNA	U53158.1 GI:1399594
Influenza A virus (A/WI/4754/1994(H1N1)) PA (PA) mRNA, partial cds	621 bp linear mRNA	U53160.1 GI:1399598
Influenza A virus (A/WI/4754/1994(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,778 bp linear mRNA	U53162.1 GI:1399602
Influenza A virus (A/WI/4754/1994(H1N1)) NP (NP) mRNA, partial cds	200 bp linear mRNA	U53164.1 GI:1399606
Influenza A virus (A/WI/4754/1994(H1N1)) neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	U53166.1 GI:1399610
Influenza A virus (A/WI/4754/1994(H1N1)) M (M) mRNA, complete cds	1,027 bp linear mRNA	U53168.1 GI:1399614

Influenza A virus (A/WI/4754/1994(H1N1)) NS (NS) mRNA, complete cds	890 bp linear mRNA	U53170.1 GI:1399618
Influenza A virus (A/WI/4755/1994(H1N1)) PB1 (PB1) mRNA, partial cds	203 bp linear mRNA	U53157.1 GI:1399592
Influenza A virus (A/WI/4755/1994(H1N1)) PB2 (PB2) mRNA, partial cds	173 bp linear mRNA	U53159.1 GI:1399596
Influenza A virus (A/WI/4755/1994(H1N1)) PA (PA) mRNA, partial cds	621 bp linear mRNA	U53161.1 GI:1399600
Influenza A virus (A/WI/4755/1994(H1N1)) hemagglutinin (HA) mRNA, complete cds	1,778 bp linear mRNA	U53163.1 GI:1399604
Influenza A virus (A/WI/4755/1994(H1N1)) NP (NP) mRNA, partial cds	215 bp linear mRNA	U53165.1 GI:1399608
Influenza A virus (A/WI/4755/1994(H1N1)) neuraminidase (NA) mRNA, partial cds	209 bp linear mRNA	U53167.1 GI:1399612
Influenza A virus (A/WI/4755/1994(H1N1)) M (M) mRNA, complete cds	1,027 bp linear mRNA	U53169.1 GI:1399616
Influenza A virus (A/WI/4755/1994(H1N1)) NS (NS) mRNA, complete cds	890 bp linear mRNA	U53171.1 GI:1399620
Influenza A virus (A/WSN/33) segment 5 nucleocapsid protein (NP) mRNA, partial cds	543 bp linear mRNA	AF306656.1 GI:11935089

Table 7. Influenza H3N2 Antigens

Strain/Protein	Length	GenBank / GI Accession No.
1. Influenza A virus (A/Aichi/2/1968(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,704 bp linear mRNA	EF614248.1 GI:148910819
2. Influenza A virus (A/Aichi/2/1968(H3N2)) hemagglutinin (HA) mRNA, partial cds	1,698 bp linear mRNA	EF614249.1 GI:148910821
3. Influenza A virus (A/Aichi/2/1968(H3N2)) hemagglutinin (HA) mRNA, partial cds	1,698 bp linear mRNA	EF614250.1 GI:148910823
4. Influenza A virus (A/Aichi/2/1968(H3N2)) hemagglutinin (HA) mRNA, partial cds	1,698 bp linear mRNA	EF614251.1 GI:148910825
5. Influenza A virus (A/Akita/1/1995(H3N2)) haemagglutinin mRNA, partial cds	1,032 bp linear mRNA	U48444.1 GI:1574989
6. Influenza A virus (A/Beijing/32/1992(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46392.1 GI:609020
7. Influenza A virus (A/Canada/33312/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501516.1 GI:21314288
8. Influenza A virus (A/Charlottesville/10/99 (H3N2)) hemagglutinin mRNA, partial cds	987 bp linear mRNA	AF297094.1 GI:11228917
9. Influenza A virus (A/Charlottesville/49/99 (H3N2)) hemagglutinin mRNA, partial cds	987 bp linear mRNA	AF297096.1 GI:11228921
10. Influenza A virus (A/Charlottesville/69/99 (H3N2)) hemagglutinin mRNA, partial cds	987 bp linear mRNA	AF297097.1 GI:11228923
11. Influenza A virus (A/Charlottesville/73/99 (H3N2)) hemagglutinin mRNA, partial cds	987 bp linear mRNA	AF297095.1 GI:11228919

12. Influenza A virus (A/England/1/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46393.1 GI:609024
13. Influenza A virus (A/England/247/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46394.1 GI:609025
14. Influenza A virus (A/England/269/93(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46395.1 GI:609027
15. Influenza A virus (A/England/284/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46396.1 GI:609029
16. Influenza A virus (A/England/286/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46397.1 GI:609031
17. Influenza A virus (A/England/289/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46398.1 GI:609033
18. Influenza A virus (A/England/328/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46399.1 GI:609035
19. Influenza A virus (A/England/346/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46400.1 GI:609037
20. Influenza A virus (A/England/347/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46401.1 GI:609039
21. Influenza A virus (A/England/42/72(H3N2)) hemagglutinin mRNA, partial cds	1,091 bp linear mRNA	AF201875.1 GI:6470274
22. Influenza A virus (A/England/471/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46402.1 GI:609041
23. Influenza A virus (A/England/67/1994(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46403.1 GI:609043
24. Influenza A virus (A/England/68/1994(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46404.1 GI:609045
25. Influenza A virus (A/England/7/1994(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46405.1 GI:609047
28. Influenza A virus (A/Guangdong/25/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46406.1 GI:609049
29. Influenza A virus (A/Hong Kong/1/68(H3N2)) hemagglutinin mRNA, partial cds	1,091 bp linear mRNA	AF201874.1 GI:6470272
30. Influenza A virus (A/Hong Kong/1/1994(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46407.1 GI:609051
31. Influenza A virus (A/Hong Kong/1143/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382319.1 GI:14487957
32. Influenza A virus (A/Hong Kong/1143/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382320.1 GI:14487959
33. Influenza A virus (A/Hong Kong/1143/99(H3N2)) neuraminidase mRNA, complete cds	1,466 bp linear mRNA	AF382329.1 GI:14487977
34. Influenza A virus (A/Hong Kong/1143/99(H3N2)) neuraminidase mRNA, complete cds	1,466 bp linear mRNA	AF382330.1 GI:14487979
35. Influenza A virus (A/Hong Kong/1144/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AY035589.1 GI:14486403
36. Influenza A virus (A/Hong Kong/1144/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382321.1 GI:14487961
37. Influenza A virus (A/Hong Kong/1144/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382322.1 GI:14487963
38. Influenza A virus (A/Hong Kong/1144/99(H3N2)) neuraminidase mRNA, complete cds	1,466 bp linear mRNA	AF382331.1 GI:14487981
39. Influenza A virus (A/Hong Kong/1144/99(H3N2)) neuraminidase mRNA, complete cds	1,466 bp linear mRNA	AF382332.1 GI:14487983
40. Influenza A virus (A/Hong Kong/1179/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AY035590.1 GI:14486405
41. Influenza A virus (A/Hong Kong/1179/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382323.1 GI:14487965

42. Influenza A virus (A/Hong Kong/1179/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382324.1 GI:14487967
43. Influenza A virus (A/Hong Kong/1180/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AY035591.1 GI:14486407
44. Influenza A virus (A/Hong Kong/1180/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382325.1 GI:14487969
45. Influenza A virus (A/Hong Kong/1180/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382326.1 GI:14487971
46. Influenza A virus (A/Hong Kong/1182/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382327.1 GI:14487973
47. Influenza A virus (A/Hong Kong/1182/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382328.1 GI:14487975
48. Influenza A virus (A/Hong Kong/2/1994(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46408.1 GI:609055
49. Influenza A virus (A/Hong Kong/23/1992(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46410.1 GI:609053
50. Influenza A virus (A/Hong Kong/34/1990(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46409.1 GI:609057
51. Influenza A virus (A/England/286/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46397.1 GI:609031
52. Influenza A virus (A/England/289/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46398.1 GI:609033
53. Influenza A virus (A/England/328/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46399.1 GI:609035
54. Influenza A virus (A/England/346/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46400.1 GI:609037
55. Influenza A virus (A/England/347/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46401.1 GI:609039
56. Influenza A virus (A/England/42/72(H3N2)) hemagglutinin mRNA, partial cds	1,091 bp linear mRNA	AF201875.1 GI:6470274
57. Influenza A virus (A/England/471/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46402.1 GI:609041
58. Influenza A virus (A/England/67/1994(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46403.1 GI:609043
59. Influenza A virus (A/England/68/1994(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46404.1 GI:609045
60. Influenza A virus (A/England/7/1994(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46405.1 GI:609047
63. Influenza A virus (A/Guandong/28/1994(H3N2)) haemagglutinin mRNA, partial cds	1,032 bp linear mRNA	U48442.1 GI:1574985
64. Influenza A virus (A/Guangdong/25/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46406.1 GI:609049
65. Influenza A virus (A/Hebei/19/1995(H3N2)) haemagglutinin mRNA, partial cds	1,032 bp linear mRNA	U48447.1 GI:1574995
66. Influenza A virus (A/Hebei/41/1994(H3N2)) haemagglutinin mRNA, partial cds	1,032 bp linear mRNA	U48441.1 GI:1574983
67. Influenza A virus (A/Hong Kong/1/68(H3N2)) hemagglutinin mRNA, partial cds	1,091 bp linear mRNA	AF201874.1 GI:6470272
68. Influenza A virus (A/Hong Kong/1/1994(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46407.1 GI:609051
69. Influenza A virus (A/Hong Kong/1143/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AY035588.1 GI:14486401
70. Influenza A virus (A/Hong Kong/1143/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382319.1 GI:14487957
71. Influenza A virus (A/Hong Kong/1143/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382320.1 GI:14487959

72. Influenza A virus (A/Hong Kong/1143/99(H3N2)) neuraminidase mRNA, complete cds	1,466 bp linear mRNA	AF382329.1 GI:14487977
73. Influenza A virus (A/Hong Kong/1143/99(H3N2)) neuraminidase mRNA, complete cds	1,466 bp linear mRNA	AF382330.1 GI:14487979
74. Influenza A virus (A/Hong Kong/1144/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AY035589.1 GI:14486403
75. Influenza A virus (A/Hong Kong/1144/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382321.1 GI:14487961
76. Influenza A virus (A/Hong Kong/1144/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382322.1 GI:14487963
77. Influenza A virus (A/Hong Kong/1144/99(H3N2)) neuraminidase mRNA, complete cds	1,466 bp linear mRNA	AF382331.1 GI:14487981
78. Influenza A virus (A/Hong Kong/1144/99(H3N2)) neuraminidase mRNA, complete cds	1,466 bp linear mRNA	AF382332.1 GI:14487983
79. Influenza A virus (A/Hong Kong/1179/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AY035590.1 GI:14486405
80. Influenza A virus (A/Hong Kong/1179/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382323.1 GI:14487965
81. Influenza A virus (A/Hong Kong/1179/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382324.1 GI:14487967
82. Influenza A virus (A/Hong Kong/1180/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AY035591.1 GI:14486407
83. Influenza A virus (A/Hong Kong/1180/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382325.1 GI:14487969
84. Influenza A virus (A/Hong Kong/1180/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382326.1 GI:14487971
85. Influenza A virus (A/Hong Kong/1182/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AY035592.1 GI:14486409
86. Influenza A virus (A/Hong Kong/1182/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382327.1 GI:14487973
87. Influenza A virus (A/Hong Kong/1182/99(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382328.1 GI:14487975
88. Influenza A virus (A/Hong Kong/2/1994(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46408.1 GI:609055
89. Influenza A virus (A/Hong Kong/23/1992(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46410.1 GI:609053
90. Influenza A virus (A/Hong Kong/34/1990(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46409.1 GI:609057
91. Influenza A virus (A/Indiana/28170/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501534.1 GI:21314324
92. Influenza A virus (A/Kinmen/618/03(H3N2)) hemagglutinin (HA) mRNA, partial cds	529 bp linear mRNA	AY961997.1 GI:68138151
93. Influenza A virus (A/Kinmen/618/03(H3N2)) neuraminidase (NA) mRNA, partial cds	383 bp linear mRNA	AY973325.1 GI:70673206
94. Influenza A virus (A/Kinmen/618/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986986.1 GI:70728099
95. Influenza A virus (A/Kinmen/621/03(H3N2)) hemagglutinin (HA) mRNA, partial cds	545 bp linear mRNA	AY962017.1 GI:68138191
96. Influenza A virus (A/Kinmen/621/03(H3N2)) neuraminidase (NA) mRNA, partial cds	386 bp linear mRNA	AY973326.1 GI:70673208
97. Influenza A virus (A/Kinmen/621/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986987.1 GI:70728101
98. Influenza A virus (A/Kinmen/639/04(H3N2)) hemagglutinin (HA) mRNA, partial cds	786 bp linear mRNA	AY962008.1 GI:68138173
99. Influenza A virus (A/Kinmen/639/04(H3N2)) neuraminidase (NA) mRNA, partial cds	381 bp linear mRNA	AY973327.1 GI:70673210

100. Influenza A virus (A/Kinmen/639/04(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986988.1 GI:70728103
101. Influenza A virus (A/Kinmen/641/04(H3N2)) hemagglutinin (HA) mRNA, partial cds	596 bp linear mRNA	AY962004.1 GI:68138165
102. Influenza A virus (A/Kinmen/641/04(H3N2)) neuraminidase (NA) mRNA, partial cds	785 bp linear mRNA	AY973328.1 GI:70673212
103. Influenza A virus (A/Kinmen/642/04(H3N2)) hemagglutinin (HA) mRNA, partial cds	576 bp linear mRNA	AY962001.1 GI:68138159
104. Influenza A virus (A/Kinmen/642/04(H3N2)) neuraminidase (NA) mRNA, partial cds	580 bp linear mRNA	AY973329.1 GI:70673214
105. Influenza A virus (A/Kinmen/642/04(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986989.1 GI:70728105
106. Influenza A virus (A/Kinmen/645/04(H3N2)) hemagglutinin (HA) mRNA, partial cds	789 bp linear mRNA	AY962009.1 GI:68138175
107. Influenza A virus (A/Kinmen/645/04(H3N2)) neuraminidase (NA) mRNA, partial cds	581 bp linear mRNA	AY973330.1 GI:70673216
108. Influenza A virus (A/Kinmen/645/04(H3N2)) nucleoprotein (NP) mRNA, partial cds	981 bp linear mRNA	AY986990.1 GI:70728107
109. Influenza A virus (A/LosAngeles/2/1987(H3N2)) polymerase protein basic 2 (PB2) mRNA, complete cds	2,341 bp linear mRNA	U62543.1 GI:1480737
110. Influenza A virus (A/Madrid/252/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46411.1 GI:609067
111. Influenza A virus (A/Michigan/22568/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501531.1 GI:21314318
112. Influenza A virus (A/Michigan/22692/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501518.1 GI:21314292
113. Influenza A virus (A/Moscow/10/99(H3N2)) partial NS1 gene for non structural protein 1 and partial NS2 gene for non structural protein 2, genomic RNA	754 bp linear mRNA	AJ519454.1 GI:31096423
114. Influenza A virus (A/ningbo/17/2002(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AY138518.1 GI:24895178
115. Influenza A virus (A/ningbo/25/2002(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AY138517.1 GI:24895169
116. Influenza A virus (A/NT/60/68/29C(H3N2)) mRNA for haemagglutinin (HA1 and HA2 genes)	1,765 bp linear mRNA	V01103.1 GI:60800
117. Influenza A virus (A/Oklahoma/323/03(H3N2)) hemagglutinin mRNA, complete cds	1,701 bp linear mRNA	DQ059385.1 GI:66933143
118. Influenza A virus (A/Oklahoma/323/03(H3N2)) neuraminidase mRNA, complete cds	1,410 bp linear mRNA	DQ059384.2 GI:75859981
119. Influenza A virus (A/Panama/2007/99(H3N2)) partial NS1 gene for non structural protein 1 and partial NS2 gene for non structural protein 2, genomic RNA	766 bp linear mRNA	AJ519458.1 GI:31096435
120. Influenza A virus (A/Pennsalvanya/20109/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501526.1 GI:21314308
121. Influenza A virus (A/Philippines/2/82(H3N2)) hemagglutinin mRNA, partial cds	1,091 bp linear mRNA	AF233691.1 GI:7331124
122. Influenza A virus (A/Pingtung/303/04(H3N2)) hemagglutinin (HA) mRNA, partial cds	767 bp linear mRNA	AY962000.1 GI:68138157
123. Influenza A virus (A/Pingtung/303/04(H3N2)) neuraminidase (NA) mRNA, partial cds	783 bp linear mRNA	AY973331.1 GI:70673218
124. Influenza A virus (A/Pingtung/303/04(H3N2)) nucleoprotein (NP) mRNA, partial cds	928 bp linear mRNA	AY986991.1 GI:70728109
125. Influenza A virus (A/Pingtung/313/04(H3N2)) hemagglutinin (HA) mRNA, partial cds	788 bp linear mRNA	AY961999.1 GI:68138155
126. Influenza A virus (A/Pingtung/313/04(H3N2)) neuraminidase (NA) mRNA, partial cds	787 bp linear mRNA	AY973332.1 GI:70673220

127. Influenza A virus (A/Pingtung/313/04(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986992.1 GI:70728111
128. Influenza A virus (A/ruddy turnstone/Delaware/142/99 (H3N2)) nonfunctional matrix protein mRNA, partial sequence	927 bp linear mRNA	AY664458.1 GI:51011862
129. Influenza A virus (A/Scotland/142/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46413.1 GI:609059
130. Influenza A virus (A/Scotland/160/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46414.1 GI:609061
131. Influenza A virus (A/Scotland/173/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46416.1 GI:609063
132. Influenza A virus (A/Scotland/174/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46415.1 GI:609065
133. Influenza A virus (A/Scotland/2/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46412.1 GI:609069
134. Influenza A virus (A/Sendai/c182/1994(H3N2)) haemagglutinin mRNA, partial cds	1,032 bp linear mRNA	U48439.1 GI:1574979
135. Influenza A virus (A/Sendai/c373/1995(H3N2)) haemagglutinin mRNA, partial cds	1,032 bp linear mRNA	U48445.1 GI:1574991
136. Influenza A virus (A/Sendai/c384/1994(H3N2)) haemagglutinin mRNA, partial cds	1,032 bp linear mRNA	U48440.1 GI:1574981
137. Influenza A virus (A/Shangdong/9/1993(H3N2)) mRNA for haemagglutinin	1,041 bp linear mRNA	Z46417.1 GI:609071
138. Influenza A virus (A/Shanghai/11/1987/X99aE high yield reassortant(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	L19416.1 GI:348117
139. Influenza A virus (A/sw/Shizuoka/110/97(H3N2)) polymerase basic 2 (PB2) mRNA, complete cds	2,280 bp linear mRNA	AF225514.1 GI:27462098
140. Influenza A virus (A/sw/Shizuoka/110/97(H3N2)) polymerase basic 1 (PB1) mRNA, complete cds	2,274 bp linear mRNA	AF225518.1 GI:27462106
141. Influenza A virus (A/sw/Shizuoka/110/97(H3N2)) polymerase acidic (PA) mRNA, complete cds	2,151 bp linear mRNA	AF225522.1 GI:27462114
142. Influenza A virus (A/sw/Shizuoka/110/97(H3N2)) nucleoprotein (NP) mRNA, complete cds	1,497 bp linear mRNA	AF225534.1 GI:27462146
143. Influenza A virus (A/sw/Shizuoka/110/97(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	AF225538.1 GI:27462154
144. Influenza A virus (A/sw/Shizuoka/110/97(H3N2)) hemagglutinin (HA1) mRNA, partial cds	984 bp linear mRNA	AF225542.1 GI:27462162
145. Influenza A virus (A/sw/Shizuoka/115/97(H3N2)) polymerase basic 2 (PB2) mRNA, complete cds	2,280 bp linear mRNA	AF225515.1 GI:27462100
146. Influenza A virus (A/sw/Shizuoka/115/97(H3N2)) polymerase basic 1 (PB1) mRNA, complete cds	2,274 bp linear mRNA	AF225519.1 GI:27462108
147. Influenza A virus (A/sw/Shizuoka/115/97(H3N2)) polymerase acidic (PA) mRNA, complete cds	2,151 bp linear mRNA	AF225523.1 GI:27462116
148. Influenza A virus (A/sw/Shizuoka/115/97(H3N2)) nucleoprotein (NP) mRNA, complete cds	1,497 bp linear mRNA	AF225535.1 GI:27462148
149. Influenza A virus (A/sw/Shizuoka/115/97(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	AF225539.1 GI:27462156
150. Influenza A virus (A/sw/Shizuoka/115/97(H3N2)) hemagglutinin (HA1) mRNA, partial cds	984 bp linear mRNA	AF225543.1 GI:27462164
151. Influenza A virus (A/sw/Shizuoka/119/97(H3N2)) polymerase basic 2 (PB2) mRNA, complete cds	2,280 bp linear mRNA	AF225516.1 GI:27462102
152. Influenza A virus (A/sw/Shizuoka/119/97(H3N2)) polymerase basic 1 (PB1) mRNA, complete cds	2,274 bp linear mRNA	AF225520.1 GI:27462110
153. Influenza A virus (A/sw/Shizuoka/119/97(H3N2)) polymerase acidic (PA) mRNA, complete cds	2,151 bp linear mRNA	AF225524.1 GI:27462118
154. Influenza A virus (A/sw/Shizuoka/119/97(H3N2)) nucleoprotein (NP) mRNA, complete cds	1,497 bp linear mRNA	AF225536.1 GI:27462150

155. Influenza A virus (A/sw/Shizuoka/119/97(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	AF225540.1 GI:27462158
156. Influenza A virus (A/sw/Shizuoka/119/97(H3N2)) hemagglutinin (HA1) mRNA, partial cds	984 bp linear mRNA	AF225544.1 GI:27462166
159. Influenza A virus (A/swine/Bakum/IDT1769/2003(H3N2)) neuraminidase mRNA, complete cds	1,410 bp linear mRNA	EU163948.1 GI:157679552
163. Influenza A virus (A/swine/Fujian/668/01(H3N2)) nonfunctional hemagglutinin mRNA, complete sequence	1,738 bp linear mRNA	AY857957.1 GI:58042507
164. Influenza A virus PB2 gene for Polymerase 2 protein, genomic RNA, strain A/Swine/Italy/1523/98	2,280 bp linear mRNA	AJ311459.1 GI:13661041
165. Influenza A virus PB1 gene for Polymerase 1 protein, genomic RNA, strain A/Swine/Italy/1523/98	2,274 bp linear mRNA	AJ311460.1 GI:13661043
166. Influenza A virus (A/swine/Italy/1523/98(H3N2)) NS1 gene for non structural protein 1 and NS2 gene for non structural protein 2, genomic RNA	821 bp linear mRNA	AJ344024.1 GI:20068146
167. Influenza A virus (A/swine/Re220/92hp(H3N2)) neuraminidase mRNA, complete cds	1,465 bp linear mRNA	EU163949.1 GI:157679554
168. Influenza A virus (A/sw/Shizuoka/120/97(H3N2)) polymerase basic 2 (PB2) mRNA, complete cds	2,280 bp linear mRNA	AF225517.1 GI:27462104
169. Influenza A virus (A/sw/Shizuoka/120/97(H3N2)) polymerase basic 1 (PB1) mRNA, complete cds	2,274 bp linear mRNA	AF225521.1 GI:27462112
170. Influenza A virus (A/sw/Shizuoka/120/97(H3N2)) polymerase acidic (PA) mRNA, complete cds	2,151 bp linear mRNA	AF225525.1 GI:27462120
171. Influenza A virus (A/sw/Shizuoka/120/97(H3N2)) nucleoprotein (NP) mRNA, complete cds	1,497 bp linear mRNA	AF225537.1 GI:27462152
172. Influenza A virus (A/sw/Shizuoka/120/97(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	AF225541.1 GI:27462160
173. Influenza A virus (A/sw/Shizuoka/120/97(H3N2)) hemagglutinin (HA1) mRNA, partial cds	984 bp linear mRNA	AF225545.1 GI:27462168
174. Influenza A virus (A/Switzerland/7729/98(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AY032978.1 GI:14161723
175. Influenza A virus (A/Switzerland/7729/98(H3N2)) hemagglutinin mRNA, complete cds	1,762 bp linear mRNA	AF382318.1 GI:14487955
176. Influenza A virus (A/Tainan/704/03(H3N2)) hemagglutinin (HA) mRNA, partial cds	528 bp linear mRNA	AY962011.1 GI:68138179
177. Influenza A virus (A/Tainan/704/03(H3N2)) neuraminidase (NA) mRNA, partial cds	384 bp linear mRNA	AY973333.1 GI:70673222
178. Influenza A virus (A/Tainan/704/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986993.1 GI:70728113
179. Influenza A virus (A/Tainan/712/03(H3N2)) hemagglutinin (HA) mRNA, partial cds	519 bp linear mRNA	AY962012.1 GI:68138181
180. Influenza A virus (A/Tainan/712/03(H3N2)) neuraminidase (NA) mRNA, partial cds	383 bp linear mRNA	AY973334.1 GI:70673224
181. Influenza A virus (A/Tainan/712/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986994.1 GI:70728115
182. Influenza A virus (A/Tainan/722/03(H3N2)) hemagglutinin (HA) mRNA, partial cds	784 bp linear mRNA	AY962005.1 GI:68138167
183. Influenza A virus (A/Tainan/722/03(H3N2)) neuraminidase (NA) mRNA, partial cds	592 bp linear mRNA	AY973335.1 GI:70673226
184. Influenza A virus (A/Tainan/722/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	936 bp linear mRNA	AY986995.1 GI:70728117
185. Influenza A virus (A/Taipei/407/03(H3N2)) hemagglutinin (HA) mRNA, partial cds	788 bp linear mRNA	AY961998.1 GI:68138153
186. Influenza A virus (A/Taipei/407/03(H3N2)) neuraminidase (NA) mRNA, partial cds	787 bp linear mRNA	AY973336.1 GI:70673228
187. Influenza A virus (A/Taipei/407/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986996.1 GI:70728119

188. Influenza A virus (A/Taipei/416/03(H3N2)) hemagglutinin (HA) mRNA, partial cds	787 bp linear mRNA	AY962007.1 GI:68138171
189. Influenza A virus (A/Taipei/416/03(H3N2)) neuraminidase (NA) mRNA, partial cds	782 bp linear mRNA	AY973337.1 GI:70673230
190. Influenza A virus (A/Taipei/416/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986997.1 GI:70728121
191. Influenza A virus (A/Taiwan/0020/98 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303703.1 GI:32330895
192. Influenza A virus (A/Taiwan/0040/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604817.1 GI:50727514
193. Influenza A virus (A/Taiwan/0045/98 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303705.1 GI:32330899
194. Influenza A virus (A/human/Taiwan/0095/96(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362820.1 GI:15055140
195. Influenza A virus (A/Taiwan/0097/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604828.1 GI:50727536
196. Influenza A virus (A/Taiwan/0104/2001 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303706.1 GI:32330901
197. Influenza A virus (A/human/Taiwan/0118/98(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362805.1 GI:15055110
198. Influenza A virus (A/Taiwan/0122/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604823.1 GI:50727526
199. Influenza A virus (A/human/Taiwan/0149/00(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362806.1 GI:15055112
200. Influenza A virus (A/Taiwan/0275/2000 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303712.1 GI:32330913
201. Influenza A virus (A/Taiwan/0275/2000 (H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AY303713.1 GI:32330915
202. Influenza A virus (A/human/Taiwan/0293/98(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362807.1 GI:15055114
203. Influenza A virus (A/Taiwan/0346/98 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303715.1 GI:32330919
204. Influenza A virus (A/Taiwan/0379/2000 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303716.1 GI:32330921
205. Influenza A virus (A/Taiwan/0379/2000 (H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AY303717.1 GI:32330923
206. Influenza A virus (A/Taiwan/0388/2001(H3N2)) hemagglutinin (HA) mRNA, partial cds	791 bp linear mRNA	AY625729.1 GI:50604415
207. Influenza A virus (A/human/Taiwan/0389/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362808.1 GI:15055116
208. Influenza A virus (A/human/Taiwan/0423/98(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362809.1 GI:15055118
209. Influenza A virus (A/Taiwan/0423/98 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303718.1 GI:32330925
210. Influenza A virus (A/human/Taiwan/0464/98(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362810.1 GI:15055120
211. Influenza A virus (A/Taiwan/0464/98 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303719.1 GI:32330927
212. Influenza A virus (A/Taiwan/0568/2001(H3N2)) hemagglutinin (HA) mRNA, partial cds	791 bp linear mRNA	AY625730.1 GI:50604440
213. Influenza A virus (A/Taiwan/0570/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604822.1 GI:50727524
214. Influenza A virus (A/Taiwan/0572/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604827.1 GI:50727534
215. Influenza A virus (A/Taiwan/0578/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604821.1 GI:50727522

216. Influenza A virus (A/Taiwan/0583/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604820.1 GI:50727520
217. Influenza A virus (A/Taiwan/0646/2000 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303722.1 GI:32330933
218. Influenza A virus (A/Taiwan/0646/2000 (H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AY303723.1 GI:32330935
219. Influenza A virus (A/human/Taiwan/0830/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362811.1 GI:15055122
220. Influenza A virus (A/Taiwan/0964/2001(H3N2)) hemagglutinin (HA) mRNA, partial cds	791 bp linear mRNA	AY625731.1 GI:50604469
221. Influenza A virus (A/human/Taiwan/1008/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362812.1 GI:15055124
222. Influenza A virus (A/Taiwan/1008/99 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303725.1 GI:32330939
223. Influenza A virus (A/Taiwan/1219/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068138.1 GI:158452149
224. Influenza A virus (A/Taiwan/1315/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068125.1 GI:158452123
225. Influenza A virus (A/Taiwan/1511/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068153.1 GI:158452179
226. Influenza A virus (A/Taiwan/1533/2003(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068119.1 GI:158452111
227. Influenza A virus (A/human/Taiwan/1537/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362813.1 GI:15055126
228. Influenza A virus (A/Taiwan/1537/99 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303728.1 GI:32330945
229. Influenza A virus (A/Taiwan/1566/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604826.1 GI:50727532
230. Influenza A virus (A/Taiwan/1568/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604819.1 GI:50727518
231. Influenza A virus (A/Taiwan/158/2003(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068116.1 GI:158452105
232. Influenza A virus (A/Taiwan/1600/96(H3N2)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138709.2 GI:4996869
233. Influenza A virus (A/Taiwan/1613/2003(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068117.1 GI:158452107
234. Influenza A virus (A/Taiwan/1651/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068148.1 GI:158452169
235. Influenza A virus (A/human/Taiwan/1748/97(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362814.1 GI:15055128
236. Influenza A virus (A/Taiwan/1748/97 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303729.1 GI:32330947
237. Influenza A virus (A/Taiwan/179/96(H3N2)) matrix protein M1 (M) mRNA, partial cds	872 bp linear mRNA	AF138707.2 GI:4996865
238. Influenza A virus (A/Taiwan/1817/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068139.1 GI:158452151
239. Influenza A virus (A/Taiwan/1904/2003(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068154.1 GI:158452181
240. Influenza A virus (A/Taiwan/1921/2003(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068155.1 GI:158452183
241. Influenza A virus (A/human/Taiwan/1986/96(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362815.1 GI:15055130
242. Influenza A virus (A/Taiwan/1990/96 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303730.1 GI:32330949
243. Influenza A virus (A/Taiwan/1990/96 (H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AY303731.1 GI:32330951

244. Influenza A virus (A/Taiwan/20/98(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139938.1 GI:4972940
245. Influenza A virus (A/Taiwan/20/98(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140627.1 GI:4972988
246. Influenza A virus (A/Taiwan/20/98(H3N2)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138715.2 GI:4996879
247. Influenza A virus (A/human/Taiwan/2031/97(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362816.1 GI:15055132
248. Influenza A virus (A/Taiwan/2034/96(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139937.1 GI:4972938
249. Influenza A virus (A/Taiwan/2034/96(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140620.1 GI:4972974
250. Influenza A virus (A/Taiwan/2034/96(H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303732.1 GI:32330953
251. Influenza A virus (A/Taiwan/2040/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604818.1 GI:50727516
252. Influenza A virus (A/Taiwan/2072/2006(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068131.1 GI:158452135
253. Influenza A virus (A/Taiwan/21/98(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139934.1 GI:4972932
254. Influenza A virus (A/Taiwan/21/98(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140624.1 GI:4972982
255. Influenza A virus (A/Taiwan/21/98(H3N2)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138716.2 GI:4996881
256. Influenza A virus (A/Taiwan/2191/96(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139932.1 GI:4972928
257. Influenza A virus (A/Taiwan/2191/96(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140622.1 GI:4972978
258. Influenza A virus (A/Taiwan/2191/96(H3N2)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138711.3 GI:156147502
259. Influenza A virus (A/Taiwan/2192/96(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139936.1 GI:4972936
260. Influenza A virus (A/Taiwan/2192/96(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140626.1 GI:4972986
261. Influenza A virus (A/Taiwan/2195/96 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303735.1 GI:32330959
262. Influenza A virus (A/Taiwan/2195/96 (H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AY303736.1 GI:32330961
263. Influenza A virus (A/Taiwan/224/98(H3N2)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138718.2 GI:4996885
264. Influenza A virus (A/human/Taiwan/2548/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AF362817.1 GI:15055134
265. Influenza A virus (A/Taiwan/268/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068120.1 GI:158452113
266. Influenza A virus (A/Taiwan/3008/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068149.1 GI:158452171
267. Influenza A virus (A/Taiwan/3075/2003(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068152.1 GI:158452177
268. Influenza A virus (A/human/Taiwan/3083/00(H3N2)) hemagglutinin (HA) mRNA, partial cds	940 bp linear mRNA	AF362818.1 GI:15055136
269. Influenza A virus (A/Taiwan/3131/2002(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604811.1 GI:50727502
270. Influenza A virus (A/Taiwan/3154/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068145.1 GI:158452163
271. Influenza A virus (A/Taiwan/3187/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068141.1 GI:158452155

272. Influenza A virus (A/Taiwan/3245/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068134.1 GI:158452141
273. Influenza A virus (A/Taiwan/3294/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068133.1 GI:158452139
274. Influenza A virus (A/Taiwan/3351/97(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139935.1 GI:4972934
275. Influenza A virus (A/Taiwan/3351/97(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140625.1 GI:4972984
276. Influenza A virus (A/Taiwan/3351/97(H3N2)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138713.2 GI:4996875
277. Influenza A virus (A/Taiwan/3351/97(H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303738.1 GI:32330965
278. Influenza A virus (A/Taiwan/3387/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068132.1 GI:158452137
279. Influenza A virus (A/Taiwan/3396/97 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303742.1 GI:32330973
280. Influenza A virus (A/Taiwan/3396/97 (H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AY303743.1 GI:32330975
281. Influenza A virus (A/Taiwan/3427/97(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139930.1 GI:4972924
282. Influenza A virus (A/Taiwan/3427/97(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140619.1 GI:4972972
283. Influenza A virus (A/Taiwan/346/98(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139940.1 GI:4972944
284. Influenza A virus (A/Taiwan/346/98(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140787.1 GI:4972992
285. Influenza A virus (A/Taiwan/346/98(H3N2)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138719.2 GI:4996887
286. Influenza A virus (A/human/Taiwan/3460/00(H3N2)) truncated hemagglutinin (HA) mRNA, partial cds	942 bp linear mRNA	AF362819.1 GI:15055138
287. Influenza A virus (A/Taiwan/3469/97(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139933.1 GI:4972930
288. Influenza A virus (A/Taiwan/3469/97(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140623.1 GI:4972980
289. Influenza A virus (A/Taiwan/3469/97(H3N2)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138714.2 GI:4996877
290. Influenza A virus (A/Taiwan/3503/97 (H3N2)) polymerase basic protein 1 (PB1) mRNA, partial cds	297 bp linear mRNA	AY303744.1 GI:32330977
291. Influenza A virus (A/Taiwan/3503/97 (H3N2)) hemagglutinin (HA) mRNA, partial cds	844 bp linear mRNA	AY303745.1 GI:32330979
292. Influenza A virus (A/Taiwan/3513/96(H3N2)) matrix protein M1 (M) mRNA, partial cds	919 bp linear mRNA	AF138712.1 GI:4928900
293. Influenza A virus (A/Taiwan/3513/97(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139931.1 GI:4972926
294. Influenza A virus (A/Taiwan/3513/97(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140621.1 GI:4972976
295. Influenza A virus (A/Taiwan/3744/2002(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604814.1 GI:50727508
296. Influenza A virus (A/human/Taiwan/3760/00(H3N2)) hemagglutinin (HA) mRNA, partial cds	940 bp linear mRNA	AF362804.1 GI:15055108
297. Influenza A virus (A/Taiwan/3896/2001 (H1N1)) hemagglutinin (HA) mRNA, partial cds	561 bp linear mRNA	AY303747.1 GI:32330983
298. Influenza A virus (A/Taiwan/4050/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604825.1 GI:50727530
299. Influenza A virus (A/Taiwan/4063/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604824.1 GI:50727528

300. Influenza A virus (A/Taiwan/41/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068137.1 GI:158452147
301. Influenza A virus (A/Taiwan/45/98(H3N2)) H3 hemagglutinin (HA) mRNA, partial cds	861 bp linear mRNA	AF139939.1 GI:4972942
302. Influenza A virus (A/Taiwan/45/98(H3N2)) N2 neuraminidase (NA) mRNA, partial cds	392 bp linear mRNA	AF140628.1 GI:4972990
303. Influenza A virus (A/Taiwan/45/98(H3N2)) matrix protein M1 (M) mRNA, partial cds	875 bp linear mRNA	AF138717.2 GI:4996883
304. Influenza A virus (A/Taiwan/4548/2003(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068114.1 GI:158452101
305. Influenza A virus (A/Taiwan/4673/2002(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604813.1 GI:50727506
306. Influenza A virus (A/Taiwan/4680/2002(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604812.1 GI:50727504
307. Influenza A virus (A/Taiwan/4735/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068136.1 GI:158452145
308. Influenza A virus (A/Taiwan/4829/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068142.1 GI:158452157
309. Influenza A virus (A/Taiwan/4836/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068130.1 GI:158452133
310. Influenza A virus (A/Taiwan/4865/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068143.1 GI:158452159
311. Influenza A virus (A/Taiwan/4883/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068121.1 GI:158452115
312. Influenza A virus (A/Taiwan/4938/2002(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604809.1 GI:50727498
313. Influenza A virus (A/Taiwan/4954/2002(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604815.1 GI:50727510
314. Influenza A virus (A/Taiwan/4963/2002(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604810.1 GI:50727500
315. Influenza A virus (A/Taiwan/4987/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068122.1 GI:158452117
316. Influenza A virus (A/Taiwan/4990/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068127.1 GI:158452127
317. Influenza A virus (A/Taiwan/5/2003(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068118.1 GI:158452109
318. Influenza A virus (A/Taiwan/5153/2002(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604816.1 GI:50727512
319. Influenza A virus (A/Taiwan/5267/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068128.1 GI:158452129
320. Influenza A virus (A/Taiwan/556/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068146.1 GI:158452165
321. Influenza A virus (A/Taiwan/5694/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068126.1 GI:158452125
322. Influenza A virus (A/Taiwan/587/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068147.1 GI:158452167
323. Influenza A virus (A/Taiwan/592/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068151.1 GI:158452175
324. Influenza A virus (A/Taiwan/7099/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604829.1 GI:50727538
325. Influenza A virus (A/Taiwan/7100/2003(H3N2)) hemagglutinin mRNA, partial cds	791 bp linear mRNA	AY604830.1 GI:50727540
326. Influenza A virus (A/Taiwan/7196/2003(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068150.1 GI:158452173
327. Influenza A virus (A/Taiwan/7568/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068135.1 GI:158452143

328. Influenza A virus (A/Taiwan/7601/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068144.1 GI:158452161
329. Influenza A virus (A/Taiwan/7681/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068124.1 GI:158452121
330. Influenza A virus (A/Taiwan/7702/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068123.1 GI:158452119
331. Influenza A virus (A/Taiwan/7873/2005(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068129.1 GI:158452131
332. Influenza A virus (A/Taiwan/8/2003(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068115.1 GI:158452103
333. Influenza A virus (A/Taiwan/93/2004(H3N2)) hemagglutinin (HA) mRNA, partial cds	750 bp linear mRNA	EU068140.1 GI:158452153
334. Influenza A virus (A/Taoyuan/108/02(H3N2)) hemagglutinin (HA) mRNA, partial cds	528 bp linear mRNA	AY962016.1 GI:68138189
335. Influenza A virus (A/Taoyuan/108/02(H3N2)) neuraminidase (NA) mRNA, partial cds	754 bp linear mRNA	AY973338.1 GI:70673232
336. Influenza A virus (A/Taoyuan/108/02(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986998.1 GI:70728123
337. Influenza A virus (A/Thailand/CU124/2006(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	EU021285.1 GI:154224724
338. Influenza A virus (A/Thailand/CU124/2006(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,701 bp linear mRNA	EU021284.1 GI:154224795
339. Influenza A virus (A/Thailand/CU228/2006(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	EU021275.1 GI:154224714
340. Influenza A virus (A/Thailand/CU228/2006(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,701 bp linear mRNA	EU021274.1 GI:154224785
341. Influenza A virus (A/Thailand/CU23/2006(H3N2)) neuraminidase (NA) mRNA, partial cds	1,347 bp linear mRNA	EU021267.1 GI:154224706
342. Influenza A virus (A/Thailand/CU23/2006(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,701 bp linear mRNA	EU021266.1 GI:154224777
343. Influenza A virus (A/Thailand/CU231/2006(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	EU021283.1 GI:154224722
344. Influenza A virus (A/Thailand/CU231/2006(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,701 bp linear mRNA	EU021282.1 GI:154224793
345. Influenza A virus (A/Thailand/CU259/2006(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	EU021279.1 GI:154224718
346. Influenza A virus (A/Thailand/CU259/2006(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,701 bp linear mRNA	EU021278.1 GI:154224789
347. Influenza A virus (A/Thailand/CU260/2006(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	EU021281.1 GI:154224720
348. Influenza A virus (A/Thailand/CU260/2006(H3N2)) hemagglutinin (HA) mRNA, partial cds	1,129 bp linear mRNA	EU021280.1 GI:154224791
349. Influenza A virus (A/Thailand/CU272/2007(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	EU021271.1 GI:154224710
350. Influenza A virus (A/Thailand/CU272/2007(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,701 bp linear mRNA	EU021270.1 GI:154224781
351. Influenza A virus (A/Thailand/CU280/2007(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	EU021273.1 GI:154224712
352. Influenza A virus (A/Thailand/CU280/2007(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,701 bp linear mRNA	EU021272.1 GI:154224783
353. Influenza A virus (A/Thailand/CU282/2007(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	EU021277.1 GI:154224716
354. Influenza A virus (A/Thailand/CU282/2007(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,701 bp linear mRNA	EU021276.1 GI:154224787
355. Influenza A virus (A/Thailand/CU32/2006(H1N1)) neuraminidase (NA) mRNA, complete cds	1,413 bp linear mRNA	EU021265.1 GI:154224704

361. Influenza A virus (A/Thailand/CU46/2006(H3N2)) neuraminidase (NA) mRNA, complete cds	1,410 bp linear mRNA	EU021269.1 GI:154224708
362. Influenza A virus (A/Thailand/CU46/2006(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,701 bp linear mRNA	EU021268.1 GI:154224779
377. Influenza A virus (A/Tottori/849AM1AL3/1994(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	U77837.1 GI:2992515
378. Influenza A virus (A/Tottori/849AM2/1994(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	U77833.1 GI:2992507
379. Influenza A virus (A/Tottori/849AM2AL3/1994(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	U77839.1 GI:2992519
380. Influenza A virus (A/Tottori/849AM4/1994(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	U77835.1 GI:2992511
382. Influenza A virus (A/Tottori/872AM2/1994(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	U77834.1 GI:2992509
383. Influenza A virus (A/Tottori/872AM2AL3/1994(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	U77840.1 GI:2992521
384. Influenza A virus (A/Tottori/872AM4/1994(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	U77836.1 GI:2992513
385. Influenza A virus (A/Tottori/872K4/1994(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	U77832.1 GI:2992505
386. Influenza A virus (A/United Kingdom/26554/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501529.1 GI:21314314
387. Influenza A virus (A/United Kingdom/34300/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501527.1 GI:21314310
388. Influenza A virus (A/Utah/20997/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501533.1 GI:21314322
389. Influenza A virus (A/Victoria/3/75) segment 5 nucleoprotein mRNA, complete cds	1,565 bp linear mRNA	AF072545.1 GI:4218933
390. Influenza A virus (A/Vienna/47/96M(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,762 bp linear mRNA	AF017270.2 GI:14286338
391. Influenza A virus (A/Vienna/47/96V(H3N2)) hemagglutinin (HA) mRNA, complete cds	1,762 bp linear mRNA	AF017272.2 GI:15004991
392. Influenza A virus (A/Vienna/81/96V(H3N2)) hemagglutinin (HA) mRNA, partial cds	1,069 bp linear mRNA	AF017271.1 GI:2407251
393. Influenza A virus (A/Virginia/21712/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501532.1 GI:21314320
394. Influenza A virus (A/Virginia/21716/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501515.1 GI:21314286
395. Influenza A virus (A/Virginia/21735/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501530.1 GI:21314316
396. Influenza A virus (A/Virginia/21743/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501524.1 GI:21314304
397. Influenza A virus (A/Virginia/21754/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501519.1 GI:21314294
398. Influenza A virus (A/Virginia/21799/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501523.1 GI:21314302
399. Influenza A virus (A/Virginia/21817/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501525.1 GI:21314306
400. Influenza A virus (A/Virginia/21822/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501520.1 GI:21314296
401. Influenza A virus (A/Virginia/21828/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501528.1 GI:21314312
402. Influenza A virus (A/Virginia/21833/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501517.1 GI:21314290
403. Influenza A virus (A/Virginia/21845/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501522.1 GI:21314300

404. Influenza A virus (A/Virginia/21847/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501535.1 GI:21314326
405. Influenza A virus (A/Virginia/G1/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AF501521.1 GI:21314298
406. Influenza A virus (A/Yilan/508/03(H3N2)) neuraminidase (NA) mRNA, partial cds	755 bp linear mRNA	AY973339.1 GI:70673234
407. Influenza A virus (A/Yilan/508/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY986999.1 GI:70728125
408. Influenza A virus (A/Yilan/513/03(H3N2)) hemagglutinin (HA) mRNA, partial cds	740 bp linear mRNA	AY962015.1 GI:68138187
409. Influenza A virus (A/Yilan/513/03(H3N2)) neuraminidase (NA) mRNA, partial cds	396 bp linear mRNA	AY973340.1 GI:70673236
410. Influenza A virus (A/Yilan/513/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY987000.1 GI:70728127
411. Influenza A virus (A/Yilan/515/03(H3N2)) hemagglutinin (HA) mRNA, partial cds	511 bp linear mRNA	AY962010.1 GI:68138177
412. Influenza A virus (A/Yilan/515/03(H3N2)) neuraminidase (NA) mRNA, partial cds	394 bp linear mRNA	AY973341.1 GI:70673238
413. Influenza A virus (A/Yilan/516/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY987001.1 GI:70728129
414. Influenza A virus (A/Yilan/518/03(H3N2)) hemagglutinin (HA) mRNA, partial cds	530 bp linear mRNA	AY962006.1 GI:68138169
415. Influenza A virus (A/Yilan/518/03(H3N2)) neuraminidase (NA) mRNA, partial cds	397 bp linear mRNA	AY973342.1 GI:70673240
416. Influenza A virus (A/Yilan/518/03(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY987002.1 GI:70728131
417. Influenza A virus (A/Yilan/538/04(H3N2)) hemagglutinin (HA) mRNA, partial cds	777 bp linear mRNA	AY962002.1 GI:68138161
418. Influenza A virus (A/Yilan/538/04(H3N2)) neuraminidase (NA) mRNA, partial cds	783 bp linear mRNA	AY973343.1 GI:70673242
419. Influenza A virus (A/Yilan/538/04(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY987003.1 GI:70728133
420. Influenza A virus (A/Yilan/549/04(H3N2)) hemagglutinin (HA) mRNA, partial cds	788 bp linear mRNA	AY962003.1 GI:68138163
421. Influenza A virus (A/Yilan/549/04(H3N2)) neuraminidase (NA) mRNA, partial cds	779 bp linear mRNA	AY973344.1 GI:70673244
422. Influenza A virus (A/Yilan/549/04(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY987004.1 GI:70728135
423. Influenza A virus (A/Yilan/557/04(H3N2)) hemagglutinin (HA) mRNA, partial cds	776 bp linear mRNA	AY962013.1 GI:68138183
424. Influenza A virus (A/Yilan/557/04(H3N2)) neuraminidase (NA) mRNA, partial cds	796 bp linear mRNA	AY973345.1 GI:70673246
425. Influenza A virus (A/Yilan/557/04(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY987005.1 GI:70728137
426. Influenza A virus (A/Yilan/566/04(H3N2)) hemagglutinin (HA) mRNA, partial cds	753 bp linear mRNA	AY962014.1 GI:68138185
427. Influenza A virus (A/Yilan/566/04(H3N2)) neuraminidase (NA) mRNA, partial cds	808 bp linear mRNA	AY973346.1 GI:70673248
428. Influenza A virus (A/Yilan/566/04(H3N2)) nucleoprotein (NP) mRNA, partial cds	882 bp linear mRNA	AY987006.1 GI:70728139
429. Influenza A virus (A/zhejiang/06/99(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AY138513.1 GI:24895131
430. Influenza A virus (A/zhejiang/10/98(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AY138515.1 GI:24895149
431. Influenza A virus (A/zhejiang/11/2002(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AY138516.1 GI:24895159

432. Influenza A virus (A/zhejiang/12/99(H3N2)) hemagglutinin-like (HA) mRNA, partial sequence	987 bp linear mRNA	AY138514.1 GI:24895141
433. Influenza A virus (A/zhejiang/8/2002(H3N2)) hemagglutinin (HA) mRNA, partial cds	987 bp linear mRNA	AY138519.1 GI:24895188
434. Influenza A virus H3N2 strain A/Akita/1/94 nonstructural protein 1 and nonstructural protein 2 mRNAs, complete cds	840 bp linear mRNA	U65670.1 GI:3929405
435. Influenza A virus H3N2 strain A/Akita/1/95 nonstructural protein 1 and nonstructural protein 2 mRNAs, complete cds	840 bp linear mRNA	U65671.1 GI:3929408
436. Influenza A virus H3N2 strain A/Shiga/20/95 nonstructural protein 1 and nonstructural protein 2 mRNAs, complete cds	840 bp linear mRNA	U65673.1 GI:3929411
437. Influenza A virus H3N2 strain A/Miyagi/69/95 nonstructural protein 1 and nonstructural protein 2 mRNAs, complete cds	840 bp linear mRNA	U65674.1 GI:3929414
438. Influenza A virus H3N2 strain A/Hebei/19/95 nonstructural protein 1 and nonstructural protein 2 mRNAs, complete cds	840 bp linear mRNA	U65672.1 GI:6468319
A/Aichi/69/1994(H3N2) haemagglutinin		U48446.1
A/Bangkok/1/1979 (H3N2) hemagglutinin (HA)		AF201843.1
A/Beijing/353/89(H3) hemagglutinin (HA)		U97740.1
A/Beijing/353/1989(H3N2) haemagglutinin		Z46391.1
A/chicken/Singapore/2002(H3N2) M2 protein		EU014143.1
A/Christ Hospital/231/82(H3N2)) hemagglutinin (HA)		U77830.1
A/duck/Eastern China/36/2002(H3N2) segment 6 neuraminidase (NA)		EU429701.1
A/duck/Eastern China/160/2003(H3N2) segment 6 neuraminidase (NA)		EU429732.1
A/duck/Eastern China/848/2003(H3N2) segment 6 neuraminidase (NA)		EU429721.1
A/duck/Eastern China/770/2003(H3N2) segment 6 neuraminidase (NA)		EU429736.1
A/duck/Eastern China/855/2003(H3N2) segment 6 neuraminidase (NA)		EU429737.1
A/duck/Eastern China/875/2003(H3N2) segment 6 neuraminidase (NA)		EU429738.1
A/duck/Eastern China/901/2003(H3N2) segment 6 neuraminidase (NA)		EU429739.1
A/duck/Eastern China/866/2003(H3N2) segment 6 neuraminidase (NA)		EU429756.1
A/duck/Eastern China/857/2003(H3N2) segment 6 neuraminidase (NA)		EU429761.1
A/duck/Eastern China/852/2003(H3N2) segment 6 neuraminidase (NA)		EU429767.1
A/duck/Eastern China/838/2003(H3N2) segment 6 neuraminidase (NA)		EU429720.1
A/duck/Eastern China/6/2004(H3N2) segment 6 neuraminidase (NA)		EU429745.1
A/duck/Eastern China/03/2005(H3N2) segment 6 neuraminidase (NA)		EU429781.1
A/duck/Eastern China/02/2006(H3N2) segment 6 neuraminidase (NA)		EU429769.1
A/duck/Eastern China/04/2006(H3N2) segment 6 neuraminidase (NA)		EU429770.1
A/duck/Eastern China/21/2006(H3N2) segment 6 neuraminidase (NA)		EU429771.1
A/duck/Eastern China/23/2006(H3N2) segment 6 neuraminidase (NA)		EU429772.1
A/duck/Eastern China/31/2006(H3N2) segment 6 neuraminidase (NA)		EU429773.1
A/duck/Eastern China/35/2006(H3N2) segment 6 neuraminidase (NA)		EU429768.1
A/duck/Eastern China/42/2006(H3N2) segment 6 neuraminidase (NA)		EU429774.1
A/duck/Eastern China/53/2006(H3N2) segment 6 neuraminidase (NA)		EU429775.1
A/duck/Eastern China/60/2006(H3N2) segment 6 neuraminidase (NA)		EU429776.1
A/duck/Eastern China/62/2006(H3N2) segment 6 neuraminidase (NA)		EU429784.1
A/duck/Eastern China/63/2006(H3N2) segment 6 neuraminidase (NA)		EU429777.1
A/duck/Eastern China/142/2006(H3N2) segment 6 neuraminidase (NA)		EU429742.1
A/Dunedin/4/1973 (H3N2) hemagglutinin (HA)		AF201842.1

Table 8. Influenza H5N1 Antigens

1. Influenza A virus (A/chicken/Burkina Faso/01.03/2006(H5N1)) mRNA for non-structural protein (ns gene)	827 bp linear mRNA	AM503036.1 GI:147846308
2. Influenza A virus (A/chicken/Burkina Faso/13.1/2006(H5N1)) partial mRNA for matrix protein 1 (m1 gene)	990 bp linear mRNA	AM503007.1 GI:147846250
3. Influenza A virus (A/chicken/Burkina Faso/13.1/2006(H5N1)) mRNA for nucleoprotein (np gene)	1,529 bp linear mRNA	AM503029.1 GI:147846294

4. Influenza A virus (A/chicken/Burkina Faso/13.1/2006(H5N1)) mRNA for non-structural protein (ns gene)	827 bp linear mRNA	AM503037.1 GI:147846310
5. Influenza A virus (A/chicken/Burkina Faso/13.1/2006(H5N1)) partial mRNA for polymerase (pa gene)	2,169 bp linear mRNA	AM503046.1 GI:147846328
6. Influenza A virus (A/chicken/Burkina Faso/13.1/2006(H5N1)) partial mRNA for polymerase basic protein 1 (pb1 gene)	2,259 bp linear mRNA	AM503056.1 GI:147846348
7. Influenza A virus (A/chicken/Burkina Faso/13.1/2006(H5N1)) partial mRNA for polymerase basic protein 2 (pb2 gene)	2,315 bp linear mRNA	AM503067.1 GI:147846859
8. Influenza A virus (A/chicken/China/1/02(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,736 bp linear mRNA	DQ023145.1 GI:66775624
9. Influenza A virus (A/chicken/China/1/02(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,509 bp linear mRNA	DQ023146.1 GI:66775626
10. Influenza A virus (A/chicken/China/1/02(H5N1)) neuraminidase (NA) mRNA, complete cds	1,379 bp linear mRNA	DQ023147.1 GI:66775628
11. Influenza A virus (A/chicken/Crimea/04/2005(H5N1)) matrix protein (M) mRNA, complete cds	999 bp linear mRNA	DQ650660.1 GI:109692767
12. Influenza A virus (A/chicken/Crimea/04/2005(H5N1)) nonstructural protein (NS) mRNA, complete cds	850 bp linear mRNA	DQ650662.1 GI:109692771
13. Influenza A virus (A/chicken/Crimea/08/2005(H5N1)) matrix protein (M) mRNA, complete cds	994 bp linear mRNA	DQ650664.1 GI:109692775
14. Influenza A virus (A/chicken/Crimea/08/2005(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,532 bp linear mRNA	DQ650666.1 GI:109692779
15. Influenza A virus (A/chicken/Crimea/08/2005(H5N1)) nonstructural protein (NS) mRNA, complete cds	850 bp linear mRNA	DQ650667.1 GI:109692781
16. Influenza A virus (A/chicken/Crimea/08/2005(H5N1)) polymerase acidic protein (PA) mRNA, complete cds	2,208 bp linear mRNA	DQ650668.1 GI:109692783
17. Influenza A virus (A/chicken/Crimea/08/2005(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,305 bp linear mRNA	DQ650670.1 GI:109692787
18. Influenza A virus (A/chicken/Dovolnoe/03/2005(H5N1)) hemagglutinin (HA) mRNA, partial cds	1,015 bp linear mRNA	DQ676838.1 GI:108782527
20. Influenza A virus (A/chicken/Guangxi/12/2004(H5N1)) polymerase PB2 mRNA, complete cds	2,341 bp linear mRNA	DQ366327.1 GI:86753731
21. Influenza A virus (A/chicken/Guangxi/12/2004(H5N1)) polymerase PB1 mRNA, complete cds	2,341 bp linear mRNA	DQ366328.1 GI:86753741
22. Influenza A virus (A/chicken/Guangxi/12/2004(H5N1)) PA protein mRNA, complete cds	2,233 bp linear mRNA	DQ366329.1 GI:86753751
23. Influenza A virus (A/chicken/Guangxi/12/2004(H5N1)) nucleocapsid mRNA, complete cds	1,565 bp linear mRNA	DQ366331.1 GI:86753771
24. Influenza A virus (A/chicken/Guangxi/12/2004(H5N1)) matrix protein mRNA, complete cds	1,027 bp linear mRNA	DQ366333.1 GI:86753791
25. Influenza A virus (A/chicken/Hong Kong/258/97(H5N1)) hemagglutinin mRNA, complete cds	1,718 bp linear mRNA	AF057291.1 GI:3068720
26. Influenza A virus (A/chicken/Hong Kong/258/97(H5N1)) neuraminidase mRNA, partial cds	1,318 bp linear mRNA	AF057292.1 GI:3068722
27. Influenza A virus (A/chicken/Hong Kong/258/97(H5N1)) nucleoprotein mRNA, complete cds	1,508 bp linear mRNA	AF057293.1 GI:3068724
28. Influenza A virus (A/Chicken/Hong Kong/728/97 (H5N1)) hemagglutinin H5 mRNA, complete cds	1,726 bp linear mRNA	AF082034.1 GI:4240435
29. Influenza A virus (A/Chicken/Hong Kong/786/97 (H5N1)) hemagglutinin H5 mRNA, complete cds	1,726 bp linear mRNA	AF082035.1 GI:4240437
30. Influenza A virus (A/chicken/Hong Kong/915/97(H5N1)) hemagglutinin H5 mRNA, complete cds	1,726 bp linear mRNA	AF082036.1 GI:4240439
31. Influenza A virus (A/chicken/Hong Kong/990/97 (H5N1)) hemagglutinin H5 mRNA, partial cds	1,091 bp linear mRNA	AF082037.1 GI:4240441
32. Influenza A virus (A/chicken/Krasnodar/01/2006(H5N1)) matrix protein 1 (M) mRNA, complete cds	1,002 bp linear mRNA	DQ676835.1 GI:108782521

33. Influenza A virus (A/chicken/Krasnodar/01/2006(H5N1)) nonstructural protein (NS) mRNA, complete cds	850 bp linear mRNA	DQ676837.1 GI:108782525
34. Influenza A virus (A/chicken/Kurgan/05/2005(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,754 bp linear mRNA	DQ449632.1 GI:90289625
35. Influenza A virus (A/chicken/Kurgan/05/2005(H5N1)) matrix protein 1 (M) mRNA, complete cds	1,002 bp linear mRNA	DQ449633.1 GI:90289627
36. Influenza A virus (A/chicken/Kurgan/05/2005(H5N1)) neuraminidase (NA) mRNA, complete cds	1,373 bp linear mRNA	DQ449634.1 GI:90289629
37. Influenza A virus (A/chicken/Kurgan/05/2005(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,540 bp linear mRNA	DQ449635.1 GI:90289631
38. Influenza A virus (A/chicken/Kurgan/05/2005(H5N1)) nonstructural protein (NS) mRNA, complete cds	850 bp linear mRNA	DQ449636.1 GI:90289633
39. Influenza A virus (A/chicken/Kurgan/05/2005(H5N1)) polymerase acidic protein (PA) mRNA, complete cds	2,208 bp linear mRNA	DQ449637.1 GI:90289635
40. Influenza A virus (A/chicken/Kurgan/05/2005(H5N1)) polymerase basic protein 1 (PB1) mRNA, complete cds	2,316 bp linear mRNA	DQ449638.1 GI:90289637
41. Influenza A virus (A/chicken/Kurgan/05/2005(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,316 bp linear mRNA	DQ449639.1 GI:90289646
42. Influenza A virus (A/chicken/Lobzenko/01/2008(H5N1)) hemagglutinin (HA) mRNA, partial cds	184 bp linear mRNA	EU447276.1 GI:168998217
43. Influenza A virus (A/chicken/Mahachkala/05/2006(H5N1)) matrix protein 1 (M) mRNA, complete cds	1,002 bp linear mRNA	DQ676831.1 GI:108782513
44. Influenza A virus (A/chicken/Mahachkala/05/2006(H5N1)) nonstructural protein (NS) mRNA, complete cds	850 bp linear mRNA	DQ676833.1 GI:108782517
45. Influenza A virus (A/chicken/Nigeria/AB13/2006(H5N1)) mRNA for nucleoprotein (np gene)	1,531 bp linear mRNA	AM503030.1 GI:147846296
46. Influenza A virus (A/chicken/Nigeria/AB13/2006(H5N1)) mRNA for non-structural protein (ns gene)	827 bp linear mRNA	AM503040.1 GI:147846316
47. Influenza A virus (A/chicken/Nigeria/AB13/2006(H5N1)) partial mRNA for polymerase (pa gene)	2,169 bp linear mRNA	AM503051.1 GI:147846338
48. Influenza A virus (A/chicken/Nigeria/AB13/2006(H5N1)) partial mRNA for polymerase basic protein 1 (pb1 gene)	2,259 bp linear mRNA	AM503060.1 GI:147846845
49. Influenza A virus (A/chicken/Nigeria/AB13/2006(H5N1)) partial mRNA for polymerase basic protein 2 (pb2 gene)	2,315 bp linear mRNA	AM503071.1 GI:147846867
70. Influenza A virus (A/chicken/Hong Kong/3123.1/2002(H5N1)) neuraminidase (NA) mRNA, partial cds	1,055 bp linear mRNA	DQ250158.1 GI:82412012
75. Influenza A virus (A/chicken/Krasnodar/01/2006(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,754 bp linear mRNA	DQ676834.1 GI:108782519
78. Influenza A virus (A/chicken/Krasnodar/01/2006(H5N1)) neuraminidase (NA) mRNA, complete cds	1,373 bp linear mRNA	DQ676836.2 GI:115520953
91. Influenza A virus (A/chicken/Lobzenko/01/2008(H5N1)) hemagglutinin (HA) mRNA, partial cds	184 bp linear mRNA	EU447276.1 GI:168998217
92. Influenza A virus (A/chicken/Mahachkala/05/2006(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,683 bp linear mRNA	DQ676830.1 GI:108782511
94. Influenza A virus (A/chicken/Mahachkala/05/2006(H5N1)) neuraminidase (NA) mRNA, complete cds	1,373 bp linear mRNA	DQ676832.1 GI:108782515
96. Influenza A virus (A/chicken/Malaysia/01/2004(H5N1)) neuramidase (NA) mRNA, partial cds	433 bp linear mRNA	DQ096567.1 GI:69145364
97. Influenza A virus (A/chicken/Nigeria/AB13/2006(H5N1)) partial mRNA for hemagglutinin (ha gene)	1,722 bp linear mRNA	AM503002.1 GI:147846240
98. Influenza A virus (A/chicken/Nigeria/AB13/2006(H5N1)) partial mRNA for neuraminidase (na gene)	1,329 bp linear mRNA	AM503020.1 GI:147846276
105. Influenza A virus (A/chicken/Nigeria/AB14/2006(H5N1)) partial mRNA for hemagglutinin (ha gene)	1,719 bp linear mRNA	AM503003.1 GI:147846242
106. Influenza A virus (A/chicken/Nigeria/AB14/2006(H5N1)) partial mRNA for matrix protein 1 (m1 gene)	953 bp linear mRNA	AM503011.1 GI:147846258

107. Influenza A virus (A/chicken/Nigeria/AB14/2006(H5N1)) partial mRNA for neuraminidase (na gene)	1,343 bp linear mRNA	AM503025.1 GI:147846286
108. Influenza A virus (A/chicken/Nigeria/AB14/2006(H5N1)) mRNA for non-structural protein (ns gene)	827 bp linear mRNA	AM503041.1 GI:147846318
109. Influenza A virus (A/chicken/Nigeria/AB14/2006(H5N1)) partial mRNA for polymerase (pa gene)	2,169 bp linear mRNA	AM503054.1 GI:147846344
110. Influenza A virus (A/chicken/Nigeria/AB14/2006(H5N1)) partial mRNA for polymerase basic protein 1 (pb1 gene)	2,259 bp linear mRNA	AM503061.1 GI:147846847
111. Influenza A virus (A/chicken/Nigeria/AB14/2006(H5N1)) partial mRNA for polymerase basic protein 2 (pb2 gene)	2,315 bp linear mRNA	AM503072.1 GI:147846869
112. Influenza A virus (A/chicken/Nigeria/AB14/2006(H5N1)) mRNA for nucleoprotein (np gene)	1,548 bp linear mRNA	AM503034.2 GI:149773117
113. Influenza A virus (A/chicken/Nigeria/BA210/2006(H5N1)) partial mRNA for neuraminidase (na gene)	1,342 bp linear mRNA	AM503022.1 GI:147846280
114. Influenza A virus (A/chicken/Nigeria/BA211/2006(H5N1)) partial mRNA for neuraminidase (na gene)	1,321 bp linear mRNA	AM503021.1 GI:147846278
115. Influenza A virus (A/chicken/Nigeria/BA211/2006(H5N1)) partial mRNA for polymerase basic protein 2 (pb2 gene)	2,315 bp linear mRNA	AM503073.1 GI:147846871
116. Influenza A virus (A/chicken/Nigeria/FA4/2006(H5N1)) partial mRNA for hemagglutinin (ha gene)	1,717 bp linear mRNA	AM503004.1 GI:147846244
117. Influenza A virus (A/chicken/Nigeria/FA4/2006(H5N1)) partial mRNA for matrix protein 1 (m1 gene)	989 bp linear mRNA	AM503013.1 GI:147846262
118. Influenza A virus (A/chicken/Nigeria/FA4/2006(H5N1)) partial mRNA for neuraminidase (na gene)	1,321 bp linear mRNA	AM503026.1 GI:147846288
119. Influenza A virus (A/chicken/Nigeria/FA4/2006(H5N1)) mRNA for non-structural protein (ns gene)	827 bp linear mRNA	AM503045.1 GI:147846326
120. Influenza A virus (A/chicken/Nigeria/FA4/2006(H5N1)) partial mRNA for polymerase (pa gene)	2,169 bp linear mRNA	AM503055.1 GI:147846346
121. Influenza A virus (A/chicken/Nigeria/FA4/2006(H5N1)) partial mRNA for polymerase basic protein 1 (pb1 gene)	2,259 bp linear mRNA	AM503064.1 GI:147846853
122. Influenza A virus (A/chicken/Nigeria/FA4/2006(H5N1)) partial mRNA for polymerase basic protein 2 (pb2 gene)	2,224 bp linear mRNA	AM503074.1 GI:147846873
123. Influenza A virus (A/chicken/Nigeria/FA6/2006(H5N1)) partial mRNA for hemagglutinin (ha gene)	1,717 bp linear mRNA	AM502998.1 GI:147846232
124. Influenza A virus (A/chicken/Nigeria/FA6/2006(H5N1)) partial mRNA for matrix protein 1 (m1 gene)	965 bp linear mRNA	AM503012.1 GI:147846260
125. Influenza A virus (A/chicken/Nigeria/FA6/2006(H5N1)) partial mRNA for neuraminidase (na gene)	1,327 bp linear mRNA	AM503023.1 GI:147846282
126. Influenza A virus (A/chicken/Nigeria/FA6/2006(H5N1)) mRNA for nucleoprotein (np gene)	1,543 bp linear mRNA	AM503031.1 GI:147846298
127. Influenza A virus (A/chicken/Nigeria/FA6/2006(H5N1)) partial mRNA for polymerase (pa gene)	2,169 bp linear mRNA	AM503052.1 GI:147846340
128. Influenza A virus (A/chicken/Nigeria/FA6/2006(H5N1)) partial mRNA for polymerase basic protein 1 (pb1 gene)	2,259 bp linear mRNA	AM503063.1 GI:147846851
129. Influenza A virus (A/chicken/Nigeria/FA7/2006(H5N1)) partial mRNA for hemagglutinin (ha gene)	1,710 bp linear mRNA	AM502999.1 GI:147846234
130. Influenza A virus (A/chicken/Nigeria/FA7/2006(H5N1)) partial mRNA for matrix protein 1 (m1 gene)	1,001 bp linear mRNA	AM503009.1 GI:147846254
131. Influenza A virus (A/chicken/Nigeria/FA7/2006(H5N1)) partial mRNA for neuraminidase (na gene)	1,331 bp linear mRNA	AM503018.1 GI:147846272
132. Influenza A virus (A/chicken/Nigeria/FA7/2006(H5N1)) mRNA for nucleoprotein (np gene)	1,531 bp linear mRNA	AM503035.1 GI:147846306
133. Influenza A virus (A/chicken/Nigeria/FA7/2006(H5N1)) mRNA for non-structural protein (ns gene)	827 bp linear mRNA	AM503042.1 GI:147846320
134. Influenza A virus (A/chicken/Nigeria/FA7/2006(H5N1)) partial mRNA for polymerase (pa gene)	2,169 bp linear mRNA	AM503049.1 GI:147846334

135. Influenza A virus (A/chicken/Nigeria/FA7/2006(H5N1)) partial mRNA for polymerase basic protein 1 (pb1 gene)	2,259 bp linear mRNA	AM503057.1 GI:147846350
136. Influenza A virus (A/chicken/Nigeria/FA7/2006(H5N1)) partial mRNA for polymerase basic protein 2 (pb2 gene)	2,315 bp linear mRNA	AM503068.1 GI:147846861
137. Influenza A virus (A/chicken/Nigeria/IF10/2006(H5N1)) partial mRNA for hemagglutinin (ha gene)	1,714 bp linear mRNA	AM503001.1 GI:147846238
138. Influenza A virus (A/chicken/Nigeria/IF10/2006(H5N1)) partial mRNA for matrix protein 1 (m1 gene)	990 bp linear mRNA	AM503010.1 GI:147846256
139. Influenza A virus (A/chicken/Nigeria/IF10/2006(H5N1)) partial mRNA for neuraminidase (na gene)	1,332 bp linear mRNA	AM503024.1 GI:147846284
140. Influenza A virus (A/chicken/Nigeria/IF10/2006(H5N1)) mRNA for non-structural protein (ns gene)	827 bp linear mRNA	AM503044.1 GI:147846324
141. Influenza A virus (A/chicken/Nigeria/IF10/2006(H5N1)) partial mRNA for polymerase (pa gene)	2,169 bp linear mRNA	AM503053.1 GI:147846342
142. Influenza A virus (A/chicken/Nigeria/IF10/2006(H5N1)) partial mRNA for polymerase basic protein 1 (pb1 gene)	2,259 bp linear mRNA	AM503059.1 GI:147846843
143. Influenza A virus (A/chicken/Nigeria/IF10/2006(H5N1)) partial mRNA for polymerase basic protein 2 (pb2 gene)	2,315 bp linear mRNA	AM503069.1 GI:147846863
144. Influenza A virus (A/chicken/Nigeria/IF10/2006(H5N1)) mRNA for nucleoprotein (np gene)	1,550 bp linear mRNA	AM503033.2 GI:149773115
145. Influenza A virus (A/chicken/Nigeria/OD8/2006(H5N1)) partial mRNA for hemagglutinin (ha gene)	1,719 bp linear mRNA	AM503005.1 GI:147846246
146. Influenza A virus (A/chicken/Nigeria/OD8/2006(H5N1)) partial mRNA for matrix protein 1 (m1 gene)	989 bp linear mRNA	AM503014.1 GI:147846264
147. Influenza A virus (A/chicken/Nigeria/OD9/2006(H5N1)) partial mRNA for hemagglutinin (ha gene)	1,720 bp linear mRNA	AM503000.1 GI:147846236
148. Influenza A virus (A/chicken/Nigeria/OD9/2006(H5N1)) partial mRNA for matrix protein 1 (m1 gene)	988 bp linear mRNA	AM503015.1 GI:147846266
149. Influenza A virus (A/chicken/Nigeria/OD9/2006(H5N1)) partial mRNA for neuraminidase (na gene)	1,330 bp linear mRNA	AM503019.1 GI:147846274
150. Influenza A virus (A/chicken/Nigeria/OD9/2006(H5N1)) mRNA for nucleoprotein (np gene)	1,531 bp linear mRNA	AM503032.1 GI:147846300
151. Influenza A virus (A/chicken/Nigeria/OD9/2006(H5N1)) mRNA for non-structural protein (ns gene)	827 bp linear mRNA	AM503043.1 GI:147846322
152. Influenza A virus (A/chicken/Nigeria/OD9/2006(H5N1)) partial mRNA for polymerase (pa gene)	2,169 bp linear mRNA	AM503050.1 GI:147846336
153. Influenza A virus (A/chicken/Nigeria/OD9/2006(H5N1)) partial mRNA for polymerase basic protein 1 (pb1 gene)	2,259 bp linear mRNA	AM503058.1 GI:147846841
154. Influenza A virus (A/chicken/Nigeria/OD9/2006(H5N1)) partial mRNA for polymerase basic protein 2 (pb2 gene)	2,315 bp linear mRNA	AM503070.1 GI:147846865
155. Influenza A virus (A/chicken/Scotland/59(H5N1)) mRNA for haemagglutinin precursor	1,768 bp linear mRNA	X07869.1 GI:60482
156. Influenza A virus (A/chicken/Scotland/59(H5N1)) N1 gene for neuraminidase, genomic RNA	1,445 bp linear mRNA	AJ416625.1 GI:39840717
161. Influenza A virus (A/chicken/zz/02/2004(H5N1)) nucleoprotein mRNA, complete cds	1,497 bp linear mRNA	DQ208502.1 GI:77158587
162. Influenza A virus (A/common coot/Switzerland/V544/2006(H5N1)) hemagglutinin (HA) gene, complete cds	1,707 bp linear mRNA	EF110519.1 GI:119394676
163. Influenza A virus (A/domestic goose/Pavlodar/1/2005(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,735 bp linear mRNA	EU190482.1 GI:158516739
164. Influenza A virus (A/duck/Eastern China/145/2003(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,401 bp linear mRNA	EU429750.1 GI:167859465
165. Influenza A virus (A/duck/Eastern China/150/2003(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,407 bp linear mRNA	EU429731.1 GI:167859427
166. Influenza A virus (A/duck/Eastern China/22/2005(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,398 bp linear mRNA	EU429783.1 GI:167859531

167. Influenza A virus (A/duck/Eastern China/304/2002(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,398 bp linear mRNA	EU429747.1 GI:167859459
168. Influenza A virus (A/duck/Eastern China/318/2002(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,401 bp linear mRNA	EU429727.1 GI:167859419
169. Influenza A virus (A/duck/Eastern China/37/2006(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,399 bp linear mRNA	EU429778.1 GI:167859521
170. Influenza A virus (A/duck/Eastern China/40/2005(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,398 bp linear mRNA	EU429757.1 GI:167859479
171. Influenza A virus (A/duck/Eastern China/48/2006(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,398 bp linear mRNA	EU429779.1 GI:167859523
172. Influenza A virus (A/duck/Eastern China/51/2005(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,398 bp linear mRNA	EU429763.1 GI:167859491
173. Influenza A virus (A/duck/Eastern China/54/2005(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,398 bp linear mRNA	EU429758.1 GI:167859481
174. Influenza A virus (A/duck/Eastern China/58/2005(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,398 bp linear mRNA	EU429764.1 GI:167859493
175. Influenza A virus (A/duck/Eastern China/59/2005(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,398 bp linear mRNA	EU429759.1 GI:167859483
176. Influenza A virus (A/duck/Eastern China/89/2005(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,398 bp linear mRNA	EU429765.1 GI:167859495
177. Influenza A virus (A/duck/Eastern China/89/2006(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,399 bp linear mRNA	EU429785.1 GI:167859535
178. Influenza A virus (A/duck/Eastern China/97/2001(H5N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,398 bp linear mRNA	EU429717.1 GI:167859399
179. Influenza A virus (A/duck/Fujian/01/2002(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585504.1 GI:47156226
180. Influenza A virus (A/duck/Fujian/01/2002(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585378.1 GI:47156310
181. Influenza A virus (A/duck/Fujian/01/2002(H5N1)) neuraminidase (NA) mRNA, complete cds	1,357 bp linear mRNA	AY585399.1 GI:47156352
182. Influenza A virus (A/duck/Fujian/01/2002(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,497 bp linear mRNA	AY585420.1 GI:47156394
183. Influenza A virus (A/duck/Fujian/01/2002(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	686 bp linear mRNA	AY585441.1 GI:47156436
184. Influenza A virus (A/duck/Fujian/13/2002(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585505.1 GI:47156228
185. Influenza A virus (A/duck/Fujian/13/2002(H5N1)) matrix protein mRNA, complete cds	761 bp linear mRNA	AY585379.1 GI:47156312
186. Influenza A virus (A/duck/Fujian/13/2002(H5N1)) neuraminidase (NA) mRNA, complete cds	1,357 bp linear mRNA	AY585400.1 GI:47156354
187. Influenza A virus (A/duck/Fujian/13/2002(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,499 bp linear mRNA	AY585421.1 GI:47156396
188. Influenza A virus (A/duck/Fujian/13/2002(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	685 bp linear mRNA	AY585442.1 GI:47156438
189. Influenza A virus (A/duck/Fujian/17/2001(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585506.1 GI:47156230
190. Influenza A virus (A/duck/Fujian/17/2001(H5N1)) matrix protein mRNA, complete cds	759 bp linear mRNA	AY585380.1 GI:47156314
191. Influenza A virus (A/duck/Fujian/17/2001(H5N1)) neuraminidase (NA) mRNA, complete cds	1,418 bp linear mRNA	AY585401.1 GI:47156356
192. Influenza A virus (A/duck/Fujian/17/2001(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,498 bp linear mRNA	AY585422.1 GI:47156398
193. Influenza A virus (A/duck/Fujian/17/2001(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	686 bp linear mRNA	AY585443.1 GI:47156440
194. Influenza A virus (A/duck/Fujian/19/2000(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585507.1 GI:47156232

195. Influenza A virus (A/duck/Fujian/19/2000(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585381.1 GI:47156316
196. Influenza A virus (A/duck/Fujian/19/2000(H5N1)) neuraminidase (NA) mRNA, complete cds	1,355 bp linear mRNA	AY585402.1 GI:47156358
197. Influenza A virus (A/duck/Fujian/19/2000(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,498 bp linear mRNA	AY585423.1 GI:47156400
198. Influenza A virus (A/duck/Fujian/19/2000(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	687 bp linear mRNA	AY585444.1 GI:47156442
199. Influenza A virus (A/duck/Guangdong/01/2001(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585508.1 GI:47156234
200. Influenza A virus (A/duck/Guangdong/01/2001(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585382.1 GI:47156318
201. Influenza A virus (A/duck/Guangdong/01/2001(H5N1)) neuraminidase (NA) mRNA, complete cds	1,414 bp linear mRNA	AY585403.1 GI:47156360
202. Influenza A virus (A/duck/Guangdong/01/2001(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,497 bp linear mRNA	AY585424.1 GI:47156402
203. Influenza A virus (A/duck/Guangdong/01/2001(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	687 bp linear mRNA	AY585445.1 GI:47156444
204. Influenza A virus (A/duck/Guangdong/07/2000(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,280 bp linear mRNA	AY585509.1 GI:47156236
205. Influenza A virus (A/duck/Guangdong/07/2000(H5N1)) matrix protein mRNA, complete cds	759 bp linear mRNA	AY585383.1 GI:47156320
206. Influenza A virus (A/duck/Guangdong/07/2000(H5N1)) neuraminidase (NA) mRNA, complete cds	1,417 bp linear mRNA	AY585404.1 GI:47156362
207. Influenza A virus (A/duck/Guangdong/07/2000(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,497 bp linear mRNA	AY585425.1 GI:47156404
208. Influenza A virus (A/duck/Guangdong/07/2000(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	690 bp linear mRNA	AY585446.1 GI:47156446
209. Influenza A virus (A/duck/Guangdong/12/2000(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585510.1 GI:47156238
210. Influenza A virus (A/duck/Guangdong/12/2000(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585384.1 GI:47156322
211. Influenza A virus (A/duck/Guangdong/12/2000(H5N1)) neuraminidase (NA) mRNA, complete cds	1,359 bp linear mRNA	AY585405.1 GI:47156364
212. Influenza A virus (A/duck/Guangdong/12/2000(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,498 bp linear mRNA	AY585426.1 GI:47156406
213. Influenza A virus (A/duck/Guangdong/12/2000(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	685 bp linear mRNA	AY585447.1 GI:47156448
214. Influenza A virus (A/duck/Guangdong/22/2002(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585511.1 GI:47156240
215. Influenza A virus (A/duck/Guangdong/22/2002(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585385.1 GI:47156324
216. Influenza A virus (A/duck/Guangdong/22/2002(H5N1)) neuraminidase (NA) mRNA, complete cds	1,412 bp linear mRNA	AY585406.1 GI:47156366
217. Influenza A virus (A/duck/Guangdong/22/2002(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,499 bp linear mRNA	AY585427.1 GI:47156408
218. Influenza A virus (A/duck/Guangdong/22/2002(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	682 bp linear mRNA	AY585448.1 GI:47156450
219. Influenza A virus (A/duck/Guangdong/40/2000(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585512.1 GI:47156242
220. Influenza A virus (A/duck/Guangdong/40/2000(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585386.1 GI:47156326
221. Influenza A virus (A/duck/Guangdong/40/2000(H5N1)) neuraminidase (NA) mRNA, partial cds	1,401 bp linear mRNA	AY585407.1 GI:47156368
222. Influenza A virus (A/duck/Guangdong/40/2000(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,499 bp linear mRNA	AY585428.1 GI:47156410

223. Influenza A virus (A/duck/Guangdong/40/2000(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	689 bp linear mRNA	AY585449.1 GI:47156452
224. Influenza A virus (A/duck/Guangxi/07/1999(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585513.1 GI:47156244
225. Influenza A virus (A/duck/Guangxi/07/1999(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585387.1 GI:47156328
226. Influenza A virus (A/duck/Guangxi/07/1999(H5N1)) neuraminidase (NA) mRNA, complete cds	1,421 bp linear mRNA	AY585408.1 GI:47156370
227. Influenza A virus (A/duck/Guangxi/07/1999(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,501 bp linear mRNA	AY585429.1 GI:47156412
228. Influenza A virus (A/duck/Guangxi/07/1999(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	687 bp linear mRNA	AY585450.1 GI:47156454
229. Influenza A virus (A/duck/Guangxi/13/2004(H5N1)) nonstructural protein 1 mRNA, complete cds	875 bp linear mRNA	DQ366342.1 GI:86753723
230. Influenza A virus (A/duck/Guangxi/13/2004(H5N1)) polymerase PB2 mRNA, complete cds	2,341 bp linear mRNA	DQ366335.1 GI:86753733
231. Influenza A virus (A/duck/Guangxi/13/2004(H5N1)) polymerase PB1 mRNA, complete cds	2,341 bp linear mRNA	DQ366336.1 GI:86753743
232. Influenza A virus (A/duck/Guangxi/13/2004(H5N1)) PA protein mRNA, complete cds	2,233 bp linear mRNA	DQ366337.1 GI:86753753
233. Influenza A virus (A/duck/Guangxi/13/2004(H5N1)) hemagglutinin mRNA, complete cds	1,776 bp linear mRNA	DQ366338.1 GI:86753763
234. Influenza A virus (A/duck/Guangxi/13/2004(H5N1)) nucleocapsid mRNA, complete cds	1,565 bp linear mRNA	DQ366339.1 GI:86753773
235. Influenza A virus (A/duck/Guangxi/13/2004(H5N1)) neuraminidase mRNA, complete cds	1,378 bp linear mRNA	DQ366340.1 GI:86753783
236. Influenza A virus (A/duck/Guangxi/13/2004(H5N1)) matrix protein mRNA, complete cds	1,027 bp linear mRNA	DQ366341.1 GI:86753793
237. Influenza A virus (A/duck/Guangxi/22/2001(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585514.1 GI:47156246
238. Influenza A virus (A/duck/Guangxi/22/2001(H5N1)) matrix protein mRNA, partial cds	757 bp linear mRNA	AY585388.1 GI:47156330
239. Influenza A virus (A/duck/Guangxi/22/2001(H5N1)) neuraminidase (NA) mRNA, complete cds	1,414 bp linear mRNA	AY585409.1 GI:47156372
240. Influenza A virus (A/duck/Guangxi/22/2001(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,498 bp linear mRNA	AY585430.1 GI:47156414
241. Influenza A virus (A/duck/Guangxi/22/2001(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	687 bp linear mRNA	AY585451.1 GI:47156456
242. Influenza A virus (A/duck/Guangxi/35/2001(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585515.1 GI:47156248
243. Influenza A virus (A/duck/Guangxi/35/2001(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585389.1 GI:47156332
244. Influenza A virus (A/duck/Guangxi/35/2001(H5N1)) neuraminidase (NA) mRNA, complete cds	1,414 bp linear mRNA	AY585410.1 GI:47156374
245. Influenza A virus (A/duck/Guangxi/35/2001(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,498 bp linear mRNA	AY585431.1 GI:47156416
246. Influenza A virus (A/duck/Guangxi/35/2001(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	685 bp linear mRNA	AY585452.1 GI:47156458
247. Influenza A virus (A/duck/Guangxi/50/2001(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585516.1 GI:47156250
248. Influenza A virus (A/duck/Guangxi/50/2001(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585398.1 GI:47156350
249. Influenza A virus (A/duck/Guangxi/50/2001(H5N1)) neuraminidase (NA) mRNA, complete cds	1,354 bp linear mRNA	AY585411.1 GI:47156376
250. Influenza A virus (A/duck/Guangxi/50/2001(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,498 bp linear mRNA	AY585432.1 GI:47156418

251. Influenza A virus (A/duck/Guangxi/50/2001(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	686 bp linear mRNA	AY585453.1 GI:47156460
252. Influenza A virus (A/duck/Guangxi/53/2002(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585517.1 GI:47156252
253. Influenza A virus (A/duck/Guangxi/53/2002(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585390.1 GI:47156334
254. Influenza A virus (A/duck/Guangxi/53/2002(H5N1)) neuraminidase (NA) mRNA, complete cds	1,361 bp linear mRNA	AY585412.1 GI:47156378
255. Influenza A virus (A/duck/Guangxi/53/2002(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,498 bp linear mRNA	AY585433.1 GI:47156420
256. Influenza A virus (A/duck/Guangxi/53/2002(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	687 bp linear mRNA	AY585454.1 GI:47156462
257. Influenza A virus (A/duck/Kurgan/08/2005(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,754 bp linear mRNA	DQ449640.1 GI:90289674
258. Influenza A virus (A/duck/Kurgan/08/2005(H5N1)) matrix protein 1 (M) mRNA, complete cds	1,002 bp linear mRNA	DQ449641.1 GI:90289689
259. Influenza A virus (A/duck/Kurgan/08/2005(H5N1)) neuraminidase (NA) mRNA, complete cds	1,373 bp linear mRNA	DQ449642.1 GI:90289708
260. Influenza A virus (A/duck/Kurgan/08/2005(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,540 bp linear mRNA	DQ449643.1 GI:90289731
261. Influenza A virus (A/duck/Kurgan/08/2005(H5N1)) nonstructural protein (NS) mRNA, complete cds	850 bp linear mRNA	DQ449644.1 GI:90289739
262. Influenza A virus (A/duck/Kurgan/08/2005(H5N1)) polymerase acidic protein (PA) mRNA, complete cds	2,208 bp linear mRNA	DQ449645.1 GI:90289756
263. Influenza A virus (A/duck/Kurgan/08/2005(H5N1)) polymerase basic protein 1 (PB1) mRNA, complete cds	2,316 bp linear mRNA	DQ449646.1 GI:90289774
264. Influenza A virus (A/duck/Kurgan/08/2005(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,316 bp linear mRNA	DQ449647.1 GI:90289783
266. Influenza A virus (A/duck/Shanghai/08/2001(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585518.1 GI:47156254
267. Influenza A virus (A/duck/Shanghai/08/2001(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585391.1 GI:47156336
268. Influenza A virus (A/duck/Shanghai/08/2001(H5N1)) neuraminidase (NA) mRNA, complete cds	1,357 bp linear mRNA	AY585413.1 GI:47156380
269. Influenza A virus (A/duck/Shanghai/08/2001(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,498 bp linear mRNA	AY585434.1 GI:47156422
270. Influenza A virus (A/duck/Shanghai/08/2001(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	685 bp linear mRNA	AY585455.1 GI:47156464
271. Influenza A virus (A/duck/Shanghai/13/2001(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585519.1 GI:47156256
272. Influenza A virus (A/duck/Shanghai/13/2001(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585392.1 GI:47156338
273. Influenza A virus (A/duck/Shanghai/13/2001(H5N1)) neuraminidase (NA) mRNA, complete cds	1,417 bp linear mRNA	AY585414.1 GI:47156382
274. Influenza A virus (A/duck/Shanghai/13/2001(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,499 bp linear mRNA	AY585435.1 GI:47156424
275. Influenza A virus (A/duck/Shanghai/13/2001(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	685 bp linear mRNA	AY585456.1 GI:47156466
276. Influenza A virus (A/duck/Shanghai/35/2002(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585520.1 GI:47156258
277. Influenza A virus (A/duck/Shanghai/35/2002(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585393.1 GI:47156340
278. Influenza A virus (A/duck/Shanghai/35/2002(H5N1)) neuraminidase (NA) mRNA, complete cds	1,363 bp linear mRNA	AY585415.1 GI:47156384
279. Influenza A virus (A/duck/Shanghai/35/2002(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,498 bp linear mRNA	AY585436.1 GI:47156426

280. Influenza A virus (A/duck/Shanghai/35/2002(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	685 bp linear mRNA	AY585457.1 GI:47156468
281. Influenza A virus (A/duck/Shanghai/37/2002(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585521.1 GI:47156260
282. Influenza A virus (A/duck/Shanghai/37/2002(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585394.1 GI:47156342
283. Influenza A virus (A/duck/Shanghai/37/2002(H5N1)) neuraminidase (NA) mRNA, complete cds	1,361 bp linear mRNA	AY585416.1 GI:47156386
284. Influenza A virus (A/duck/Shanghai/37/2002(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,497 bp linear mRNA	AY585437.1 GI:47156428
285. Influenza A virus (A/duck/Shanghai/37/2002(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	685 bp linear mRNA	AY585458.1 GI:47156470
286. Influenza A virus (A/duck/Shanghai/38/2001(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,282 bp linear mRNA	AY585522.1 GI:47156262
287. Influenza A virus (A/duck/Shanghai/38/2001(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585395.1 GI:47156344
288. Influenza A virus (A/duck/Shanghai/38/2001(H5N1)) neuraminidase (NA) mRNA, complete cds	1,355 bp linear mRNA	AY585417.1 GI:47156388
289. Influenza A virus (A/duck/Shanghai/38/2001(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,499 bp linear mRNA	AY585438.1 GI:47156430
290. Influenza A virus (A/duck/Shanghai/38/2001(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	692 bp linear mRNA	AY585459.1 GI:47156472
291. Influenza A virus (A/duck/Sheyang/1/2005(H5N1)) nonstructural protein (NS) mRNA, complete cds	875 bp linear mRNA	DQ354059.1 GI:87128643
292. Influenza A virus (A/duck/Tuva/01/2006(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,748 bp linear mRNA	DQ851291.1 GI:112820195
293. Influenza A virus (A/duck/Tuva/01/2006(H5N1)) matrix protein 1 (M1) mRNA, complete cds	991 bp linear mRNA	DQ851292.1 GI:112820197
294. Influenza A virus (A/duck/Tuva/01/2006(H5N1)) neuraminidase (NA) mRNA, complete cds	1,364 bp linear mRNA	DQ851293.1 GI:112820199
295. Influenza A virus (A/duck/Tuva/01/2006(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,531 bp linear mRNA	DQ851294.1 GI:112820201
296. Influenza A virus (A/duck/Tuva/01/2006(H5N1)) nonstructural protein (NS) mRNA, complete cds	842 bp linear mRNA	DQ851295.1 GI:112820203
297. Influenza A virus (A/duck/Vietnam/1/2005(H5N1)) nonstructural protein 1 mRNA, complete cds	890 bp linear mRNA	DQ366310.1 GI:86753715
298. Influenza A virus (A/duck/Vietnam/1/2005(H5N1)) polymerase PB2 mRNA, complete cds	2,341 bp linear mRNA	DQ366303.1 GI:86753725
299. Influenza A virus (A/duck/Vietnam/1/2005(H5N1)) polymerase PB1 mRNA, complete cds	2,341 bp linear mRNA	DQ366304.1 GI:86753735
300. Influenza A virus (A/duck/Vietnam/1/2005(H5N1)) PA protein mRNA, complete cds	2,233 bp linear mRNA	DQ366305.1 GI:86753745
301. Influenza A virus (A/duck/Vietnam/1/2005(H5N1)) hemagglutinin mRNA, complete cds	1,779 bp linear mRNA	DQ366306.1 GI:86753755
302. Influenza A virus (A/duck/Vietnam/1/2005(H5N1)) nucleocapsid mRNA, complete cds	1,565 bp linear mRNA	DQ366307.1 GI:86753765
303. Influenza A virus (A/duck/Vietnam/1/2005(H5N1)) neuraminidase mRNA, complete cds	1,401 bp linear mRNA	DQ366308.1 GI:86753775
304. Influenza A virus (A/duck/Vietnam/1/2005(H5N1)) matrix protein mRNA, complete cds	1,027 bp linear mRNA	DQ366309.1 GI:86753785
305. Influenza A virus (A/duck/Vietnam/8/05(H5N1)) nonstructural protein 1 mRNA, complete cds	890 bp linear mRNA	DQ366326.1 GI:86753719
306. Influenza A virus (A/duck/Vietnam/8/05(H5N1)) polymerase PB2 mRNA, complete cds	2,341 bp linear mRNA	DQ366319.1 GI:86753729
307. Influenza A virus (A/duck/Vietnam/8/05(H5N1)) polymerase PB1 mRNA, complete cds	2,341 bp linear mRNA	DQ366320.1 GI:86753739

308. Influenza A virus (A/duck/Vietnam/8/05(H5N1)) PA protein mRNA, complete cds	2,233 bp linear mRNA	DQ366321.1 GI:86753749
309. Influenza A virus (A/duck/Vietnam/8/05(H5N1)) hemagglutinin mRNA, complete cds	1,779 bp linear mRNA	DQ366322.1 GI:86753759
310. Influenza A virus (A/duck/Vietnam/8/05(H5N1)) nucleocapsid mRNA, complete cds	1,565 bp linear mRNA	DQ366323.1 GI:86753769
311. Influenza A virus (A/duck/Vietnam/8/05(H5N1)) neuraminidase mRNA, complete cds	1,401 bp linear mRNA	DQ366324.1 GI:86753779
312. Influenza A virus (A/duck/Vietnam/8/05(H5N1)) matrix protein mRNA, complete cds	1,027 bp linear mRNA	DQ366325.1 GI:86753789
313. Influenza A virus (A/duck/Yangzhou/232/2004(H5N1)) nonfunctional nonstructural protein (NS) mRNA, complete sequence	876 bp linear mRNA	DQ354060.1 GI:87128645
314. Influenza A virus (A/duck/Zhejiang/11/2000(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585523.1 GI:47156264
315. Influenza A virus (A/duck/Zhejiang/11/2000(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585396.1 GI:47156346
316. Influenza A virus (A/duck/Zhejiang/11/2000(H5N1)) neuraminidase (NA) mRNA, complete cds	1,352 bp linear mRNA	AY585418.1 GI:47156390
317. Influenza A virus (A/duck/Zhejiang/11/2000(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,498 bp linear mRNA	AY585439.1 GI:47156432
318. Influenza A virus (A/duck/Zhejiang/11/2000(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	687 bp linear mRNA	AY585460.1 GI:47156474
319. Influenza A virus (A/duck/Zhejiang/52/2000(H5N1)) polymerase basic protein 2 (PB2) mRNA, complete cds	2,281 bp linear mRNA	AY585524.1 GI:47156266
320. Influenza A virus (A/duck/Zhejiang/52/2000(H5N1)) matrix protein mRNA, complete cds	760 bp linear mRNA	AY585397.1 GI:47156348
321. Influenza A virus (A/duck/Zhejiang/52/2000(H5N1)) neuraminidase (NA) mRNA, complete cds	1,423 bp linear mRNA	AY585419.1 GI:47156392
322. Influenza A virus (A/duck/Zhejiang/52/2000(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,499 bp linear mRNA	AY585440.1 GI:47156434
323. Influenza A virus (A/duck/Zhejiang/52/2000(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	686 bp linear mRNA	AY585461.1 GI:47156476
324. Influenza A virus (A/Egypt/0636-NAMRU3/2007(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,749 bp linear mRNA	EF382359.1 GI:124244205
325. Influenza A virus (A/goosander/Switzerland/V82/06 (H5N1)) hemagglutinin (HA) gene, complete cds	1,707 bp linear mRNA	EF110518.1 GI:119394674
326. Influenza A virus (A/goose/Guangdong/1/96/(H5N1)) hemagglutinin mRNA, complete cds	1,707 bp linear mRNA	AF148678.1 GI:5007022
327. Influenza A virus (A/Goose/Huadong/1/2000(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,779 bp linear mRNA	DQ201829.1 GI:76786306
328. Influenza A virus (A/Goose/Huadong/1/2000(H5N1)) neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	DQ201830.1 GI:76786308
329. Influenza A virus (A/goose/Hungary/2823/2/2007(H5N1)) polymerase PB1 (PB1) mRNA, partial cds	2,287 bp linear mRNA	EF446768.1 GI:126428373
330. Influenza A virus (A/goose/Hungary/2823/2/2007(H5N1)) polymerase PB2 (PB2) mRNA, partial cds	2,274 bp linear mRNA	EF446769.1 GI:126428375
331. Influenza A virus (A/goose/Hungary/2823/2/2007(H5N1)) polymerase PA (PA) mRNA, complete cds	2,175 bp linear mRNA	EF446770.1 GI:126428377
332. Influenza A virus (A/goose/Hungary/2823/2/2007(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,735 bp linear mRNA	EF446771.1 GI:126428379
333. Influenza A virus (A/goose/Hungary/2823/2/2007(H5N1)) nucleocapsid protein (NP) mRNA, partial cds	1,473 bp linear mRNA	EF446772.1 GI:126428381
334. Influenza A virus (A/goose/Hungary/2823/2/2007(H5N1)) neuraminidase (NA) mRNA, partial cds	1,311 bp linear mRNA	EF446773.1 GI:126428383
335. Influenza A virus (A/goose/Hungary/2823/2/2007(H5N1)) matrix protein 1 (M1) mRNA, partial cds	971 bp linear mRNA	EF446774.1 GI:126428385

336. Influenza A virus (A/goose/Hungary/2823/2/2007(H5N1)) nonstructural protein 1 (NS1) mRNA, partial cds	795 bp linear mRNA	EF446775.1 GI:126428387
337. Influenza A virus (A/goose/Hungary/3413/2007(H5N1)) polymerase PB1 (PB1) mRNA, partial cds	2,277 bp linear mRNA	EF446776.1 GI:126428389
338. Influenza A virus (A/goose/Hungary/3413/2007(H5N1)) polymerase PB2 (PB2) mRNA, partial cds	2,274 bp linear mRNA	EF446777.1 GI:126428391
339. Influenza A virus (A/goose/Hungary/3413/2007(H5N1)) polymerase PA (PA) mRNA, partial cds	2,163 bp linear mRNA	EF446778.1 GI:126428393
340. Influenza A virus (A/goose/Hungary/3413/2007(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,722 bp linear mRNA	EF446779.1 GI:126428395
341. Influenza A virus (A/goose/Hungary/3413/2007(H5N1)) nucleocapsid protein (NP) mRNA, partial cds	1,463 bp linear mRNA	EF446780.1 GI:126428397
342. Influenza A virus (A/goose/Hungary/3413/2007(H5N1)) neuraminidase (NA) mRNA, partial cds	1,289 bp linear mRNA	EF446781.1 GI:126428399
343. Influenza A virus (A/goose/Hungary/3413/2007(H5N1)) matrix protein 1 (M1) mRNA, partial cds	955 bp linear mRNA	EF446782.1 GI:126428401
344. Influenza A virus (A/goose/Hungary/3413/2007(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	805 bp linear mRNA	EF446783.1 GI:126428403
345. Influenza A virus (A/goose/jiangsu/131/2002(H5N1)) nonfunctional nonstructural protein (NS) mRNA, complete sequence	877 bp linear mRNA	DQ354061.1 GI:87128646
346. Influenza A virus (A/goose/Jiangsu/220/2003(H5N1)) nonstructural protein (NS) mRNA, complete cds	875 bp linear mRNA	DQ354062.1 GI:87128647
347. Influenza A virus (A/goose/Krasnoozerka/627/2005(H5N1)) hemagglutinin (HA) mRNA, complete cds	1,754 bp linear mRNA	DQ676840.1 GI:108782531
348. Influenza A virus (A/goose/Krasnoozerka/627/2005(H5N1)) nucleoprotein (NP) mRNA, complete cds	1,530 bp linear mRNA	DQ676841.1 GI:108782533
349. Influenza A virus (A/goose/Krasnoozerka/627/2005(H5N1)) nonstructural protein (NS) mRNA, complete cds	850 bp linear mRNA	DQ676842.1 GI:108782535
350. Influenza A virus (A/goose/Vietnam/3/05(H5N1)) nonstructural protein 1 mRNA, complete cds	890 bp linear mRNA	DQ366318.1 GI:86753717
351. Influenza A virus (A/goose/Vietnam/3/05(H5N1)) polymerase PB2 mRNA, complete cds	2,341 bp linear mRNA	DQ366311.1 GI:86753727
352. Influenza A virus (A/goose/Vietnam/3/05(H5N1)) polymerase PB1 mRNA, complete cds	2,341 bp linear mRNA	DQ366312.1 GI:86753737
353. Influenza A virus (A/goose/Vietnam/3/05(H5N1)) PA protein mRNA, complete cds	2,233 bp linear mRNA	DQ366313.1 GI:86753747
354. Influenza A virus (A/goose/Vietnam/3/05(H5N1)) hemagglutinin mRNA, complete cds	1,779 bp linear mRNA	DQ366314.1 GI:86753757
355. Influenza A virus (A/goose/Vietnam/3/05(H5N1)) nucleocapsid mRNA, complete cds	1,565 bp linear mRNA	DQ366315.1 GI:86753767
356. Influenza A virus (A/goose/Vietnam/3/05(H5N1)) neuraminidase mRNA, complete cds	1,401 bp linear mRNA	DQ366316.1 GI:86753777
357. Influenza A virus (A/goose/Vietnam/3/05(H5N1)) matrix protein mRNA, complete cds	1,027 bp linear mRNA	DQ366317.1 GI:86753787
358. Influenza A virus (A/gull/Pennsylvania/4175/83(H5N1)) hemagglutinin H5 mRNA, partial cds	1,700 bp linear mRNA	AF082043.1 GI:4240453
360. Influenza A virus (A/HongKong/156/97(H5N1)) neuraminidase mRNA, complete cds	1,388 bp linear mRNA	AF028708.1 GI:2865377
361. Influenza A virus (A/HongKong/156/97(H5N1)) hemagglutinin mRNA, complete cds	1,741 bp linear mRNA	AF028709.1 GI:2865379
362. Influenza A virus (A/HongKong/156/97(H5N1)) nucleoprotein mRNA, complete cds	1,549 bp linear mRNA	AF028710.1 GI:2865381
363. Influenza A virus (A/hooded vulture/Burkina Faso/1/2006(H5N1)) partial mRNA for nucleoprotein (np gene)	1,451 bp linear mRNA	AM503028.1 GI:147846292
364. Influenza A virus (A/hooded vulture/Burkina Faso/1/2006(H5N1)) mRNA for non-structural protein (ns gene)	827 bp linear mRNA	AM503038.1 GI:147846312

365. Influenza A virus (A/hooded vulture/Burkina Faso/1/2006(H5N1)) partial mRNA for polymerase (pa gene)	2,169 bp linear mRNA	AM503047.1 GI:147846330
366. Influenza A virus (A/hooded vulture/Burkina Faso/1/2006(H5N1)) partial mRNA for polymerase basic protein 1 (pb1 gene)	1,686 bp linear mRNA	AM503065.1 GI:147846855
367. Influenza A virus (A/hooded vulture/Burkina Faso/2/2006(H5N1)) partial mRNA for matrix protein 1 (m1 gene)	977 bp linear mRNA	AM503006.1 GI:147846248
368. Influenza A virus (A/hooded vulture/Burkina Faso/2/2006(H5N1)) partial mRNA for neuraminidase (na gene)	1,336 bp linear mRNA	AM503017.1 GI:147846270
369. Influenza A virus (A/hooded vulture/Burkina Faso/2/2006(H5N1)) partial mRNA for nucleoprotein (np gene)	1,499 bp linear mRNA	AM503027.1 GI:147846290
370. Influenza A virus (A/hooded vulture/Burkina Faso/2/2006(H5N1)) mRNA for non-structural protein (ns gene)	827 bp linear mRNA	AM503039.1 GI:147846314
371. Influenza A virus (A/hooded vulture/Burkina Faso/2/2006(H5N1)) partial mRNA for polymerase (pa gene)	2,169 bp linear mRNA	AM503048.1 GI:147846332
372. Influenza A virus (A/hooded vulture/Burkina Faso/2/2006(H5N1)) partial mRNA for polymerase basic protein 1 (pb1 gene)	2,259 bp linear mRNA	AM503062.1 GI:147846849
373. Influenza A virus (A/hooded vulture/Burkina Faso/2/2006(H5N1)) partial mRNA for polymerase basic protein 2 (pb2 gene)	2,315 bp linear mRNA	AM503066.1 GI:147846857
374. Influenza A virus (A/Indonesia/CDC177/2005(H5N1)) M2 protein mRNA, complete cds	294 bp linear mRNA	EU014135.1 GI:151336850
375. Influenza A virus (A/Indonesia/CDC298/2005(H5N1)) M2 protein mRNA, complete cds	294 bp linear mRNA	EU014138.1 GI:151336856
376. Influenza A virus (A/Indonesia/CDC485/2006(H5N1)) M2 protein mRNA, complete cds	294 bp linear mRNA	EU014136.1 GI:151336852
377. Influenza A virus (A/Indonesia/CDC530/2006(H5N1)) M2 protein mRNA, complete cds	294 bp linear mRNA	EU014134.1 GI:151336848
378. Influenza A virus (A/Indonesia/CDC535/2006(H5N1)) M2 protein mRNA, complete cds	294 bp linear mRNA	EU014133.1 GI:151336846
379. Influenza A virus (A/Indonesia/CDC540/2006(H5N1)) M2 protein mRNA, complete cds	294 bp linear mRNA	EU014132.1 GI:151336844
380. Influenza A virus (A/Indonesia/CDC561/2006(H5N1)) M2 protein mRNA, complete cds	294 bp linear mRNA	EU014137.1 GI:151336854
381. Influenza A virus (A/Indonesia/CDC60/2005(H5N1)) M2 protein mRNA, complete cds	294 bp linear mRNA	EU014139.1 GI:151336858
382. Influenza A virus (A/mallard/Wisconsin/428/75(H5N1)) hemagglutinin mRNA, partial cds	996 bp linear mRNA	U79453.1 GI:1840071
383. Influenza A virus (A/ostrich/VRLCU/Egypt/2011(H5N1)) segment 4 hemagglutinin (HA) mRNA, partial cds	441 bp linear mRNA	JN157759.1 GI:338223304
384. Influenza A virus (A/quail/yunnan/092/2002(H5N1)) nonstructural protein (NS) mRNA, complete cds	875 bp linear mRNA	DQ354063.1 GI:87128649
385. Influenza A virus (A/R(Turkey/Ontario/7732/66-Bellamy/42)(H5N1)) HA mRNA for hemagglutinin, partial cds	1,472 bp linear mRNA	AB241613.1 GI:82581222
386. Influenza A virus (A/Thailand/LFPN-2004/2004(H5N1)) neuraminidase mRNA, complete cds	1,350 bp linear mRNA	AY679513.1 GI:50843945
387. Influenza A virus (A/Thailand/LFPN-2004/2004(H5N1)) hemagglutinin mRNA, complete cds	1,704 bp linear mRNA	AY679514.1 GI:50843949
388. Influenza A virus (A/tiger/Thailand/CU-T4/04(H5N1)) polymerase basic protein 2 (PB2) mRNA, partial cds	534 bp linear mRNA	DQ017251.1 GI:65329524
389. Influenza A virus (A/tiger/Thailand/CU-T5/04(H5N1)) polymerase basic protein 2 (PB2) mRNA, partial cds	582 bp linear mRNA	DQ017252.1 GI:65329536
390. Influenza A virus (A/tiger/Thailand/CU-T6/04(H5N1)) polymerase basic protein 2 (PB2) mRNA, partial cds	564 bp linear mRNA	DQ017253.1 GI:65329553
391. Influenza A virus (A/tiger/Thailand/CU-T8/04(H5N1)) polymerase basic protein 2 (PB2) mRNA, partial cds	582 bp linear mRNA	DQ017254.1 GI:65329568
392. Influenza A virus (A/turkey/England/250/2007(H5N1)) hemagglutinin (HA) mRNA, partial cds	1,695 bp linear mRNA	EF441263.1 GI:129307104

393. Influenza A virus (A/turkey/England/250/2007(H5N1)) matrix protein (M) mRNA, partial cds	943 bp linear mRNA	EF441264.1 GI:129307106
394. Influenza A virus (A/turkey/England/250/2007(H5N1)) nonstructural protein 1 (NS1) mRNA, complete cds	812 bp linear mRNA	EF441265.1 GI:129307109
395. Influenza A virus (A/turkey/England/250/2007(H5N1)) polymerase PA (PA) mRNA, complete cds	2,185 bp linear mRNA	EF441266.1 GI:129307111
396. Influenza A virus (A/turkey/England/250/2007(H5N1)) polymerase PB2 (PB2) mRNA, partial cds	2,272 bp linear mRNA	EF441267.1 GI:129307113
397. Influenza A virus (A/turkey/England/250/2007(H5N1)) nucleocapsid (NP) mRNA, partial cds	1,396 bp linear mRNA	EF441268.1 GI:129307115
398. Influenza A virus (A/turkey/England/250/2007(H5N1)) polymerase PB1 (PB1) mRNA, partial cds	2,288 bp linear mRNA	EF441269.1 GI:129307117
399. Influenza A virus (A/turkey/England/250/2007(H5N1)) neuraminidase (NA) mRNA, partial cds	1,276 bp linear mRNA	EF441270.1 GI:129307119
A/chicken/Burkina Faso/13.1/2006(H5N1) neuraminidase (NA)		AM503016.1
A/chicken/Crimea/04/2005(H5N1) neuraminidase (NA)		DQ650661.1
A/chicken/Crimea/04/2005(H5N1) hemagglutinin		DQ650659.1
A/chicken/Crimea/08/2005(H5N1) polymerase basic protein 1 (PB1)		DQ650669.1
A/chicken/Crimea/08/2005(H5N1) neuraminidase (NA)		DQ650665.1
A/chicken/Crimea/08/2005(H5N1) hemagglutinin (HA)		DQ650663.1
A/chicken/Guangxi/12/2004(H5N1) nonstructural protein 1		DQ366334.1
A/chicken/Guangxi/12/2004(H5N1) neuraminidase		DQ366332.1
A/chicken/Guangxi/12/2004(H5N1) hemagglutinin		DQ366330.1
A/duck/Kurgan/08/2005(H5N1) nucleoprotein (NP)		DQ449643.1

Table 9. Other Influenza A Antigens (H1N*, H2N*, H3N*)

Strain/Protein	Length	GenBank / GI Accession Nos.
H1N*		
Influenza A virus (A/duck/Hong Kong/193/1977(H1N2)) nucleoprotein (NP) mRNA, partial cds	1,402 bp linear mRNA	U49097.1 GI:1912392
Influenza A virus (A/duck/Hong Kong/193/1977(H1N2)) polymerase (PB1) mRNA, partial cds	258 bp linear mRNA	U48285.1 GI:1912374
Influenza A virus (A/England/2/2002(H1N2)) partial NS1 gene for non structural protein 1 and partial NS2 gene for non structural protein 2, genomic RNA	795 bp linear mRNA	AJ519455.1 GI:31096426
Influenza A virus (A/England/3/02(H1N2)) partial mRNA for nucleoprotein (np gene)	384 bp linear mRNA	AJ489497.1 GI:27526856
Influenza A virus (A/England/3/02(H1N2)) partial mRNA for polymerase subunit 2 (pb2 gene)	442 bp linear mRNA	AJ489488.1 GI:27526838
Influenza A virus (A/England/5/02(H1N2)) partial mRNA for nucleoprotein (np gene)	384 bp linear mRNA	AJ489498.1 GI:27526858
Influenza A virus (A/England/5/02(H1N2)) partial mRNA for polymerase subunit 2 (pb2 gene)	442 bp linear mRNA	AJ489489.1 GI:27526840
Influenza A virus (A/England/57/02(H1N2)) partial mRNA for nucleoprotein (np gene)	384 bp linear mRNA	AJ489499.1 GI:27526860
Influenza A virus (A/England/57/02(H1N2)) partial mRNA for polymerase subunit 2 (pb2 gene)	442 bp linear mRNA	AJ489492.1 GI:27526846
Influenza A virus (A/England/691/01(H1N2)) partial mRNA for nucleoprotein (np gene)	384 bp linear mRNA	AJ489496.1 GI:27526854
Influenza A virus (A/England/73/02(H1N2)) partial mRNA for nucleoprotein (np gene)	384 bp linear mRNA	AJ489500.1 GI:27526862
Influenza A virus (A/England/73/02(H1N2)) partial mRNA for polymerase subunit 2 (pb2 gene)	442 bp linear mRNA	AJ489493.1 GI:27526848

Influenza A virus (A/England/90/02(H1N2)) partial mRNA for nucleoprotein (np gene)	384 bp linear mRNA	AJ489501.1 GI:27526864
Influenza A virus (A/England/90/02(H1N2)) partial mRNA for polymerase subunit 2 (pb2 gene)	442 bp linear mRNA	AJ489490.1 GI:27526842
Influenza A virus (A/England/97/02(H1N2)) partial mRNA for nucleoprotein (np gene)	384 bp linear mRNA	AJ489502.1 GI:27526866
Influenza A virus (A/England/97/02(H1N2)) partial mRNA for polymerase subunit 2 (pb2 gene)	442 bp linear mRNA	AJ489491.1 GI:27526844
Influenza A virus (A/England/627/01(H1N2)) partial mRNA for nucleoprotein (np gene)	384 bp linear mRNA	AJ489494.1 GI:27526850
Influenza A virus (A/England/627/01(H1N2)) partial mRNA for polymerase subunit 2 (pb2 gene)	442 bp linear mRNA	AJ489485.1 GI:27526832
Influenza A virus (A/England/691/01(H1N2)) partial mRNA for polymerase subunit 2 (pb2 gene)	442 bp linear mRNA	AJ489487.1 GI:27526836
Influenza A virus (A/Egypt/96/2002(H1N2)) partial NS1 gene for non structural protein 1 and partial NS2 gene for non structural protein 2, genomic RNA	747 bp linear mRNA	AJ519457.1 GI:31096432
Influenza A virus (A/Israel/6/2002(H1N2)) partial NS1 gene for non structural protein 1 and partial NS2 gene for non structural protein 2, genomic RNA	773 bp linear mRNA	AJ519456.1 GI:31096429
Influenza A virus (A/Saudi Arabia/2231/2001(H1N2)) partial NS1 gene for non structural protein 1 and partial NS2 gene for non structural protein 2, genomic RNA	772 bp linear mRNA	AJ519453.1 GI:31096420
Influenza A virus (A/Scotland/122/01(H1N2)) partial mRNA for nucleoprotein (np gene)	384 bp linear mRNA	AJ489495.1 GI:27526852
Influenza A virus (A/Scotland/122/01(H1N2)) partial mRNA for polymerase subunit 2 (pb2 gene)	442 bp linear mRNA	AJ489486.1 GI:27526834
Influenza A virus (A/swine/Bakum/1832/2000(H1N2)) hemagglutinin (HA) mRNA, partial cds	832 bp linear mRNA	AY861443.1 GI:57791765
Influenza A virus (A/swine/Bakum/1832/2000(H1N2)) neuraminidase mRNA, partial cds	467 bp linear mRNA	AY870645.1 GI:58042754
Influenza A virus (A/swine/Cotes d'Armor/0040/2007(H1N2)) segment 4 partial mRNA	1,039 bp linear mRNA	AM503547.1 GI:225578611
Influenza A virus (A/swine/Cotes d'Armor/0136_17/2006(H1N2)) partial mRNA for haemagglutinin precursor (HA1 gene)	1,136 bp linear mRNA	AM490224.3 GI:222062921
Influenza A virus (A/swine/England/72685/96(H1N2)) haemagglutinin precursor, mRNA, complete cds	1,778 bp linear mRNA	AF085417.1 GI:3831770
Influenza A virus (A/swine/England/17394/96(H1N2)) haemagglutinin precursor, mRNA, complete cds	1,778 bp linear mRNA	AF085416.1 GI:3831768
Influenza A virus (A/swine/England/690421/95(H1N2)) haemagglutinin precursor, mRNA, complete cds	1,778 bp linear mRNA	AF085415.1 GI:3831766
Influenza A virus (A/swine/England/438207/94(H1N2)) haemagglutinin precursor, mRNA, complete cds	1,778 bp linear mRNA	AF085414.1 GI:3831764
Influenza A virus (A/Swine/Korea/CY02/02(H1N2)) neuraminidase (NA) mRNA, complete cds	1,427 bp linear mRNA	AY129157.1 GI:24286064
Influenza A virus (A/Swine/Korea/CY02/02(H1N2)) matrix protein (M) mRNA, complete cds	952 bp linear mRNA	AY129158.1 GI:24286066
Influenza A virus (A/Swine/Korea/CY02/02(H1N2)) nucleoprotein (NP) mRNA, complete cds	1,542 bp linear mRNA	AY129159.1 GI:24286069
Influenza A virus (A/Swine/Korea/CY02/02(H1N2)) nonstructural protein (NS) mRNA, complete cds	842 bp linear mRNA	AY129160.1 GI:24286081
Influenza A virus (A/Swine/Korea/CY02/02(H1N2)) polymerase acidic protein 2 (PA) mRNA, complete cds	2,165 bp linear mRNA	AY129161.1 GI:24286087
Influenza A virus (A/Swine/Korea/CY02/02(H1N2)) polymerase subunit 1 (PB1) mRNA, complete cds	2,274 bp linear mRNA	AY129162.1 GI:24286096
Influenza A virus (A/Swine/Korea/CY02/02(H1N2)) polymerase subunit 2 (PB2) mRNA, complete cds	2,334 bp linear mRNA	AY129163.1 GI:24286100
Influenza A virus (A/swine/Scotland/410440/94(H1N2)) haemagglutinin precursor, mRNA, complete cds	1,778 bp linear mRNA	AF085413.1 GI:3831762

Influenza A virus (A/swine/Spain/80598-LP4/2007(H1N2)) matrix protein 2 (M2) mRNA, partial cds	291 bp linear mRNA	EU305436.1 GI:168830657
Influenza A virus (A/Switzerland/3100/2002(H1N2)) partial HA gene for Haemagglutinin, genomic RNA	975 bp linear mRNA	AJ517813.1 GI:38422519
Influenza A virus (A/duck/Hong Kong/717/1979(H1N3)) nucleoprotein (NP) mRNA, partial cds	1,387 bp linear mRNA	U49095.1 GI:1912388
Influenza A virus (A/duck/Hong Kong/717/1979(H1N3)) polymerase (PB1) mRNA, partial cds	265 bp linear mRNA	U48281.1 GI:1912366
Influenza A virus (A/herring gull/New Jersey/780/86 (H1N3)) nonfunctional matrix protein mRNA, partial sequence	971 bp linear mRNA	AY664422.1 GI:51011826
Influenza A virus (A/mallard/Alberta/42/77(H1N6)) nonfunctional matrix protein mRNA, partial sequence	997 bp linear mRNA	AY664426.1 GI:51011830
Influenza A virus (A/swine/England/191973/92(H1N7)) matrix protein M1 mRNA, complete cds	1,020 bp linear mRNA	U85985.1 GI:1835733
Influenza A virus (A/swine/England/191973/92(H1N7)) nucleoprotein mRNA, complete cds	1,524 bp linear mRNA	U85987.1 GI:1835737
Influenza A virus (A/swine/England/191973/92(H1N7)) neuraminidase mRNA, complete cds	1,458 bp linear mRNA	U85988.1 GI:1835739
Influenza A virus (A/swine/England/191973/92(H1N7)) haemagglutinin HA mRNA, partial cds	1,698 bp linear mRNA	U85986.1 GI:1835735
H2N*		
Influenza A virus (A/ruddy turnstone/Delaware/81/93 (H2N1)) nonfunctional matrix protein mRNA, partial sequence	917 bp linear mRNA	AY664465.1 GI:51011869
Influenza A virus (A/ruddy turnstone/Delaware/34/93 (H2N1)) nonfunctional matrix protein mRNA, partial sequence	968 bp linear mRNA	AY664429.1 GI:51011833
Influenza A virus (A/Shorebird/Delaware/122/97(H2N1)) nonfunctional matrix protein mRNA, partial sequence	925 bp linear mRNA	AY664466.1 GI:51011870
Influenza A virus (A/shorebird/Delaware/138/97 (H2N1)) nonfunctional matrix protein mRNA, partial sequence	958 bp linear mRNA	AY664454.1 GI:51011858
Influenza A virus (A/shorebird/Delaware/111/97 (H2N1)) nonfunctional matrix protein mRNA, partial sequence	958 bp linear mRNA	AY664457.1 GI:51011861
Influenza A virus (A/shorebird/Delaware/24/98 (H2N1)) nonfunctional matrix protein mRNA, partial sequence	979 bp linear mRNA	AY664442.1 GI:51011846
Influenza virus type A/Leningrad/134/17/57 (H2N2) PA RNA, complete cds	2,233 bp linear mRNA	M81579.1 GI:324935
Influenza A virus (STRAIN A/MALLARD/NEW YORK/6750/78) partial mRNA for PA protein	2,151 bp linear mRNA	AJ243994.1 GI:5918195
Influenza A virus (A/X-7(F1)/(H2N2)) neuraminidase mRNA, complete cds	1,467 bp linear mRNA	M11205.1 GI:323969
Influenza A virus (A/mallard/Alberta/77/77 (H2N3)) nonfunctional matrix protein mRNA, partial sequence	1,009 bp linear mRNA	AY664425.1 GI:51011829
Influenza A virus (A/mallard/Alberta/226/98(H2N3)) nonfunctional matrix protein mRNA, partial sequence	968 bp linear mRNA	AY664447.1 GI:51011851
Influenza A virus (A/sanderling/New Jersey/766/86 (H2N7)) nonfunctional matrix protein mRNA, partial sequence	846 bp linear mRNA	AY664477.1 GI:51011881
Influenza A virus (A/laughing gull/New Jersey/798/86 (H2N7)) nonfunctional matrix protein mRNA, partial sequence	907 bp linear mRNA	AY664471.1 GI:51011875
Influenza A virus (A/herring gull/Delaware/471/1986(H2N7)) nonfunctional matrix protein mRNA, partial sequence	960 bp linear mRNA	AY664440.1 GI:51011844
Influenza A virus (A/ruddy turnstone/Delaware/142/98 (H2N8)) nonfunctional matrix protein mRNA, partial sequence	1,011 bp linear mRNA	AY664423.1 GI:51011827
Influenza A virus (A/pintail/Alberta/293/77 (H2N9)) nonfunctional matrix protein mRNA, partial sequence	906 bp linear mRNA	AY664473.1 GI:51011877
Influenza A virus (A/blue-winged teal/Alberta/16/97 (H2N9)) nonfunctional matrix protein mRNA, partial sequence	961 bp linear mRNA	AY664449.1 GI:51011853
Influenza A virus (A/Laughing gull/New Jersey/75/85 (H2N9)) nonfunctional matrix protein mRNA, partial sequence	952 bp linear mRNA	AY664437.1 GI:51011841
Influenza A virus (A/mallard/Alberta/205/98 (H2N9)) nonfunctional matrix protein mRNA, partial sequence	959 bp linear mRNA	AY664450.1 GI:51011854

H3N*		
Influenza A virus (A/duck/Eastern China/267/2003(H3N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	EU429755.1 GI:167859475
Influenza A virus (A/duck/Eastern China/253/2003(H3N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	EU429754.1 GI:167859473
Influenza A virus (A/duck/Eastern China/252/2003(H3N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	EU429753.1 GI:167859471
Influenza A virus (A/duck/Eastern China/243/2003(H3N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	EU429752.1 GI:167859469
Influenza A virus (A/duck/Eastern China/262/2003(H3N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	EU429734.1 GI:167859433
Influenza A virus (A/duck/Eastern China/233/2003(H3N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,459 bp linear mRNA	EU429733.1 GI:167859431
Influenza A virus (A/duck/Eastern China/213/2003(H3N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	EU429723.1 GI:167859411
Influenza A virus (A/duck/Eastern China/341/2003(H3N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	EU429719.1 GI:167859403
Influenza A virus (A/duck/Eastern China/01/2002(H3N1)) segment 6 neuraminidase (NA) mRNA, complete cds	1,458 bp linear mRNA	EU429718.1 GI:167859401
Influenza A virus (A/mallard/Alberta/22/76 (H3N6)) nonfunctional matrix protein mRNA, partial sequence	1,013 bp linear mRNA	AY664434.1 GI:51011838
Influenza A virus (A/mallard/Alberta/199/99(H3N6)) nonfunctional matrix protein mRNA, partial sequence	970 bp linear mRNA	AY664443.1 GI:51011847
Influenza A virus (A/shorebird/Delaware/222/97 (H3N6)) nonfunctional matrix protein mRNA, partial sequence	922 bp linear mRNA	AY664461.1 GI:51011865
Influenza A virus (A/Duck/Hokkaido/8/80 (H3N8)) hemagglutinin precursor, mRNA, partial cds	984 bp linear mRNA	AF079570.1 GI:3414978
Influenza A virus (A/Duck/Hokkaido/8/80 (H3N8)) nucleoprotein mRNA, complete cds	1,497 bp linear mRNA	AF079571.1 GI:3414980
Influenza A virus (A/duck/Ukraine/1/1963(H3N8)) segment 6 neuraminidase (NA) mRNA, complete cds	1,461 bp linear mRNA	EU429797.1 GI:167859559
Influenza A virus (A/duck/Eastern China/19/2004(H3N8)) segment 6 neuraminidase (NA) mRNA, complete cds	1,460 bp linear mRNA	EU429698.1 GI:167859361
Influenza A virus (A/duck/Eastern China/90/2004(H3N8)) segment 6 neuraminidase (NA) mRNA, complete cds	1,460 bp linear mRNA	EU429700.1 GI:167859365
Influenza A virus (A/duck/Eastern China/18/2005(H3N8)) segment 6 neuraminidase (NA) mRNA, complete cds	1,460 bp linear mRNA	EU429787.1 GI:167859539
Influenza A virus (A/duck/Eastern China/119/2005(H3N8)) segment 6 neuraminidase (NA) mRNA, complete cds	1,460 bp linear mRNA	EU429788.1 GI:167859541
Influenza A virus (A/equine/Argentina/1/96(H3N8)) hemagglutinin precursor (HA1) mRNA, partial cds	1,061 bp linear mRNA	AF197246.1 GI:6651512
Influenza A virus (A/equine/Argentina/2/94(H3N8)) hemagglutinin precursor (HA1) mRNA, partial cds	1,061 bp linear mRNA	AF197245.1 GI:6651510
Influenza A virus (A/equine/Argentina/1/95(H3N8)) hemagglutinin precursor (HA1) mRNA, partial cds	1,061 bp linear mRNA	AF197244.1 GI:6651508
Influenza A virus HA partial gene for haemagglutinin, genomic RNA, strain A/equine/Berlin/3/89(H3N8)	1,026 bp linear mRNA	AJ223194.1 GI:2780201
Influenza A virus HA partial gene for haemagglutinin, genomic RNA, strain A/equine/Berlin/4/89(H3N8)	1,006 bp linear mRNA	AJ223195.1 GI:2780203
Influenza A virus (A/equine/Florida/1/94(H3N8)) hemagglutinin precursor (HA1) mRNA, partial cds	1,061 bp linear mRNA	AF197242.1 GI:6651504
Influenza A virus (A/equine/Grobois/1/98(H3N8)) nonstructural protein NS1 mRNA, complete cds	695 bp linear mRNA	AY328471.1 GI:32966577
Influenza A virus (A/equi 2/Gotland/01(H3N8)) hemagglutinin HA1 subunit mRNA, partial cds	473 bp linear mRNA	AY919314.1 GI:60250543
Influenza A virus (A/eq/Kentucky/81(H3N8)) hemagglutinin mRNA, complete cds	1,763 bp linear mRNA	U58195.1 GI:1377873
Influenza A virus (A/equine/Kentucky/9/95(H3N8)) hemagglutinin precursor (HA1) mRNA, partial cds	1,061 bp linear mRNA	AF197247.1 GI:6651514

Influenza A virus (A/equine/Kentucky/1/96(H3N8)) hemagglutinin precursor (HA1) mRNA, partial cds	1,061 bp linear mRNA	AF197248.1 GI:6651516
Influenza A virus (A/equine/Kentucky/1/97(H3N8)) hemagglutinin precursor (HA1) mRNA, partial cds	1,061 bp linear mRNA	AF197249.1 GI:6651518
Influenza A virus (A/equine/Kentucky/1/98(H3N8)) hemagglutinin precursor (HA1) mRNA, partial cds	1,061 bp linear mRNA	AF197241.1 GI:6651502
Influenza A virus (A/equine/Santiago/85(H3N8)) nucleoprotein mRNA, complete cds	1,497 bp linear mRNA	AY383753.1 GI:37223511
Influenza A virus (A/equine/Santiago/85(H3N8)) hemagglutinin mRNA, complete cds	1,698 bp linear mRNA	AY383755.1 GI:37223515
Influenza A virus (A/equine/Santiago/85(H3N8)) neuraminidase mRNA, complete cds	1,413 bp linear mRNA	AY383754.1 GI:37223513
Influenza A virus (A/equine/Saskatoon/1/90(H3N8)) hemagglutinin precursor (HA1) mRNA, partial cds	1,061 bp linear mRNA	AF197243.1 GI:6651506
Influenza A virus (A/mallard/Alberta/I14/97 (H3N8)) nonfunctional matrix protein mRNA, partial sequence	1,010 bp linear mRNA	AY664432.1 GI:51011836
Influenza A virus (A/mallard/Alberta/167/98 (H3N8)) nonfunctional matrix protein mRNA, partial sequence	961 bp linear mRNA	AY664489.1 GI:51011893
Influenza A virus (A/pintail/Alberta/37/99(H3N8)) nonfunctional matrix protein mRNA, partial sequence	970 bp linear mRNA	AY664445.1 GI:51011849
Influenza A virus (A/sanderling/Delaware/65/99 (H3N8)) nonfunctional matrix protein mRNA, partial sequence	922 bp linear mRNA	AY664455.1 GI:51011859

Table 10. Other Influenza A Antigens (H4N*-H13N*)

SEQ ID NO	Strain/Protein	GenBank Access No.
H4N*		
1215	A/chicken/Singapore/1992(H4N1) M2 protein	EU014144.1
1216	A/mallard/Alberta/47/98(H4N1) nonfunctional matrix protein	AY664488.1
1217	A/duck/Hong Kong/412/1978(H4N2) polymerase (PB1)	U48279.1
1218	A/mallard/Alberta/300/77 (H4N3) nonfunctional matrix protein	AY664480.1
1219	A/Duck/Czechoslovakia/56(H4N6) segment 4 hemagglutinin	AF290436.1
1220	A/duck/Eastern China/376/2004(H4N6) segment 6neuraminidase (NA)	EU429792.1
1221	A/duck/Eastern China/01/2007(H4N6) segment 6 neuraminidase (NA)	EU429790.1
1222	A/duck/Eastern China/216/2007(H4N6) segment 6 neuraminidase (NA)	EU429789.1
1223	A/duck/Eastern China/166/2004(H4N6) segment 6 neuraminidase (NA)	EU429746.1
1224	A/duck/Eastern China/02/2003(H4N6) segment 6 neuraminidase (NA)	EU429713.1
1225	A/duck/Eastern China/160/2002(H4N6) segment 6 neuraminidase (NA)	EU429706.1
1226	A/mallard/Alberta/111/99(H4N6) nonfunctional matrix protein	AY664482.1
1227	A/mallard/Alberta/213/99 (H4N6) nonfunctional matrix protein	AY664460.1
1228	A/mallard/Alberta/30/98 (H4N6) nonfunctional matrix protein	AY664484.1
1229	A/blue-winged teal/Alberta/96/76 (H4N8) nonfunctional matrix protein	AY664420.1
H5N*		
1230	A/chicken/Florida/25717/1993(H5N2) hemagglutinin	U05332.1
1231	A/chicken/Hidalgo/26654-1368/1994(H5N2) hemagglutinin (HA)	U37172.1
1232	A/chicken/Jalisco/14585-660/1994(H5N2) hemagglutinin (HA)	U37181.1
1233	A/chicken/Mexico/26654-1374/1994(H5N2) hemagglutinin (HA)	U37173.1

1234	A/chicken/Mexico/31381-3/1994(H5N2) hemagglutinin (HA)	U37176.1
1235	A/chicken/Mexico/31381-6/1994(H5N2) hemagglutinin (HA)	U37175.1
1236	A/chicken/Mexico/31381-4/1994(H5N2) hemagglutinin (HA)	U37174.1
1237	A/chicken/Mexico/31381-5/1994(H5N2) hemagglutinin (HA)	U37169.1
1238	A/chicken/Mexico/31381-8/1994(H5N2) hemagglutinin (HA)	U37170.1
1239	A/Chicken/Mexico/31381-Avilab/94(H5N2)hemagglutinin (HA)	L46585.1
1240	A/chicken/Mexico/31382-1/1994(H5N2)hemagglutinin (HA)	U37168.1
1241	A/chicken/Mexico/31381-2/1994(H5N2) hemagglutinin (HA)	U37167.1
1242	A/chicken/Mexico/31381-1/1994(H5N2) hemagglutinin (HA)	U37166.1
1243	A/chicken/Mexico/31381-7/1994(H5N2) hemagglutinin (HA)	U37165.1
1244	A/chicken/Pennsylvania/13609/1993(H5N2) hemagglutinin	U05331.1
1245	A/chicken/Pennsylvania/1/1983(H5N2) hemagglutinin esterase precursor	M18001.1
1246	A/chicken/Pennsylvania/1370/1983(H5N2) hemagglutinin esterase precursor	M10243.1
1247	A/Chicken/Puebla/8623-607/94(H5N2) hemagglutinin (HA)	L46586.1
1248	A/chicken/Puebla/14586-654/1994(H5N2) hemagglutinin (HA)	U37180.1
1249	A/chicken/Puebla/14585-622/1994(H5N2) hemagglutinin (HA)	U37179.1
1250	A/chicken/Puebla/8623-607/1994(H5N2)hemagglutinin (HA)	U37178.1
1251	A/chicken/Puebla/8624-604/1994(H5N2) hemagglutinin (HA)	U37177.1
1252	A/Chicken/Queretaro/14588-19/95(H5N2) hemagglutinin (HA)	L46587.1
1253	A/chicken/Queretaro/7653-20/95(H5N2) hemagglutinin (HA)	U79448.1
1254	A/chicken/Queretaro/26654-1373/1994(H5N2) hemagglutinin (HA)	U37171.1
1255	A/chicken/Queretaro/14588-19/1994(H5N2)hemagglutinin (HA)	U37182.1
1256	A/chicken/Singapore/98(H5N2) matrix protein 2 (M2)	EF682127.1
1257	A/chicken/Taiwan/1209/03(H5N2) hemagglutinin protein (HA)	AY573917.1
1258	A/chicken/Taiwan/1209/03(H5N2) neuraminidase	AY573918.1
1259	A/duck/Eastern China/64/2004(H5N2) segment 6 neuraminidase (NA)	EU429791.1
1260	A/duck/Eastern China/264/2002(H5N2) segment 6 neuraminidase (NA)	EU429744.1
1261	A/duck/Eastern China/01/2001(H5N2) segment 6 neuraminidase (NA)	EU429728.1
1262	A/duck/Eastern China/06/2000(H5N2) segment 6 neuraminidase (NA)	EU429722.1
1263	A/duck/Hong Kong/342/78(H5N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107452.1
1264	A/duck/Hong Kong/342/78(H5N2) hemagglutinin precursor	U20475.1
1265	A/duck/Michigan/80(H5N2) hemagglutinin 1 chain	U20474.1
1266	A/duck/Michigan/80(H5N2) hemagglutinin	U79449.1
1267	A/duck/MN/1564/81(H5N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107467.1
1268	A/duck/Mongolia/54/2001(H5N2) hemagglutinin (HA)	AB241614.2
1269	A/duck/Primorie/2621/01(H5N2) hemagglutinin (HA)	AJ621811.3
1270	A/duck/Primorie/2621/01(H5N2)nucleoprotein (NP)	AJ621812.1
1271	A/duck/Primorie/2621/01(H5N2) nonstructural protein (NS)	AJ621813.1
1272	A/duck/Pennsylvania/84(H5N2) hemagglutinin 1chain	U20473.1
1273	A/duck/Potsdam/1402-6/86(H5N2) hemagglutinin H5	AF082042.1
1274	A/emu/Texas/39442/93(H5N2) hemaglutinin	U28920.1
1275	A/emu/Texas/39442/93(H5N2) hemaglutinin	U28919.1
1276	A/mallard/Alberta/645/80(H5N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107471.1
1277	A/mallard/AR/1C/2001(H5N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107463.1

1278	A/mallard/NY/189/82(H5N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107465.1
1279	A/mallard/MN/25/80(H5N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107473.1
1280	A/mallard/MI/18/80(H5N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107470.1
1281	A/mallard/Ohio/345/88(H5N2) hemagglutinin	U79450.1
1282	A/parrot/CA/6032/04(H5N2) polymerase basic protein 2 (PB2)	DQ256390.1
1283	A/parrot/CA/6032/04(H5N2) polymerase basic protein 1 (PB1)	DQ256389.1
1284	A/parrot/CA/6032/04(H5N2) matrix protein (M)	DQ256384.2
1285	A/parrot/CA/6032/04(H5N2) hemagglutinin (HA)	DQ256383.1
1286	A/parrot/CA/6032/04(H5N2) neuraminidase (NA)	DQ256385.1
1287	A/parrot/CA/6032/04(H5N2) polymerase basic protein 2 (PB2)	DQ256390.1
1288	A/parrot/CA/6032/04(H5N2) nucleoprotein (NP)	DQ256386.1
1289	A/parrot/CA/6032/04(H5N2)) polymerase (PA)	DQ256388.1
1290	A/ruddy turnstone/Delaware/244/91 (H5N2) nonfunctional matrix protein	AY664474.1
1291	A/ruddy turnstone/Delaware/244/91 (H5N2)	U05330.1
1292	A/turkey/Colorado/72(H5N2) hemagglutinin 1 chain (HA)	U20472.1
1293	A/turkey/England/N28/73 (H5N2) hemagglutinin	AY500365.1
1294	A/turkey/TX/14082/81(H5N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107464.1
1295	A/turkey/MN/1704/82(H5N2)) matrix protein 1 (M) and matrix protein 2 (M)	DQ107472.1
1296	A/turkey/Minnesota/10734/95(H5N2)) hemagglutinin	U79455.1
1297	A/turkey/Minnesota/3689-1551/81(H5N2) hemagglutinin	U79454.1
1298	A/chicken/Singapore/1997(H5N3) M2 protein	EU014141.1
1299	A/duck/Hokkaido/299/04(H5N3) hemagglutinin (HA)	AB241626.1
1300	A/duck/Hokkaido/193/04(H5N3) hemagglutinin (HA)	AB241625.1
1301	A/duck/Hokkaido/101/04(H5N3) hemagglutinin (HA)	AB241624.1
1302	A/duck/Hokkaido/447/00(H5N3) hemagglutinin (HA)	AB241620.1
1303	A/duck/Hokkaido/69/00(H5N3) hemagglutinin (HA)	AB241619.1
1304	A/duck/Hong Kong/205/77(H5N3) hemagglutinin H5	AF082038.1
1305	A/duck/Hong Kong/698/79(H5N3) hemagglutinin H5	AF082039.1
1306	A/duck/Hong Kong/308/78(H5N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107457.1
1307	A/duck/Hong Kong/825/80(H5N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107455.1
1308	A/duck/Hong Kong/820/80(H5N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107453.1
1309	A/duck/Hong Kong/205/77(H5N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107456.1
1310	A/Duck/Ho Chi Minh/014/78(H5N3) segment 4 hemagglutinin	AF290443.1
1311	A/duck/Jiangxi/6151/2003(H5N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107451.1
1312	A/duck/Malaysia/F119-3/97(H5N3) hemagglutinin	AF303057.1
1313	A/duck/Miyagi/54/76(H5N3)hemagglutinin (HA)	AB241615.1
1314	A/duck/Mongolia/596/01(H5N3) hemagglutinin HA)	AB241622.1
1315	A/duck/Mongolia/500/01(H5N3)hemagglutinin (HA)	AB241621.1
1316	A/duck/Primorie/2633/01(H5N3) matrix protein (M1)	AJ621810.1
1317	A/duck/Primorie/2633/01(H5N3)nucleoprotein (NP)	AJ621808.1
1318	A/duck/Primorie/2633/01(H5N3)hemagglutinin (HA)	AJ621807.1
1319	A/duck/Primorie/2633/01(H5N3)nucleoprotein (NP)	AJ621809.1
1320	A/goose/Hong Kong/23/78(H5N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107454.1
1321	A/mallard/Wisconsin/169/75(H5N3) hemagglutinin	U79452.1
1322	A/swan/Hokkaido/51/96(H5N3)hemagglutinin (HA)	AB241617.1

1323	A/swan/Hokkaido/4/96(H5N3) hemagglutinin (HA)	AB241616.1
1324	A/turkey/CA/6878/79(H5N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107459.1
1325	A/tern/South Africa/61(H5N3) hemagglutinin precursor (HA)	U20460.1
1326	A/gull/Delaware/5/2000(H5N4) matrix protein 1 (M) and matrix protein 2 (M)	DQ107459.1
1327	A/gull/Delaware/4/2000(H5N4) matrix protein 1 (M) and matrix protein 2 (M)	DQ107458.1
1328	A/shorebird/Delaware/109/2000(H5N4) matrix protein 1 (M)	DQ107460.1
1329	A/shorebird/Delaware/243/2000(H5N4) matrix protein 1 (M) and matrix protein 2 (M)	DQ107462.1
1330	A/shorebird/Delaware/230/2000(H5N4) matrix protein 1 (M) and matrix protein 2 (M)	DQ107461.1
1331	A/mallard/Wisconsin/34/75(H5N6) hemagglutinin	U79451.1
1332	A/duck/Potsdam/2216-4/1984(H5N6) hemagglutinin H5	AF082041.1
1333	A/shorebird/Delaware/207/98 (H5N8) nonfunctional matrix protein	AY664456.1
1334	A/shorebird/Delaware/27/98 (H5N8) nonfunctional matrix protein	AY664453.1
1335	A/herring gull/Delaware/281/98 (H5N8) nonfunctional matrix protein	AY664452.1
1336	A/mallard/Ohio/556/1987(H5N9) hemagglutinin (HA)	U67783.2
1337	A/turkey/Wisconsin/68(H5N9) hemagglutinin	U79456.1
H6N*		
1338	A/blue-winged teal/Alberta/685/82(H6N1) matrix protein 1 (M) and matrix protein 2 (M)	DQ107448.1
1339	A/chicken/Taiwan/7-5/99(H6N1) nucleocapsid protein (NP)	AF261750.1
1340	A/chicken/Taiwan/7-5/99(H6N1) matrix protein	AF262213.1
1341	A/chicken/Taiwan/7-5/99(H6N1) nonstructural protein	AF262212.1
1342	A/chicken/Taiwan/7-5/99(H6N1) polymerase (PA)	AF262211.1
1343	A/chicken/Taiwan/7-5/99(H6N1) polymerase subunit PB1	AF262210.1
1344	A/chicken/Taiwan/7-5/99(H6N1) nucleocapsid protein (NP)	AF261750.1
1345	A/chicken/Taiwan/ns2/99(H6N1) segment 4 hemagglutinin (HA1)	AF310985.1
1346	A/chicken/Taiwan/na3/98(H6N1) segment 4 hemagglutinin (HA1)	AF310984.1
1347	A/chicken/Taiwan/7-5/99(H6N1) segment 4 hemagglutinin (HA1)	AF310983.1
1348	A/duck/Hong Kong/D73/76(H6N1) matrix protein 1 (M) and matrix protein 2 (M)	DQ107432.1
1349	A/duck/Taiwan/9/23-3/2000(H6N1) matrix protein 1 (M) and matrix protein 2 (M)	DQ107407.1
1350	A/pheasant/Hong Kong/FY479/2000(H6N1) matrix protein 1 (M) and matrix protein 2 (M)	DQ107409.1
1351	A/pheasant/Hong Kong/SSP44/2002(H6N1) matrix protein 1 (M) and matrix protein 2 (M)	DQ107412.1
1352	A/quail/Hong Kong/YU421/2002(H6N1) matrix protein 1 (M) and matrix protein 2 (M)	DQ107414.1
1353	A/avian/NY/17150-7/2000(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107423.1
1354	A/chicken/CA/285/2003(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107429.1
1355	A/chicken/CA/375TR/2002(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107428.1
1356	A/chicken/CA/203/2003(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107426.1
1357	A/chicken/NY/101250-7/2001(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107419.1
1358	A/chicken/CA/625/2002(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107418.1
1359	A/Chicken/California/0139/2001(H6N2)nucleoprotein (NP)	AF474070.1
1360	A/Chicken/California/650/2000(H6N2) nucleoprotein (NP)	AF474069.1
1361	A/Chicken/California/9420/2001(H6N2) neuraminidase N2 (N2)	AF474048.1
1362	A/Chicken/California/9174/2001(H6N2) neuraminidase N2 (N2)	AF474047.1
1363	A/Chicken/California/8892/2001(H6N2)neuraminidase N2 (N2)	AF474046.1
1364	A/Chicken/California/6643/2001(H6N2) neuraminidase N2 (N2)	AF474045.1
1365	A/Chicken/California/1316/2001(H6N2)neuraminidase N2 (N2)	AF474044.1

1366	A/Chicken/California/0139/2001(H6N2) neuraminidase N2 (N2)	AF474043.1
1367	A/Chicken/California/1002/2000(H6N2) neuraminidase N2 (N2)	AF474042.1
1368	A/Chicken/California/650/2000(H6N2) neuraminidase N2 (N2)	AF474041.1
1369	A/Chicken/California/465/2000(H6N2) neuraminidase N2 (N2)	AF474040.1
1370	A/Chicken/California/431/2000(H6N2) neuraminidase N2 (N2)	AF474039.1
1371	A/Chicken/California/6643/2001(H6N2) hemagglutinin H6 (H6)	AF474035.1
1372	A/Chicken/California/431/2000(H6N2) hemagglutinin H6 (H6)	AF474029.1
1373	A/Chicken/California/9420/2001(H6N2) hemagglutinin H6 (H6)	AF474038.1
1374	A/Chicken/California/9174/2001(H6N2) hemagglutinin H6 (H6)	AF474037.1
1375	A/Chicken/California/8892/2001(H6N2) hemagglutinin H6 (H6)	AF474036.1
1376	A/Chicken/California/1316/2001(H6N2) hemagglutinin H6 (H6)	AF474034.1
1377	A/Chicken/California/0139/2001(H6N2) hemagglutinin H6 (H6)	AF474033.1
1378	A/Chicken/California/1002/2000(H6N2) hemagglutinin H6 (H6)	AF474032.1
1379	A/Chicken/California/650/2000(H6N2) hemagglutinin H6 (H6)	AF474031.1
1380	A/Chicken/California/465/2000(H6N2) hemagglutinin H6 (H6)	AF474030.1
1381	A/cornish cross/CA/139/2001(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107424.1
1382	A/duck/Eastern China/164/2002(H6N2) segment 6 neuraminidase (NA)	EU429762.1
1383	A/duck/Eastern China/729/2003(H6N2) segment 6 neuraminidase (NA)	EU429760.1
1384	A/duck/Eastern China/262/2002(H6N2) segment 6 neuraminidase (NA)	EU429743.1
1385	A/duck/Eastern China/74/2006(H6N2) segment 6 neuraminidase (NA)	EU429741.1
1386	A/duck/Eastern China/161/2002(H6N2) segment 6 neuraminidase (NA)	EU429740.1
1387	A/duck/Hong Kong/960/80(H6N2)) matrix protein 1 (M) and matrix protein 2 (M)	DQ107435.1
1388	A/duck/Hong Kong/D134/77(H6N2)) matrix protein 1 (M) and matrix protein 2 (M)	DQ107433.1
1389	A/duck/CA/10221/2002(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107421.1
1390	A/duck/Shantou/5540/2001(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107431.1
1391	A/guinea fowl/Hong Kong/SSP99/2002(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107413.1
1392	A/mallard/NY/016/83(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107449.1
1393	A/mallard/NY/046/83(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107450.1
1394	A/pintail/Alberta/644/81(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107445.1
1395	A/quail/Hong Kong/SF792/2000(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107410.1
1396	A/ruddy turnstone/Delaware/106/98 (H6N2) nonfunctional matrix protein	AY664439.1
1397	A/Shorebird/Delaware/127/97(H6N2) nonfunctional matrix protein	AY664467.1
1398	A/shorebird/Delaware/124/2001(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107417.1
1399	A/shorebird/Delaware/208/2001(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107427.1
1400	A/turkey/CA/527/2002(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107420.1
1401	A/turkey/CA/1623CT/2002(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107425.1
1402	A/turkey/MN/836/80(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107440.1
1403	A/turkey/MN/735/79(H6N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107437.1
1404	A/chicken/Hong Kong/17/77(H6N4)) matrix protein 1 (M) and matrix protein 2 (M)	DQ107436.1
1405	A/chicken/Hong Kong/CSW106/2001(H6N4) matrix protein 1 (M) and matrix protein 2 (M)	DQ107406.1
1406	A/gull/Delaware/18/2000(H6N4) matrix protein 1 (M) and matrix protein 2 (M)	DQ107415.1
1407	A/pheasant/Hong Kong/CSW2573/2001(H6N4) matrix protein 1 (M) and matrix protein 2 (M)	DQ107411.1
1408	A/quail/Hong Kong/CSW106/2001(H6N4) matrix protein 1 (M) and matrix protein 2 (M)	DQ107430.1
1409	A/Shorebird/Delaware/194/98(H6N4) nonfunctional matrix protein	AY664424.1
1410	A/shorebird/Delaware/259/2000(H6N4) matrix protein 1 (M) and matrix protein 2 (M)	DQ107416.1

1411	A/shearwater/Australia/1/1972(H6N5) segment 6 neuraminidase (NA)	EU429794.1
1412	A/shearwater/Australia/1/1972(H6N5) polymerase A (PA)	L25832.1
1413	A/pintail/Alberta/1040/79(H6N5) matrix protein 1 (M) and matrix protein 2 (M)	DQ107439.1
1414	A/blue-winged teal/MN/993/80(H6N6)) matrix protein 1 (M) and matrix protein 2 (M)	DQ107441.1
1415	A/duck/NY/83779/2002(H6N6) matrix protein 1 (M) and matrix protein 2 (M)	DQ107422.1
1416	A/duck/MN/1414/81(H6N6) matrix protein 1 (M) and matrix protein 2 (M)	DQ107444.1
1417	A/mallard/Alberta/289/82(H6N6) matrix protein 1 (M) and matrix protein 2 (M)	DQ107447.1
1418	A/mallard duck/MN/1041/80(H6N6) matrix protein 1 (M) and matrix protein 2 (M)	DQ107442.1
1419	A/pintail/Alberta/189/82(H6N6) matrix protein 1 (M) and matrix protein 2 (M)	DQ107446.1
1420	A/sanderling/Delaware/1258/86(H6N6) nonfunctional matrix protein	AY664436.1
1421	A/blue-winged teal/Alberta/368/78(H6N8)) matrix protein 1 (M) and matrix protein 2 (M)	DQ107438.1
1422	A/ruddy turnstone/Delaware/105/98 (H6N8) nonfunctional matrix protein	AY664428.1
1423	A/domestic duck/NY/81(H6N8)) matrix protein (M)	DQ107443.1
1424	A/duck/Eastern China/163/2002(H6N8) segment 6 neuraminidase (NA)	EU429786.1
1425	A/duck/Hong Kong/D182/77(H6N9) matrix protein 1 (M) and matrix protein 2 (M)	DQ107434.1
1426	A/chicken/Hong Kong/SF3/2001(H6) matrix protein 1 (M) and matrix protein 2 (M)	DQ107408.1
H7N*		
1427	A/African starling/England/983/79(H7N1) neuraminidase (N1)	AJ416629.1
1428	A/Afri.Star./Eng-Q/938/79(H7N1) hemagglutinin precurosr	AF149295.1
1429	A/chicken/Italy/1067/99(H7N1) matrix protein 1 (M1)	AJ416630.1
1430	A/chicken/Italy/1067/99(H7N1) neuraminidase (N1)	AJ416627.1
1431	A/chicken/Italy/4575/99 (H7N1) hemagglutinin (HA)	AJ493469.1
1432	A/chicken/Italy/13474/99(H7N1) haemagglutinin (HA)	AJ491720.1
1433	A/chicken/Italy/445/1999(H7N1)	AX537385.1
1434	A/Chicken/Italy/267/00(H7N1) hemagglutinin (HA)	AJ493215.1
1435	A/Chicken/Italy/13489/99(H7N1) hemagglutinin (HA)	AJ493214.1
1436	A/Chicken/Italy/13307/99(H7N1) hemagglutinin (HA)	AJ493212.1
1437	A/chicken/Singapore/1994(H7N1) M2 protein	EU014140.1
1438	A/duck/Hong Kong/301/78(H7N1) matrix protein 1 (M) and matrix protein 2 (M)	DQ107475.1
1439	A/Hong Kong/301/78(H7N1) hemagglutinin (HA)	AY672090.1
1440	A/fowl plaguq virus/Rostock/34 (H7N1) NP protein	AJ243993.1
1441	A/fowl plaguq virus/Rostock/34 (H7N1) PA protein	AJ243992.1
1442	A/fowl plaguq virus/Rostock/34 (H7N1) PB2 protein	AJ243991.1
1443	A/fowl plaguq virus/Rostock/34 (H7N1) PB1 protein	AJ243990.1
1444	A/ostrich/South Africa/5352/92(H7N1) hemagglutinin precursor (HA)	U20458.1
1445	A/rhea/North Carolina/39482/93(H7N1) hemagglutinin precursor (HA)	U20468.1
1446	A/turkey/Italy/3775/99 (H7N1) hemagglutinin (HA)	AJ493472.1
1447	A/turkey/Italy/4603/99 (H7N1) hemagglutinin (HA)	AJ493471.1
1448	A/turkey/Italy/4602/99 (H7N1) hemagglutinin (HA)	AJ493470.1
1449	A/turkey/Italy/4169/99 (H7N1) hemagglutinin (HA)	AJ493468.1
1450	A/turkey/Italy/4073/99 (H7N1) hemagglutinin (HA)	AJ493467.1
1451	A/turkey/Italy/3889/99 (H7N1) hemagglutinin (HA)	AJ493466.1
1452	A/turkey/Italy/12598/99(H7N1) haemagglutinin (HA)	AJ489520.1
1453	A/turkey/Italy/4580/99(H7N1) haemagglutinin (HA)	AJ416628.1

1454	A/Turkey/Italy/335/00(H7N1) haemagglutinin (HA)	AJ493217.1
1455	A/Turkey/Italy/13468/99(H7N1) haemagglutinin (HA)	AJ493216.1
1456	A/Turkey/Italy/13467/99(H7N1) haemagglutinin (HA)	AJ493213.1
1457	A/chicken/CT/9407/2003(H7N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107478.1
1458	A/chicken/NY/116124/2003(H7N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107479.1
1459	A/chicken/PA/143586/2002(H7N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107477.1
1460	A/duck/Hong Kong/293/78(H7N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107474.1
1461	A/duck/Hong Kong/293/78(H7N2) hemagglutinin precursor (HA)	U20461.1
1462	A/laughing gull/Delaware/2838/87 (H7N2) nonfunctional matrix protein	AY664427.1
1463	A/pheasant/NJ/30739-9/2000(H7N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107481.1
1464	A/ruddy turnstone/Delaware/130/99 (H7N2) onfunctional matrix protein	AY664451.1
1465	A/unknown/149717-12/2002(H7N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107480.1
1466	A/unknown/NY/74211-5/2001(H7N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107476.1
1467	A/unknown/149717-12/2002(H7N2) matrix protein 1 (M) and matrix protein 2(M)	DQ107480.1
1468	A/unknown/NY/74211-5/2001(H7N2) matrix protein 1(M) and matrix protein 2 (M)	DQ107476.1
1469	A/chicken/British Columbia/CN7-3/04 (H7N3) hemagglutinin (HA)	AY644402.1
1470	A/chicken/British Columbia/CN7-3/04 (H7N3) matrix protein (M1)	AY677732.1
1471	A/chicken/Italy/270638/02(H7N3) hemagglutinin (HA)	EU158111.1
1472	A/gadwall/MD/3495/83(H7N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107488.1
1473	A/mallard/Alberta/22/2001(H7N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107482.1
1474	A/mallard/Alberta/699/81(H7N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107487.1
1475	A/pintail/Alberta/25/2001(H7N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107483.1
1476	A/Quail/Arkansas/16309-7/94 (H7N3) hemagglutinin protein subunit 1 precursor (HA1)	AF072401.1
1477	A/ruddy turnstone/New Jersey/65/85(H7N3) nonfunctional matrix protein	AY664433.1
1478	A/turkey/England/63(H7N3) hemagglutinin precursor (HA)	U20462.1
1479	A/Turkey/Colorado/13356/91 (H7N3) hemagglutinin protein subunit 1 precursor (HA1)	AF072400.1
1480	A/turkey/MN/1200/80(H7N3)) matrix protein 1 (M) and matrix protein 2 (M)	DQ107486.1
1481	A/turkey/MN/1818/82(H7N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107489.1
1482	A/turkey/Minnesota/1237/80(H7N3) hemagglutinin precursor (HA)	U20466.1
1483	A/turkey/TX/1/79(H7N3) matrix protein 1 (M) and matrix protein 2 (M)	DQ107484.1
1484	A/Turkey/Oregon/71(H7N3) hemagglutinin	AF497557.1
1485	A/Turkey/Utah/24721-10/95 (H7N3) hemagglutinin protein subunit 1 precursor (HA1)	AF072402.1
1486	A/softbill/South Africa/142/92(H7N4) hemagglutinin precursor (HA)	U20464.1
1487	A/ruddy turnstone/Delaware/2770/87 (H7N5) nonfunctional matrix protein	AY664476.1
1488	A/chicken/Brescia/1902(H7N7) hemagglutinin 1 chain (HA)	U20471.1
1489	A/chicken/Jena/1816/87(H7N7) hemagglutinin precursor (HA)	U20469.1
1490	A/chicken/Leipzig/79(H7N7) hemagglutinin precursor (HA)	U20459.1
1491	A/duck/Heinersdorf/S495/6/86(H7N7) hemagglutinin precursor (HA)	U20465.1
1492	A/equine/Prague/1/56 (H7N7) neuraminidase	U85989.1
1493	A/equine/Santiago/77(H7N7) nucleoprotein	AY383752.1
1494	A/equine/Santiago/77(H7N7) neuraminidase	AY383757.1
1495	A/equine/Santiago/77(H7N7) hemagglutinin	AY383756.1
1496	A/FPV/Weybridge(H7N7) matrix protein	M38299.1
1497	A/goose/Leipzig/187/7/1979(H7N7) hemagglutinin	L43914.1
1498	A/goose/Leipzig/192/7/1979(H7N7) hemagglutinin	L43915.1

1499	A/goose/Leipzig/137/8/1979(H7N7) hemagglutinin	L43913.1
1500	A/ruddy turnstone/Delaware/134/99 (H7N7) nonfunctional matrix protein	AY664468.1
1501	A/seal/Mass/1/80 H7N7 recombinant	S73497.1
1502	A/swan/Potsdam/63/6/81(H7N7) hemagglutinin precursor (HA)	U20467.1
1503	A/tern/Potsdam/342/6/79(H7N7) hemagglutinin precursor (HA)	U20470.1
1504	A/pintail/Alberta/121/79(H7N8) matrix protein 1 (M) and matrix protein 2 (M)	DQ107485.1
1505	A/Turkey/Minnesota/38429/88(H7N9) hemagglutinin	AF497551.1
H8N*		
1506	A/turkey/Ontario/6118/1968(H8N4) segment 6 neuraminidase (NA)	EU429793.1
1507	A/Mallard Duck/Alberta/357/84(H8N4) segment 4 hemagglutinin (HA1)	AF310988.1
1508	A/Pintail Duck/Alberta/114/79(H8N4) segment 4 hemagglutinin (HA1)	AF310987.1
1509	A/duck/Eastern China/01/2005(H8N4) segment 6 neuraminidase (NA)	EU429780.1
1510	A/Red Kont/Delaware/254/94(H8N4) segment 4 hemagglutinin (HA1)	AF310989.1
H9N*		
1511	A/chicken/Amioz/1527/03(H9N2) nucleoprotein	DQ116511.1
1512	A/chicken/Amioz/1527/03(H9N2) neuraminidase	DQ116081.1
1513	A/chicken/Amioz/1527/03(H9N2) hemagglutinin	DQ108911.1
1514	A/chicken/Alonim/1953/104(H9N2) hemagglutinin	DQ108928.1
1515	A/chicken/Alonim/1552/03(H9N2) hemagglutinin	DQ108914.1
1516	A/chicken/Alonim/1552/03(H9N2) nucleoprotein	DQ116514.1
1517	A/chicken/Alonim/1965/04(H9N2) hemagglutinin	DQ108929.1
1518	A/Chicken/Anhui/1/98(H9N2) hemagglutinin (HA)	AF461511.1
1519	A/Chicken/Beijing/1/95(H9N2) nonfunctional matrix protein	AF536719.1
1520	A/Chicken/Beijing/1/95(H9N2) nucleoprotein (NP)	AF536699.1
1521	A/Chicken/Beijing/1/95(H9N2) nonfunctional nonstructural protein	AF536729.1
1522	A/Chicken/Beijing/1/95(H9N2) segment 6 neuraminidase (NA)	AF536709.1
1523	A/Chicken/Beijing/2/97(H9N2) nucleoprotein (NP)	AF536700.1
1524	A/Chicken/Beijing/2/97(H9N2) nonfunctional matrix protein	AF536720.1
1525	A/Chicken/Beijing/2/97(H9N2) nonfunctional nonstructural protein	AF536730.1
1526	A/Chicken/Beijing/2/97(H9N2) segment 6 neuraminidase (NA)	AF536710.1
1527	A/Chicken/Beijing/1/97(H9N2) hemagglutinin (HA)	AF461530.1
1528	A/Chicken/Beijing/3/99(H9N2) nonfunctional matrix protein	AF536721.1
1529	A/Chicken/Beijing/3/99(H9N2) nucleoprotein (NP)	AF536701.1
1530	A/Chicken/Beijing/3/99(H9N2) nonfunctional nonstructural protein	AF536731.1
1531	A/Chicken/Beijing/3/99(H9N2) segment 6 neuraminidase (NA)	AF536711.1
1532	A/chicken/Beit Alfa/1282/03(H9N2)hemagglutinin	DQ104476.1
1533	A/chicken/Beit-Aran/29/05(H9N2) hemagglutinin	DQ108931.1
1534	A/chicken/Bnei Darom/1557/03(H9N2) hemagglutinin	DQ108915.1
1535	A/chicken/Ein Habsor/1808/04(H9N2) hemagglutinin	DQ108925.1
1536	A/Chicken/Gangxi/2/00(H9N2) hemagglutinin (HA)	AF461514.1
1537	A/Chicken/Gangxi/1/00(H9N2) hemagglutinin (HA)	AF461513.1
1538	A/chicken/Gan Shomron/1465/03(H9N2) hemagglutinin	DQ104480.1
1539	A/chicken/Gan Shomron/1292/03(H9N2) hemagglutinin	DQ104478.1

1540	A/chicken/Gan_Shomron/1465/03(H9N2) nucleoprotein	DQ116506.1
1541	A/chicken/Gan_Shomron/1465/03(H9N2) neuraminidase	DQ116077.1
1542	A/chicken/Gan Shomron/1543/04(H9N2) nucleoprotein	DQ116512.1
1543	A/chicken/Gan Shomron/1543/04(H9N2) hemagglutinin	DQ108912.1
1544	A/Chicken/Guangdong/97(H9N2) nonfunctional matrix protein	AF536722.1
1545	A/Chicken/Guangdong/97(H9N2) nucleoprotein (NP)	AF536702.1
1546	A/Chicken/Guangdong/97(H9N2) nonfunctional nonstructural protein	AF536732.1
1547	A/Chicken/Guangdong/97(H9N2) segment 6 neuraminidase (NA)	AF536712.1
1548	A/Chicken/Gansu/1/99(H9N2) hemagglutinin (HA)	AF461512.1
1549	A/chicken/Gujrat/India/3697/2004(H9N2) polymerase basic 2 (PB2)	DQ979865.1
1550	A/chicken/Haryana/India/2424/2004(H9N2) polymerase basic 2 (PB2)	DQ979862.1
1551	A/Chicken/Henan/98(H9N2) nonfunctional matrix protein	AF536726.1
1552	A/Chicken/Henan/98(H9N2) nucleoprotein (NP)	AF536706.1
1553	A/Chicken/Henan/98(H9N2) nonfunctional nonstructural protein	AF536736.1
1554	A/Chicken/Henan/2/98(H9N2) hemagglutinin (HA)	AF461517.1
1555	A/Chicken/Henan/1/99(H9N2) hemagglutinin (HA)	AF461516.1
1556	A/Chicken/Henan/98(H9N2) segment 6 neuraminidase (NA)	AF536716.1
1557	A/Chicken/Hebei/1/96(H9N2) nonfunctional matrix protein	AF536723.1
1558	A/Chicken/Hebei/1/96(H9N2) segment 6 nonfunctional neuraminidase protein	AF536713.1
1559	A/Chicken/Hebei/1/96(H9N2) nucleoprotein (NP)	AF536703.1
1560	A/Chicken/Hebei/1/96(H9N2) nonfunctional nonstructural protein	AF536733.1
1561	A/Chicken/Hebei/1/96(H9N2) segment 6 nonfunctional neuraminidase protein	AF536713.1
1562	A/Chicken/Hebei/2/00(H9N2) hemagglutinin (HA)	AF461531.1
1563	A/Chicken/Hebei/2/98(H9N2) nonfunctional matrix protein	AF536724.1
1564	A/Chicken/Hebei/2/98(H9N2) nucleoprotein (NP)	AF536704.1
1565	A/Chicken/Hebei/2/98(H9N2) nonfunctional nonstructural protein	AF536734.1
1566	A/Chicken/Hebei/2/98(H9N2) segment 6 neuraminidase (NA)	AF536714.1
1567	A/Chicken/Hebei/1/00(H9N2) hemagglutinin (HA)	AF461515.1
1568	A/Chicken/Hebei/3/98(H9N2) nucleoprotein (NP)	AF536705.1
1569	A/Chicken/Hebei/3/98(H9N2) nonfunctional matrix protein	AF536725.1
1570	A/Chicken/Hebei/3/98(H9N2) nonfunctional onstructural protein	AF536735.1
1571	A/Chicken/Hebei/3/98(H9N)) segment 6 neuraminidase (NA)	AF536715.1
1572	A/chicken/Hong Kong/FY313/2000(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107508.1
1573	A/chicken/Hong Kong/WF208/2001(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107513.1
1574	A/chicken/Hong Kong/NT471/2002(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107514.1
1575	A/chicken/Hong Kong/WF2/99(H9N2) hemagglutinin	AY206677.1
1576	A/chicken/Iarah/1376/03(H9N2) nucleoprotein	DQ116504.1
1577	A/chicken/Iarah/1376/03(H9N2) neuraminidase	DQ116075.1
1578	A/chicken/Iarah/1376/03(H9N2) hemagglutinin	DQ108910.1
1579	A/chicken/India/2793/2003(H9N2) hemagglutinin (HA)	AY336597.1
1580	A/chicken/Iran/101/1998(H9N2) matrix protein 2 (M2)	EU477375.1
1581	A/Chicken/Jiangsu/1/99(H9N)) hemagglutinin (HA)	AF461509.1
1582	A/Chicken/Jiangsu/2/98(H9N2) hemagglutinin (HA)	AF461510.1
1583	A/chicken/Kfar Monash/636/02(H9N2) hemagglutinin	DQ104464.1
1584	A/chicken/Kalanit/1966/06.12.04(H9N2) hemagglutinin	DQ108930.1

1585	A/chicken/Kalanit/1946/04(H9N2) hemagglutinin	DQ108927.1
1586	A/chicken/Korea/S4/2003(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107517.1
1587	A/Chicken/Korea/MS96/96(H9N2) matrix protein 1 and 2 (M)	AF203788.1
1588	A/Chicken/Korea/MS96/96(H9N2) neuraminidase subtype 2	AF203786.1
1589	A/Chicken/Korea/MS96/96(H9N2) nucleoprotein	AF203787.1
1590	A/Chicken/Liaoning/99(H9N2) nonfunctional matrix protein	AF536727.1
1591	A/Chicken/Liaoning/1/00(H9N2) hemagglutinin (HA)	AF461518.1
1592	A/Chicken/Liaoning/99(H9N2) nucleoprotein (NP)	AF536707.1
1593	A/Chicken/Liaoning/99(H9N2) nonfunctional matrix protein	AF536727.1
1594	A/Chicken/Liaoning/99(H9N2) nonfunctional onstructural protein	AF536737.1
1595	A/Chicken/Liaoning/2/00(H9N2) hemagglutinin (HA)	AF461519.1
1596	A/chicken/Liaoning/99(H9N2) segment 6 neuraminidase (NA)	AF536717.1
1597	A/chicken/Mudanjiang/0823/2000(H9N2) nucleoprotein (NP)	AY496851.1
1598	A/Chicken/Mudanjiang/0823/2000 (H9N2) nonstructural protein	AY631868.1
1599	A/Chicken/Mudanjiang/0823/00 (H9N2) hemagglutinin (HA)	AY513715.1
1600	A/chicken/Mudanjiang/0823/2000(H9N2) matrix protein (M1)	AY496852.1
1601	A/chicken/Mudanjiang/0823/2000(H9N2) nucleoprotein (np)	AY496851.1
1602	A/chicken/Maale HaHamisha/90658/00(H9N2) hemagglutinin	DQ104472.1
1603	A/chicken/Maanit/1477/03(H9N2) hemagglutinin	DQ104483.1
1604	A/chicken/Maanit/1291/03(H9N2) hemagglutinin	DQ104477.1
1605	A/chicken/Maanit/1275/03(H9N2) hemagglutinin	DQ104457.1
1606	A/chicken/Maanit/1477/03(H9N2) nucleoprotein	DQ116508.1
1607	A/chicken/Netohah/1373/03 (H9N2) nucleoprotein	DQ116503.1
1608	A/chicken/Netohah/1373/03 (H9N2) neuraminidase	DQ116074.1
1609	A/chicken/Netohah/1373/03 (H9N2) hemagglutinin	DQ108909.1
1610	A/chicken/Neve Ilan/1504/03(H9N2) hemagglutinin	DQ104484.1
1611	A/chicken/Neve_Ilan/1504/03(H9N2) nucleoprotein	DQ116509.1
1612	A/chicken/Neve_Ilan/1504/03(H9N2) neuraminidase	DQ116079.1
1613	A/chicken/Orissa/India/2317/2004(H9N2) polymerase basic 2 (PB2)	DQ979861.1
1614	A/chicken/Pardes-Hana-Carcur/1475/03(H9N2) hemagglutinin	DQ104482.1
1615	A/chicken/Pardes-Hana-Carcur/1475/03(H9N2) neuraminidase	DQ116078.1
1616	A/chicken/Saar/1456/03(H9N2) hemagglutinin	DQ104479.1
1617	A/chicken/Sde_Uziahu/1747/04(H9N2) neuraminidase	DQ116068.1
1618	A/chicken/Sede Uzziyyahu/1651/04(H9N2) hemagglutinin	DQ108923.1
1619	A/chicken/Sde Uziahu/1747/04(H9N2)	DQ108905.1
1620	A/chicken/Singapore/1998(H9N2) M2 protein	EU014142.1
1621	A/chicken/Singapore/1998(H9N2) M2 protein	EU014142.1
1622	A/Chicken/Shandong/98(H9N2) nonfunctional matrix protein	AF536728.1
1623	A/Chicken/Shandong/1/98(H9N2) hemagglutinin (HA)	AF461520.1
1624	A/Chicken/Shandong/98(H9N2) nucleoprotein (NP)	AF536708.1
1625	A/Chicken/Shandong/98(H9N2) nonfunctional nonstructural protein	AF536738.1
1626	A/Chicken/Shandong/98(H9N2) segment 6 neuraminidase (NA)	AF536718.1
1627	A/Chicken/Shandong/2/99(H9N2) hemagglutinin (HA)	AF461521.1
1628	A/chicken/Shandong/1/02(H9N2) neuraminidase (NA)	AY295761.1
1629	A/Chicken/Shanghai/F/98(H9N2) hemagglutinin	AF461532.1

1630	A/Chicken/Shanghai/1/02(H9N2) hemagglutinin	AY281745.1
1631	A/Chicken/Shanghai/2/99(H9N2)) hemagglutinin (HA)	AF461522.1
1632	A/Chicken/Shanghai/3/00(H9N2)) hemagglutinin (HA)	AF461523.1
1633	A/Chicken/Shanghai/F/98(H9N2) hemagglutinin (HA)	AY743216.1
1634	A/Chicken/Shanghai/4-2/01(H9N2) hemagglutinin (HA)	AF461525.1
1635	A/Chicken/Shanghai/4-1/01(H9N2) hemagglutinin (HA)	AF461524.1
1636	A/Chicken/Shanghai/4/01(H9N2) hemagglutinin (HA)	AY083841.1
1637	A/Chicken/Shanghai/3/01(H9N2) hemagglutinin HA)	AY083840.1
1638	A/chicken/Talmei_Elazar/1304/03(H9N2)nucleoprotein	DQ116530.1
1639	A/chicken/Talmei_Elazar/1304/03(H9N2) neuraminidase	DQ116072.1
1640	A/Chicken/Tianjing/2/96(H9N2) hemagglutinin	AF461527.1
1641	A/Chicken/Tianjing/1/96(H9N2) hemagglutinin (HA)	AF461526.1
1642	A/chicken/Tel Adashim/811/01 (H9N2) hemagglutinin	DQ104467.1
1643	A/chicken/Tel Adashim/811/01 (H9N2) nucleoprotein	DQ116527.1
1644	A/ck/Tel_Adashim/811/01(H9N2) neuraminidase	DQ116064.1
1645	A/chicken/Tel Adashim/812/01 (H9N2) nucleoprotein	DQ116528.1
1646	A/chicken/Tel Adashim/812/01 (H9N2) hemagglutinin	DQ104468.1
1647	A/ck/Tel_Adashim/812/01(H9N2) neuraminidase	DQ116065.1
1648	A/chicken/Tel Adashim/786/01 (H9N2) nucleoprotein	DQ116524.1
1649	A/chicken/Tel Adashim/809/01 (H9N2) hemagglutinin	DQ104465.1
1650	A/chicken/Tel Adashim/809/01 (H9N2) nucleoprotein	DQ116525.1
1651	A/chicken/Tel Adashim/1469/03 (H9N2) nucleoprotein	DQ116507.1
1652	A/chicken/Tel Adashim/1469/303(H9N2) hemagglutinin	DQ104481.1
1653	A/chicken/Tel Adashim/1506/03 (H9N2) neuraminidase	DQ116080.1
1654	A/chicken/Tel Adashim/1506/03(H9N2) hemagglutinin	DQ104474.1
1655	A/chicken/Tel Adashim/1506/03 (H9N2) nucleoprotein	DQ116510.1
1656	A/chicken/Tel Adashim/1332/03(H9N2) nucleoprotein	DQ116501.1
1657	A/chicken/Tel Adashim/1321/03(H9N2) nucleoprotein	DQ116500.1
1658	A/chicken/Tel Adashim/1332/03(H9N2) hemagglutinin	DQ108907.1
1659	A/chicken/Tel Adashim/1321/03(H9N2) hemagglutinin	DQ108906.1
1660	A/chicken/Telmond/1308/03(H9N2) nucleoprotein	DQ116499.1
1661	A/chicken/Telmond/1308/03(H9N2) neuraminidase	DQ116073.1
1662	A/chicken/Telmond/1308/03(H9N2) hemagglutinin	DQ108921.1
1663	A/chicken/Tzrofa/1568/04(H9N2) nucleoprotein	DQ116519.1
1664	A/chicken/Tzrofa/1568/04(H9N2) hemagglutinin	DQ108919.1
1665	A/chicken/UP/India/2544/2004(H9N2) polymerase basic 2 (PB2)	DQ979864.1
1666	A/chicken/UP/India/2543/2004(H9N2) polymerase basic 2 (PB2)	DQ979863.1
1667	A/chicken/Wangcheng/4/2001(H9N2) nucleoprotein	AY268949.1
1668	A/chicken/Ysodot/1362/03(H9N2) nucleoprotein	DQ116502.1
1669	A/chicken/Ysodot/1362/03(H9N2) hemagglutinin	DQ108908.1
1670	A/Chicken/Yunnan/2/00(H9N2) hemagglutinin (HA)	AF461529.1
1671	A/Chicken/Yunnan/1/99(H9N2) hemagglutinin (HA)	AF461528.1
1672	A/duck/Eastern China/01/2000(H9N2) segment 6 neuraminidase (NA)	EU429725.1
1673	A/duck/Eastern China/48/2001(H9N2) segment 6 neuraminidase (NA)	EU429707.1
1674	A/duck/Eastern China/66/2003(H9N2) segment 6 neuraminidase (NA)	EU429699.1

1675	A/duck/Eastern China/80/2004(H9N2) segment 6 neuraminidase (NA)	EU429726.1
1676	A/duck/Hong Kong/448/78(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107494.1
1677	A/duck/Hong Kong/448/78(H9N2) hemagglutinin precursor	AY206673.1
1678	A/duck/Hong Kong/366/78(H9N2) hemagglutinin precursor	AY206674.1
1679	A/duck/Hong Kong/784/79(H9N2)) matrix protein 1(M) and matrix protein 2 (M)	DQ107496.1
1680	A/duck/Hong Kong/702/79(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107495.1
1681	/duck/Hong Kong/702/79(H9N2) hemagglutinin precursor	AY206672.1
1682	A/duck/Hong Kong/610/79(H9N2) hemagglutinin precursor	AY206680.1
1683	A/duck/Hong Kong/552/79(H9N2) hemagglutinin precursor	AY206679.1
1684	A/duck/Hong Kong/644/79(H9N2) hemagglutinin precursor	AY206678.1
1685	A/duck/Korea/S13/2003(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107518.1
1686	A/duck/Nanchang/4-361/2001(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107511.1
1687	A/duck/NY/83793/2002(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107499.1
1688	A/goose/MN/5733-1243/80(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107492.1
1689	A/geese/Tel Adashim/829/01(H9N2) hemagglutinin	DQ104469.1
1690	A/geese/Tel Adashim/830/01(H9N2 hemagglutinin	DQ104470.1
1691	A/ostrich/Eshkol/1436/03(H9N2) neuraminidase	DQ116076.1
1692	A/ostrich/Eshkol/1436/03(H9N2) nucleoprotein	DQ116505.1
1693	A/pigeon/Hong Kong/WF286/2000(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107509.1
1694	A/quail/Hong Kong/YU415/2002(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107516.1
1695	A/quail/Hong Kong/SSP225/2001(H9) matrix protein 1 (M) and matrix protein 2 (M)	DQ107512.1
1696	A/quail/Hong Kong/YU1495/2000(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107510.1
1697	A/quail/Hong Kong/A28945/88(H9N2) hemagglutinin precursor	AY206675.1
1698	A/shorebird/Delaware/276/99 (H9N2) nonfunctional matrix protein	AY664464.1
1699	A/shorebird/Delaware/113/2001(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107505.1
1700	A/silky chicken/Hong Kong/WF266/2002(H9N2) matrix protein 2 (M) and matrix protein 1 (M)	DQ107515.1
1701	A/shorebird/Delaware/77/2001(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107497.1
1702	A/guinea fowl/Hong Kong/WF10/99(H9N2) hemagglutinin precursor	AY206676.1
1703	A/swine/Hangzhou/1/2006(H9N2) nucleocapsid protein (NP)	DQ907704.1
1704	A/swine/Hangzhou/1/2006(H9N2)) matrix protein 1 (M1)	EF055887.1
1705	A/swine/Hangzhou/1/2006(H9N2)) nonstructural protein 1 (NS1)	DQ823385.1
1706	A/Sw/ShanDong/1/2003(H9N2) hemagglutinin (HA)	AY294658.1
1707	A/turkey/CA/6889/80(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107491.1
1708	A/turkey/TX/28737/81(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107493.1
1709	A/turkey/MN/511/78(H9N2) matrix protein 1 (M) and matrix protein 2 (M)	DQ107490.1
1710	A/turkey/Beit Herut/1267/03(H9N2) hemagglutinin	DQ104485.1
1711	A/turkey/Beit HaLevi/1009/02(H9N2) hemagglutinin	DQ104473.1
1712	A/turkey/Beit Herut/1265/03(H9N2) hemagglutinin	DQ104456.1
1713	A/turkey/Beit_HaLevi/1562/03(H9N2) nucleoprotein	DQ116515.1
1714	A/turkey/Beit_HaLevi/1566/04(H9N2) nucleoprotein	DQ116517.1
1715	A/turkey/Beit_HaLevi/1562/03(H9N2) neuraminidase	DQ116083.1
1716	A/turkey/Beit_HaLevi/1566/04(H9N2) neuraminidase	DQ116084.1
1717	A/turkey/Beit_Herut/1267/03(H9N2) neuraminidase	DQ116070.1
1718	A/turkey/Beit_Herut/1265/03(H9N2) neuraminidase	DQ116069.1
1719	A/turkey/Beit HaLevi/1566/04(H9N2) hemagglutinin	DQ108917.1

1720	A/turkey/Bezat/89/05(H9N2) hemagglutinin	DQ108922.1
1721	A/turkey/Brosh/1276/03(H9N2) hemagglutinin	DQ104458.1
1722	A/turkey/Brosh/1276/03(H9N2) neuraminidase	DQ116071.1
1723	A/turkey/Emek Hefer/1272/03(H9N2) hemagglutinin	DQ104475.1
1724	A/turkey/Ein Habsor/1804/04(H9N2) hemagglutinin	DQ108924.1
1725	A/turkey/Ein Tzurim/1172/02(H9N2) hemagglutinin	DQ104451.1
1726	A/turkey/Ein Tzurim/1738/04(H9N2) hemagglutinin	DQ108920.1
1727	A/turkey/Ein_Tzurim/1738/04(H9N2) neuraminidase	DQ116085.1
1728	A/turkey/Gyvat Haim Ehud/1544/03(H9N2)hemagglutinin	DQ108913.1
1729	A/turkey/Givat Haim/810/01 (H9N2) hemagglutinin	DQ104466.1
1730	A/turkey/Givat Haim/810/01 (H9N2) nucleoprotein	DQ116526.1
1731	A/turkey/Givat Haim/868/02(H9N2) hemagglutinin	DQ104471.1
1732	A/turkey/Givat Haim/622/02(H9N2) hemagglutinin	DQ104462.1
1733	A/turkey/Givat_Haim/965/02(H9N2) nucleoprotein	DQ116498.1
1734	A/turkey/Gyvat_Haim_Ehud/1544/03(H9N2) nucleoprotein	DQ116513.1
1735	A/turkey/Gyvat_Haim_Ehud/1544/03(H9N2) neuraminidase	DQ116082.1
1736	A/tk/Givat_Haim/810/25.12.01(H9N2) neuraminidase	DQ116063.1
1737	A/turkey/Givat_Haim/622/02(H9N2)) neuraminidase	DQ116060.1
1738	A/turkey/Givat_Haim/965/02(H9N2) neuraminidase	DQ116057.1
1739	A/turkey/Hod_Ezyon/699/02(H9N2) neuraminidase	DQ116062.1
1740	A/turkey/Mishmar Hasharon/619/02 (H9N2) hemagglutinin	DQ104461.1
1741	A/turkey/Mishmar_Hasharon/619/02(H9N2) neuraminidase	DQ116059.1
1742	A/turkey/Kfar_Vitkin/616/02(H9N2) neuraminidase	DQ116058.1
1743	A/turkey/Kfar Vitkin/616/02 (H9N2) hemagglutinin	DQ104460.1
1744	A/turkey/Kfar Vitkin/615/02 (H9N2)hemagglutinin	DQ104459.1
1745	A/turkey/Kfar Vitkin/615/02 (H9N2) nucleoprotein	DQ116520.1
1746	A/turkey/Kfar_Vitkin/616/02(H9N2)) nucleoprotein	DQ116521.1
1747	A/turkey/Kfar Warburg/1224/03(H9N2) hemagglutinin	DQ104455.1
1748	A/tk/Kfar_Vitkin/615/02(H9N)) neuraminidase	DQ116067.1
1749	A/turkey/Mishmar_Hasharon/619/02(H9N2) nucleoprotein	DQ116522.1
1750	A/turkey/Naharia/1013/02(H9N2) hemagglutinin	DQ104449.1
1751	A/turkey/Nahalal/1547/04(H9N2) hemagglutinin	DQ108932.1
1752	A/turkey/Neve Ilan/90710/00 (H9N2) nucleoprotein	DQ116529.1
1753	A/tk/Neve_Ilan/90710/00(H9N2) neuraminidase	DQ116066.1
1754	A/turkey/Qevuzat_Yavne/1242/03(H9N2) neuraminidase	DQ116086.1
1755	A/turkey/Sapir/1199/02(H9N2) hemagglutinin	DQ104452.1
1756	A/turkey/Shadmot Dvorah/1567/04(H9N2) nucleoprotein	DQ116518.1
1757	A/turkey/Shadmot Dvorah/1567/04(H9N2) hemagglutinin	DQ108918.1
1758	A/turkey/Tzur Moshe/1565/04(H9N2) nucleoprotein	DQ116516.1
1759	A/turkey/Tzur Moshe/1565/04(H9N2) hemagglutinin	DQ108916.1
1760	A/turkey/Yedidia/625/02 (H9N2) hemagglutinin	DQ104463.1
1761	A/turkey/Yedidia/625/02 (H9N2) nucleoprotein	DQ116523.1
1762	A/turkey/Yedidia/625/02 (H9N2) neuraminidase	DQ116061.1
1763	A/turkey/Yedidia/911/02(H9N2) hemagglutinin	DQ104448.1
1764	A/turkey/Avigdor/1215/03(H9N2) hemagglutinin	DQ104454.1

1765	A/turkey/Avigdor/1209/03(H9N2) hemagglutinin	DQ104453.1
1766	A/turkey/Avichail/1075/02(H9N2) hemagglutinin	DQ104450.1
1767	A/turkey/Avigdor/1920/04(H9N2) hemagglutinin	DQ108926.1
1768	A/pintail/Alberta/49/2003(H9N5) matrix protein 1 (M) and matrix protein 2 (M)	DQ107498.1
1769	A/red knot/Delaware/2552/87 (H9N5) nonfunctional matrix protein	AY664472.1
1770	A/duck/Hong Kong/147/77(H9N6) hemagglutinin precursor	AY206671.1
1771	A/shorebird/Delaware/270/2001(H9N7) matrix protein 1 (M) and matrix protein 2 (M)	DQ107504.1
1772	A/shorebird/Delaware/277/2000(H9N7) matrix protein 1 (M) and matrix protein 2 (M)	DQ107507.1
1773	A/shorebird/Delaware/275/2001(H9N7)) matrix protein 2 (M) and matrix protein 1 (M)	DQ107506.1
1774	A/ruddy turnstone/Delaware/116/98 (H9N8) nonfunctional matrix protein	AY664435.1
1775	A/shorebird/Delaware/141/2002(H9N9) matrix protein 1 (M) and matrix protein 2 (M)	DQ107503.1
1776	A/ruddy turnstone/Delaware/103/2002(H9N9) matrix protein 1 (M) and matrix protein 2 (M)	DQ107502.1
1777	A/shorebird/Delaware/29/2002(H9N9) matrix protein 1 (M) and matrix protein 2 (M)	DQ107501.1
1778	A/shorebird/Delaware/18/2002(H9N9) matrix protein 1 (M) and matrix protein 2 (M)	DQ107500.1
1779	A/ruddy turnstone/Delaware/259/98 (H9N9) nonfunctional matrix protein	AY664469.1
H10N*		
1780	A/duck/Eastern China/527/2003(H10N3) segment 6 neuraminidase (NA)	EU429716.1
1781	A/duck/Eastern China/495/2003(H10N3) segment 6 neuraminidase (NA)	EU429715.1
1782	A/duck/Eastern China/372/2003(H10N3) segment 6 neuraminidase (NA)	EU429714.1
1783	A/duck/Eastern China/488/2003(H10N3) segment 6 neuraminidase (NA)	EU429712.1
1784	A/duck/Eastern China/453/2002(H10N3) segment 6 neuraminidase (NA)	EU429711.1
1785	A/duck/Eastern China/412/2003(H10N3) segment 6 neuraminidase (NA)	EU429710.1
1786	A/duck/Eastern China/404/2003(H10N3) segment 6 neuraminidase (NA)	EU429709.1
1787	A/duck/Eastern China/397/2003(H10N3) segment 6 neuraminidase (NA)	EU429708.1
1788	A/duck/Eastern China/502/2003(H10N3) segment 6 neuraminidase (NA)	EU429705.1
1789	A/duck/Eastern China/395/2003(H10N3) segment 6 neuraminidase (NA)	EU429704.1
1790	A/duck/Eastern China/356/2003(H10N3) segment 6 neuraminidase (NA)	EU429703.1
1791	A/duck/Eastern China/368/2003(H10N3) segment 6 neuraminidase (NA)	EU429702.1
1792	A/chicken/Singapore/1993(H10N5) M2 protein	EU014145.1
1793	A/red knot/Delaware/2561/87 (H10N5) nonfunctional matrix protein	AY664441.1
1794	A/chicken/Germany/N/1949(H10N7) segment 6 neuraminidase (NA)	EU429796.1
1795	A/ruddy turnstone/Delaware/2764/87 (H10N7) nonfunctional matrix protein	AY664462.1
1796	A/mallard/Alberta/71/98 (H10N7) nonfunctional matrix protein	AY664485.1
1797	A/mallard/Alberta/90/97 (H10N7) nonfunctional matrix protein	AY664446.1
1798	A/mallard/Alberta/110/99(H10N7) nonfunctional matrix protein	AY664481.1
1799	A/mallard/Alberta/297/77 (H10N7) nonfunctional matrix protein	AY664430.1
1800	A/mallard/Alberta/223/98 (H10N8) nonfunctional matrix protein	AY664486.1
H11N*		
1801	A/ruddy turnstone/New Jersey/51/85 (H11N1) nonfunctional matrix protein	AY664479.1
1802	A/duck/Nanchang/1749/1992(H11N2) nucleoprotein (NP)	U49094.1
1803	A/duck/Hong Kong/62/1976(H11N2) polymerase (PB1)	U48280.1
1804	A/duck/Yangzhou/906/2002(H11N2) hemagglutinin	DQ080993.1
1805	A/shorebird/Delaware/86/99 (H11N2) nonfunctional matrix protein	AY664463.1

1806	A/ruddy turnstone/Delaware Bay/2762/1987(H11N2)polymerase PB2 (PB2)	CY126279.1
1807	A/ruddy turnstone/Delaware/2762/87 (H11N2) nonfunctional matrix protein	AY664459.1
1808	A/ruddy turnstone/Delaware Bay/2762/1987(H11N2) polymerase PB1 (PB1) and PB1-F2 protein (PB1-F2)	CY126278.1
1809	A/ruddy turnstone/Delaware/2589/87 (H11N4) nonfunctional matrix protein	AY664478.1
1810	A/duck/England/1/1956(H11N6) segment 6 neuraminidase (NA)	EU429795.1
1811	A/mallard/Alberta/125/99 (H11N6) nonfunctional matrix protein	AY664483.1
1812	A/duck/Memphis/546/1974(H11N9) segment 6 neuraminidase (NA)	EU429798.1
1813	A/mallard/Alberta/122/99 (H11N9) nonfunctional matrix protein	AY664444.1
H12N*		
1814	A/Mallard Duck/Alberta/342/83(H12N1) segment 4 hemagglutinin (HA1)	AF310991.1
1815	A/ruddy turnstone/Delaware/67/98(H12N4) nonfunctional matrix protein	AY664470.1
1816	A/Ruddy Turnstone/Delaware/67/98(H12N4) segment 4 hemagglutinin (HA1)	AF310990.1
1817	A/mallard/Alberta/52/97 (H12N5) nonfunctional matrix protein	AY664448.1
1818	A/mallard/Alberta/223/77 (H12N5) nonfunctional matrix protein	AY664431.1
1819	A/Laughing Gull/New Jersey/171/92(H12N5) segment 4 hemagglutinin (HA1)	AF310992.1
1820	A/ruddy turnstone/Delaware/265/98 (H12N8) nonfunctional matrix protein	AY664438.1
H13N*		
1821	A/herring gull/New Jersey/782/86 (H13N2) nonfunctional matrix protein	AY664475.1
1822	A/shorebird/Delaware/224/97 (H13N6) nonfunctional matrix protein	AY664421.1
other		
1823	A/PR/8/34 (H1N1) x A/England/939/69 (H3N2) PB1 protein	AJ564806.1
1824	A/PR/8/34 (H1N1) x A/England/939/69 (H3N2)PB2 protein	AJ564804.1
1825	A/duck/Czechslovakia/56(H4N6) x A/USSR/90/77(H1N1)) neuraminidase (NA)	EU643639.1
1826	A/duck/Czechslovakia/56(H4N6) x A/USSR/90/77(H1N1)) neuraminidase (NA)	EU643638.1
1827	A/duck/Ukraine/63(H3N8) x A/USSR/90/77(H1N1)) neuraminidase (NA)	EU643637.1
1828	A/duck/Ukraine/63(H3N8) x A/USSR/90/77(H1N1)) neuraminidase (NA)	EU643636.1
1829	RCB1-XXI: A/USSR/90/77(H1N1)xA/Duck/Czechoslov 56 (H4N6) segment 4 hemagglutinin	AF290438.1
1830	RCB1: A/USSR/90/77(H1N1)xA/Duck/Czechoslov 56 (H4N6) hemagglutinin	AF290437.1
1831	PX14-XIII (A/USSR/90/77(H1N1)xA/Pintail Duck/Primorie/695/76(H2N3)) segment 4 hemagglutinin	AF290442.1
1832	PX14(A/USSR/90/77(H1N1)xA/Pintail Duck/Primorie/695/76(H2N3)) segment 4 hemagglutinin	AF290441.1
1833	PX8-XIII(A/USSR/90/77(H1N1)xA/Pintail Duck/Primorie/695/76(H2N3)) segment 4 hemagglutinin	
1834	PX8(A/USSR/90/77(H1N1)xA/Pintail Duck/Primorie/695/76(H2N3)) segment 4 hemagglutinin	AF290439.1
1835	A/swine/Schleswig-Holstein/1/93 hemagglutinin (HA)	U72669.1
1836	A/swine/England/283902/93 hemagglutinin (HA)	U72668.1
1837	A/swine/England/195852/92 hemagglutinin (HA)	U72667.1
1838	A/swine/England/117316/86 hemagglutinin (HA)	U72666.1
1839	A/turkey/Germany/2482/90) hemagglutinin (HA)	U96766.1

Table 11. Influenza B Antigens

Strain/Protein	GenBank Access No.
B/Daeku/47/97 hemagglutinin	AF521237.1
B/Daeku/45/97 hemagglutinin	AF521236.1
B/Daeku/10/97 hemagglutinin	AF521221.1
B/Daeku/9/97 hemagglutinin	AF521220.1
B/Gyeonggi/592/2005 neuraminidase	DQ231543.1
B/Gyeonggi/592/2005 hemagglutinin	DQ231538.1
B/Hong Kong/5/72 neuraminidase	AF305220.1
B/Hong Kong/5/72 hemagglutinin	AF305219.1
B/Hong Kong/157/99 hemagglutinin	AF387503.1
B/Hong Kong/157/99 hemagglutinin	AF387502.1
B/Hong Kong/156/99 hemagglutinin	AF387501.1
B/Hong Kong/156/99 hemagglutinin	AF387500.1
B/Hong Kong/147/99 hemagglutinin	AF387499.1
B/Hong Kong/147/99 hemagglutinin	AF387498.1
B/Hong Kong/110/99 hemagglutinin	AF387497.1
B/Hong Kong/110/99 hemagglutinin	AF387496.1
B/Incheon/297/2005 hemagglutinin	DQ231539.1
B/Incheon/297/2005 neuraminidase	DQ231542.1
B/Lee/40 polymerase protein (PB1)	D00004.1
B/Michigan/22572/99 hemagglutinin	AY129961.1
B/Michigan/22723/99 hemagglutinin (HA)	AY112992.1
B/Michigan/22631/99 hemagglutinin (HA)	AY112991.1
B/Michigan/22587/99 hemagglutinin (HA)	AY112990.1
B/New York/20139/99 hemagglutinin	AY129960.1
B/Panama/45/90 nucleoprotein	AF005739.1
B/Panama/45/90 polymerase (PA)	AF005738.1
B/Panama/45/90 polymerase (PB2)	AF005737.1
B/Panama/45/90 polymerase (PB1)	AF005736.1
B/Pusan/250/99 hemagglutinin	AF521218.1
B/Pusan/255/99 hemagglutinin	AF521226.1
B/Pusan/270/99 hemagglutinin	AF521219.1
B/Pusan/285/99 hemagglutinin	AF521217.1
B/Riyadh/01/2007 segment 8 nuclear export protein (NEP) and non structural protein 1 (NS1)	GU135839.1
B/Seoul/6/88 hemagglutinin	AF521238.1
B/Seoul/12/88 hemagglutinin	AF521239.1
B/Seoul/1/89 hemagglutinin	AF521230.1
B/Seoul/37/91 hemagglutinin	AF521229.1
B/Seoul/38/91 hemagglutinin	AF521227.1
B/Seoul/40/91 hemagglutinin	AF521235.1
B/Seoul/41/91 hemagglutinin	AF521228.1
B/Seoul/13/95 hemagglutinin	AF521225.1
B/Seoul/12/95 hemagglutinin	AF521223.1
B/Seoul/17/95 hemagglutinin	AF521222.1

B/Seoul/21/95 hemagglutinin	AF521224.1
B/Seoul/16/97 hemagglutinin	AF521233.1
B/Seoul/19/97 hemagglutinin	AF521231.1
B/Seoul/28/97 hemagglutinin	AF521234.1
B/Seoul/31/97 hemagglutinin	AF521232.1
B/Seoul/232/2004 neuraminidase	DQ231541.1
B/Seoul/1163/2004 neuraminidase	DQ231540.1
B/Seoul/1163/2004 hemagglutinin	DQ231537.1
B/Sichuan/379/99 hemagglutinin (HA)	AF319590.1
B/Sichuan/38/2000 hemagglutinin (HA)	AF319589.1
B/South Carolina/25723/99 hemagglutinin	AY129962.1
B/Switzerland/4291/97 hemagglutinin	AF387505.1
B/Switzerland/4291/97 hemagglutinin	AF387504.1
B/Taiwan/21706/97 nonstructural protein 1 (NS1)	AF492479.1
B/Taiwan/21706/97 hemagglutinin (HA)	AF026162.1
B/Taiwan/3143/97 nonstructural protein 1 (NS1)	AF492478.1
B/Taiwan/3143/97 haemagglutinin (HA)	AF026161.1
B/Taiwan/2026/99 nonstructural protein 1 (NS1)	AF492481.1
B/Taiwan/2026/99 hemagglutinin	AY604741.1
B/Taiwan/2027/99 nonstructural protein 1 (NS1)	AF492480.1
B/Taiwan/2027/99 hemagglutinin	AY604742.1
B/Taiwan/1243/99 nonstructural protein NS1(NS1)	AF380504.1
B/Taiwan/1243/99 hemagglutinin	AY604740.1
B/Taiwan/2195/99 hemagglutinin	AY604743.1
B/Taiwan/2195/99 nonstructural protein 1 (NS1)	AF492482.1
B/Taiwan/1293/2000 nonstructural protein NS1(NS1)	AF380509.1
B/Taiwan/1293/00 hemagglutinin	AY604746.1
B/Taiwan/1293/2000 hemagglutinin (HA)	AF492477.1
B/Taiwan/1265/2000 nonstructural protein NS1 (NS1)	AF380508.1
B/Taiwan/1265/00 hemagglutinin	AY604745.1
B/Taiwan/4184/2000 nonstructural protein NS1 (NS1)	AF380507.1
B/Taiwan/4184/00 hemagglutinin (HA)	AY604750.1
B/Taiwan/31511/2000 nonstructural protein NS1 (NS1)	AF380505.1
B/Taiwan/31511/00 hemagglutinin (HA)	AY604748.1
B/Taiwan/12192/2000 hemagglutinin	AY604747.1
B/Taiwan/41010/00 hemagglutinin (HA)	AY604749.1
B/Taiwan/41010/2000 nonstructural protein NS1 (NS1)	AF380506.1
B/Taiwan/0409/00 hemagglutinin (HA)	AY604744.1
B/Taiwan/202/2001 nonstructural protein 1 (NS1)	AF380512.1
B/Taiwan/202/2001 hemagglutinin (HA)	AF366076.1
B/Taiwan/11515/2001 nonstructural protein 1 (NS1)	AF380511.1
B/Taiwan/11515/01 hemagglutinin	AY604754.1
B/Taiwan/11515/2001 hemagglutinin (HA)	AF366075.1
B/Taiwan/1103/2001 nonstructural protein NS1 (NS1)	AF380510.1
B/Taiwan/1103/01 hemagglutinin	AY604755.1

B/Taiwan/114/2001 hemagglutinin (HA), HA-4 allele	AF492476.1
B/Taiwan/2805/2001 hemagglutinin (HA)	AF400581.1
B/Taiwan/2805/01 hemagglutinin (HA)	AY604752.1
B/Taiwan/0114/01 hemagglutinin (HA)	AY604753.1
B/Taiwan/0202/01 hemagglutinin (HA)	AY604751.1
B/Taiwan/4119/02 hemagglutinin (HA)	AY604778.1
B/Taiwan/4602/02 hemagglutinin (HA)	AY604777.1
B/Taiwan/1950 /02 hemagglutinin (HA)	AY604776.1
B/Taiwan/1949/02 hemagglutinin (HA)	AY604775.1
B/Taiwan/1584 /02 hemagglutinin (HA)	AY604774.1
B/Taiwan/1561 /02 hemagglutinin (HA)	AY604773.1
B/Taiwan/ 1536/02 hemagglutinin (HA)	AY604772.1
B/Taiwan/1534 /02 hemagglutinin (HA)	AY604771.1
B/Taiwan/1503 /02 hemagglutinin (HA)	AY604770.1
B/Taiwan/1502/02 hemagglutinin (HA)	AY604769.1
B/Taiwan/1013 /02 hemagglutinin (HA)	AY604768.1
B/Taiwan/0993 /02 hemagglutinin (HA)	AY604766.1
B/Taiwan/0932 /02 hemagglutinin (HA)	AY604765.1
B/Taiwan/0927/02 hemagglutinin (HA)	AY604764.1
B/Taiwan/0880 /02 hemagglutinin (HA)	AY604763.1
B/Taiwan/0874/02 hemagglutinin (HA)	AY604762.1
B/Taiwan/0730 /02 hemagglutinin (HA)	AY604761.1
B/Taiwan/0722/02 hemagglutinin (HA)	AY604760.1
B/Taiwan/0702 /02 hemagglutinin (HA)	AY604759.1
B/Taiwan/0654/02 hemagglutinin (HA)	AY604758.1
B/Taiwan/0600/02 hemagglutinin (HA)	AY604757.1
B/Taiwan/0409 /02 hemagglutinin (HA)	AY604756.1
B/Taiwan/0879/02 nonfunctional hemagglutinin	AY604767.1
B/Taiwan/ 3532/03 hemagglutinin (HA)	AY604794.1
B/Taiwan/2551 /03 hemagglutinin (HA)	AY604793.1
B/Taiwan/ 1618/03 hemagglutinin (HA)	AY604792.1
B/Taiwan/ 1574/03 hemagglutinin (HA)	AY604791.1
B/Taiwan/1013 /03 hemagglutinin (HA)	AY604790.1
B/Taiwan/0833 /03 hemagglutinin (HA)	AY604789.1
B/Taiwan/0735 /03 hemagglutinin (HA)	AY604788.1
B/Taiwan/0699/03 hemagglutinin (HA)	AY604787.1
B/Taiwan/0684/03 hemagglutinin (HA)	AY604786.1
B/Taiwan/0616 /03 hemagglutinin (HA)	AY604785.1
B/Taiwan/0615 /03 hemagglutinin (HA)	AY604784.1
B/Taiwan/0610 /03 hemagglutinin (HA)	AY604783.1
B/Taiwan/0576 /03 hemagglutinin (HA)	AY604782.1
B/Taiwan/0569/03 hemagglutinin (HA)	AY604781.1
B/Taiwan/0562/03 hemagglutinin (HA)	AY604780.1
B/Taiwan/0002 /03 hemagglutinin (HA)	AY604779.1
B/Taiwan/773/2004 hemagglutinin (HA)	EU068195.1

B/Taiwan/187/2004 hemagglutinin (HA)	EU068194.1
B/Taiwan/3892/2004 hemagglutinin (HA)	EU068193.1
B/Taiwan/562/2004 hemagglutinin (HA)	EU068191.1
B/Taiwan/234/2004 hemagglutinin (HA)	EU068188.1
B/Taiwan/4897/2004 hemagglutinin (HA)	EU068186.1
B/Taiwan/8579/2004 hemagglutinin (HA)	EU068184.1
B/Taiwan/184/2004 hemagglutinin (HA)	EU068183.1
B/Taiwan/647/2005 hemagglutinin (HA)	EU068196.1
B/Taiwan/877/2005 hemagglutinin (HA)	EU068198.1
B/Taiwan/521/2005 hemagglutinin (HA)	EU068189.1
B/Taiwan/1064/2005 hemagglutinin (HA)	EU068192.1
B/Taiwan/3722/2005 hemagglutinin (HA)	EU068197.1
B/Taiwan/5049/2005 hemagglutinin (HA)	EU068190.1
B/Taiwan/5011/2005 hemagglutinin (HA)	EU068187.1
B/Taiwan/4659/2005 hemagglutinin (HA)	EU068185.1
B/Taiwan/25/2005 hemagglutinin (HA)	EU068182.1
B/Taiwan/1037/2005 hemagglutinin (HA)	EU068181.1
B/Taiwan/62/2005 hemagglutinin (HA)	EU068180.1
B/Taiwan/591/2005 hemagglutinin (HA)	EU068179.1
B/Taiwan/649/2005 hemagglutinin (HA)	EU068178.1
B/Taiwan/4554/2005 hemagglutinin (HA)	EU068177.1
B/Taiwan/987/2005 hemagglutinin (HA)	EU068176.1
B/Taiwan/2607/2006 hemagglutinin (HA)	EU068175.1
B/Vienna/1/99 hemagglutinin	AF387495.1
B/Vienna/1/99 hemagglutinin	AF387494.1
B/Vienna/1/99 hemagglutinin	AF387493.1
B/Vienna/1/99 hemagglutinin	AF387492.1

Table 12. Influenza C Antigens

Strain/Protein	**GenBank Access No.**
C/JHB/1/66) hemagglutinin-esterase-fusion protein (HEF) mRNA, complete cds.	AY880247.1
STRAIN C/ANN ARBOR/1/50) persistent variant segment 7 non-structural protein 1 (NS1) mRNA, complete cds	AF102027.1
(STRAIN C/ANN ARBOR/1/50) wild type segment 7 non-structural protein 1 (NS1) mRNA, complete cds	AF102026.1
(C/JHB/1/66) hemagglutinin-esterase-fusion protein (HEF) mRNA, complete cds	AY880247.1
(STRAIN C/BERLIN/1/85) mRNA for basic polymerase 2 precursor	X55992.1

Table 13: H7 Hemagglutinin Amino Acid Sequences

SEQ ID NO	Accession No / Strain / Protein	Amino Acid Sequence

2284	AAM19228 A/turkey/Minnesota/3842 9/1988 1988// HA 20335017	ACVLVEAKGDKICLGHHAVVNGTKVNTLTEKGIEVVNATETVETANIGKICTQGKRPTDLGQCGLLG TLIGPPQCDQFLEFESDLIIERREGNDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYSGIITNGAT SACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIVWGIHHSGSTTEQTKLYGSGNK LITVESSKYQQSFTPSPGARPQVNGESGRIDFHWMLLDPNDTVTFTFNGAFIAPDRASFFKGESLG VQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQPSLLLATGMRNVPENPKTRGL FGAIAGFIEKDGGSHYG
2285	AAY46211 A/mallard/Sweden/91/20 02 2002// HA 66394828	MNTQILVFALVAIIPINADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSRGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYS GIRTNGAPSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRNDPALIIWGIHHSGSTTEQTK LYGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDRASFL RGKSMGIQSGVQIDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLLLATGMKNVPEIP KGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQF ELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLREN AEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASC FILLAIAMGLVFMCVKNGNMRCTICI
2286	ABI84694 A/turkey/Minnesota/1/198 8 1988/07/13 HA 115278573	MNTQILVFIACVLVEAKGDKICLGHHAVVNGTKVNTLTEKGIEVVNATETVETANIGKICTQGKRPTDL GQCGLLGTLIGPPQCDQFLEFESDLIIERREGNDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIVWGIHHSGSTTEQT KLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWMLLDPNDTVTFTFNGAFIAPDRAS FFKGESLGVQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQPSLLLATGMRNVP ENPKTRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTNQ QFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLR ENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHAQYRAESLQNRIQIDPVKLSSGYKDIILWFSFGAS CFLLLAIAMGLVFICIKNGNMRCTICI
2287	ABS89409 A/blue-winged teal/Ohio/566/2006 2006// HA 155016324	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRPTDL GQCGLLGTLIGPPQCDQFLEFDTDLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHSGSATEQT KLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRAS FFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNVP ENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQL RENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQNRIQIDPVRLSSGYKDIILWFSFGA SCFLLLAIAMGLVFICIKNGNMRCTICI
2288	ACD03594 A/ruddy turnstone/DE/1538/2000 2000// HA 187384848	MNTQILAFIACMLVGVRGDKICLGHHAVANGTKVNTLTEKGIEVVNATETVESANIKKICTQGKRPTD LGQCGLLGTLIGPPQCDQFLEFDSDLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTY SGIRTNGATSACRRLGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHSGSANEQ TKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRA SFFRGESLGIQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNV PENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QQFELMDNEFNEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQL RENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQNRIQIDPVKLSSGYKDIILWFSFGA SCFLLLAIAMGLIFICIKNGNMRCTICI
2289	BAH22785 A/duck/Mongolia/119/200 8 2008// HA 223717820	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIGKETMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHHSGSTTEQTK LYGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASF LRGKSMGIQSGVQVDANCEGDCYHNGGTIISNLPFQNINSRTVGKCPRYVKQESLLLATGMKNVPE IPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIERTNQQ FELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSNGYKDVILWFSFGAS CFILLAIAMGLVFICVKNGNMRCTICI
2290	CAY39406 A/Anas crecca/Spain/1460/2008 2008/01/26 HA 254674376	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVD LGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHHSGSTTEQT KLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNV PEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTN QQFELIDNEFTEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMNKLYERVKRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGA SCFILLAIAMGLVFICVKNGNMRCTICI

2291	ACX53683 A/goose/Czech Republic/1848-K9/2009 2009/02/04 HA 260907763	MNIQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERRGGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHHSGSTTEQTK LYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASF LKGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVP ELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVKRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQINPVKLSSGYKDVILWFSFGA SCFILLAIAMGLVFICVKNGNMRCTICI
2292	ACZ48625 A/turkey/Minnesota/3842 9/1988 1988// HA 269826341	MNTQILVFIACVLVEAKGDKICLGHHAVVNGTKVNTLTEKGIEVVNATETVETANIGKICTQGKRPTDL GQCGLLGTLIGPPQCDQFLEFESDLIIERREGNDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIVWGIHHSGSTTEQT KLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWMLLDPNDTVTFTFNGAFIAPDRAS FFKGESLGVQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQPSLLLATGMRNVP ENPKTRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTNQ QFEL
2293	ADC29485 A/mallard/Spain/08.0099 1.3/2005 2005/11/ HA 284927336	STQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQH TIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQ IDPVKLSSGYKDVILWFSFGASCFILL
2294	ADK71137 A/blue-winged teal/Guatemala/CIP049-01/2008 2008/02/07 HA 301333785	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRPTDL GQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYS GIRTNGATSACRRSGSSSYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHSGSATEQT KLYGSGNKLITVGSSKYQQSFTPSPGTRPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRAS FLRGKSLGIQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNVP ENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QHFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQL RENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQNRIQIDPVKLSSGYKDIILWFSFGA SCFLLLAIAMGLVFICIKNGNMRCTICI
2295	ADK71148 A/blue-winged teal/Guatemala/CIP049-02/2008 2008/03/05 HA 301333804	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNXTETVETANIKKICTHGKRPTDL GQCGLLGTLIGPPQCDRFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHSGSATEQT KLYGSGNKLITVGSSKYQQSFTPSPGTRPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRAS FLRGKSLGIQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNVP ENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVKQL RENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQNRIQIDPVKLSSGYKDIILWFSFGA SCFLLLAIAMGLVFICIKNGNMRCTICI
2296	ADN34727 A/goose/Czech Republic/1848-T14/2009 2009/02/04 HA 307141869	MNIQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERRGGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYS GIRTNGXTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHHSGSTTEQTK LYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASF LKGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVP ELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVKRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQINPVKLSSGYKDVILWFSFGA SCFILLAIAMGLVFICVKNGNMRCTICI
2297	AEK84760 A/wild bird/Korea/A14/2011 2011/02/ HA 341610308	PAFIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESL MLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQIT GKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMN KLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGY KDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2298	AEK84761 A/wild bird/Korea/A3/2011 2011/02/ HA 341610310	ILVFALVAIIPTNANKIGLGHHAVSNGTKVNTLTERGVEVFNATETVERTNVPRICSKGKKTVDLGQC GLRGTITGPPQCDQFLKFSPDLIIERQKGSDVCYPGKFVNEKPLRQILRESGGIDKETMGFAYNGIKT NGPPIACRKSGSSFYAKMKWLLSNTDKAAFPQMTKSYKNTRRNPALIVWGIHHSGSTTKQTKLYGI GSNLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIPPDRASFLRGK SMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVPELPKG KGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELI DNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLRENAE EDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFIL LAIAMGLVFICVKNGNMRCTICI

2299	AEK84763 A/wild bird/Korea/A9/2011 2011/02/ HA 341610314	ILVFALVAIIPTNANKIGLGHHAVSNGTKVNTLTERGVEFFNATETVEPTNVPRICSKGKKTVDLGQC GLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEKALRQILRESGGIDKETMGFAYSGIKT NGPPIACRKSGSSFYAKMKWLLSNTDKAAFPQMTKSYKNIRRDPALIVWGIHHSGSTTKQTNLYGIG SNLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFIFNGAFIAPDRASFLIGKSM GIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVPELPKGRG LFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDN EFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLRENAEED GTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAI AMGLVFICVKNGNMRCTICI
2300	AEK84765 A/spot-billed duck/Korea/447/2011 2011/04/ HA 341610318	LVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVDLGQC GLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYSGIRT NGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGIHHSGSTTEQTKLYG SGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDRASFLRG KSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVPEPPK GRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFE LIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLRENA EEDGTGCFEIFHKCDDDCMARIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFI LLAIAMGLVFICVKNGNMRCTICI
2301	AEM98291 A/wild duck/Mongolia/1-241/2008 2008/04/ HA 344196120	SILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVDLGQ CGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYSGIR TNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHHSGSTTEQTKLYG SGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRG KSMGIQSGVQVDANCEGDCYHSGGSIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVPELPK GRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFE LIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVKRQLRENA EEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQINPVKLSSGYKDVILWFSFGASCFI LLAIAMGLVFICVKNGNMRCTI
2302	AFM09439 A/emperor goose/Alaska/44063-061/2006 2006/05/23 HA 390535062	QILAFIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETVNIKKICTQGKRPTDLGQC GLLGTLIGPPQCDQFLEFDADLIIERRKGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYSGIRT NGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHSGSATEQTKLYG SGNKLITVGSSKYQQSFVPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPERASFFRG ESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNVPENPK TRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTNQQFEL IDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQLRENAE EDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQNRIQINPVKLSSGYKDIILWFSFGASCFLLL AIAMGLVFICIKNGNMRCTICI
2303	AFV33945 A/guinea fowl/Nebraska/17096-1/2011 2011/04/05 HA 409676820	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERRIEVVNATETVETANIKKICTQGKRPTDL GQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRNKPALIVWGVHHSGSATEQT KLYGSGSKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRAS FFRGESLGVQSDVPLDSGCEGDCFHKGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNVP ENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEIWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQLR ENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESLQNRIQIDPVKLSSGYKDIILWFSFGAS CFLLLAIAMGLVFICIKNGNMRCTICI
2304	AFV33947 A/goose/Nebraska/17097-4/2011 2011/04/05 HA 409676827	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRPTDL GQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIVWGVHHSASATEQT KLYGSGSKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRAS FFRGESLGVQSDVPLDSGCEGDCFHKGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNVP ENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEIWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQLR ENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESLQNRIQIDPVKLSSGYKDIILWFSFGAS CFLLLAIAMGLVFICIKNGNMRCTICI

2305	AFX85260 A/ruddy turnstone/Delaware Bay/220/1995 1995/05/21 HA 423514912	MNTQILAFIACMLIGINGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKRICTQGKRPIDLG QCGLLGTLIGPPQCDQFLEFDSDLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYSGI RTNGATSACIRLGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHSGSANEQTKLY GSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRASFFR GESLGVQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGRCPRYVKQTSLLLATGMKNVPEN PKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTNQQ FELIDNEFNEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQLREN AEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQNRIQIDPVKLSSGYKDIILWFSFGASCF LLLAIAMGLVFICIKNGNMRCTICI
2306	AGE08098 A/northern shoverl/Mississippi/11OS 145/2011 2011/01/08 HA 444344488	MNTQILTLIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRPTDL GQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHSGSATEQT KLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRAS FFRGESLGVQSDVPLDSGCEGDCFHNGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNVP ENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQL RENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESLQNRIQIDPVKLSSGYKDIILWFSFGA SCFLLLAIAMGLVFICIKNGNMRCTICI
2307	AGI60301 A/Hangzhou/1/2013 2013/03/24 HA 475662454	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGISGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASF LRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPE IPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQ FELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLREN AEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASC FILLAIVMGLVFICVKNGNMRCTICI
2308	AGI60292 A/Shanghai/4664T/2013 2013/03/05 HA 476403560	MNTQILVFALIAIIPANADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCHHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2309	AGJ72861 A/chicken/Zhejiang/DTID-ZJU01/2013 2013/04/ HA 479280294	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGGEVVNATETVERTNIPRICSKGKKTVDL GQGGPRGTITGPPQCDQFLEFSADLIMERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQT KLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRA SFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNV PEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTN QQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGA SCFILLAIVMGLVFICVKNGNMRCTICI
2310	AGJ73503 A/Nanjing/1/2013 2013/03/28 HA 479285761	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI

2311	BAN16711 A/duck/Gunma/466/2011 2011// HA 482661571	MNTQVLVFALMAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTV DLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFT YSGIRTNGTTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIAWGIHHSGSTTE QTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDR ASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKN VPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKT NQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQ LRENAEEDDTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSF GASCFILLAIAMGLVFICVKNGNMRCTICI
2312	AGK84857 A/Hangzhou/2/2013 2013/04/01 HA 485649824	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQITKSYKNTRKSPALIVWGIHHSVSTAEQTKL YGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASF LRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPE IPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQ FELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLREN AEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASC FILLAIVMGLVFICVKNGNMRCTICI
2313	AGL44438 A/Shanghai/02/2013 2013/03/05 HA 496493389	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2314	AGL33692 A/Shanghai/4655T/2013 2013/02/26 HA 491874175	GMIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVIN WTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDD CMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNM RCTICI
2315	AGL33693 A/Shanghai/4659T/2013 2013/02/27 HA 491874186	GMIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVIN WTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDD CMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNM RCTICI
2316	AGL95088 A/Taiwan/S02076/2013 2013/04/22 HA 501485301	VFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDLGQCGLL GTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGIRTNGA TSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLYGSGN KLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRGKS MGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEIPKGR GLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELID NEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEED GTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAI VMGLVFICVKNGNMR
2317	AGL95098 A/Taiwan/T02081/2013 2013/04/22 HA 501485319	LVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDLGQCGL LGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGIRTNG ATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLYGSG NKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRGK SMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEIPKG RGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELI DNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEE DGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILL AIVMGLVFICVKNGNMRCT
2318	AGM53883 A/Shanghai/5083T/2013 2013/04/20 HA 507593986	GFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITE VWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNN TYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCT
2319	AGM53884 A/Shanghai/5180T/2013 2013/04/23 HA 507593988	AQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNA ELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKY REEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2320	AGM53885 A/Shanghai/5240T/2013 2013/04/25 HA 507593990	QNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSY NAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHS KYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCT
2321	AGM53886 A/Shanghai/4842T/2013 2013/04/13 HA 507593992	NAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYN AELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSK YREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCT
2322	AGM53887 A/Shanghai/4701T/2013 2013/04/06 HA 507593994	NAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYN AELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSK YREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTIC
2323	AGN69462 A/Wuxi/2/2013 2013/03/31 HA 511105778	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGSTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGSKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2324	AGN69474 A/Wuxi/1/2013 2013/03/31 HA 511105798	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLINGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2325	AGO51387 A/Jiangsu/2/2013 2013/04/20 HA 514390990	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYRXEAMXBXIQIDPVKLSSGYKDVXJWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2326	BAN59726 A/duck/Mongolia/147/2008 2008/08/29 HA 519661951	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIGKETMGFTYS GIRTNGATSACRRSRSSFYAEMKWLLSNTDNAAFPQMTRSYKNTRKDPALIIWGIHHSGSTTEQTK LYGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASF LRGKSMGIQSGVQVDANCEGDCYHNGGTIISNLPFQNINSRTVGKCPRYVKQESLLLATGMKNVPE IPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIERTNQQ FELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSNGYKDVILWFSFGAS CFILLAIAMGLVFICVKNGNMRCTICI
2327	BAN59727 A/duck/Mongolia/129/2010 2010// HA 519661954	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERINVPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHHSGSTTEQTK LYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASF LRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVP ELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVKRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQINPVKLSSGYKDVILWFSFGA SCFILLAIAMGLVFICVKNGNMRCTICI

2328	AGQ80952 A/duck/Jiangxi/3096/2009 2009// HA 523788794	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTSIPRICSKGKRAVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQTTKSYKNTRKDPALIIWGIHHSGSTTEQTKL YGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFL RGKSMGIQSGVQVDANCEGDCYHNGGTIISNLPFQNINSRAVGKCPRYVKQESLLLATGMKNVPEI PKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQ FELIDNEFTEVERQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIAMGLVFICVKNGNMRCTICI
2329	AGQ80989 A/duck/Jiangxi/3257/2009 2009// HA 523788868	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTSIPRICSKGKRAVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQTTKSYKNTRKDPALIIWGIHHSGSTTEQTKL YGSGNKLITVGXSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFL RGKSMGIQSGVQVDANCEGDCYHNGGTIISNLPFQNINSRAVGKCPRYVKQESLLLATGMKNVPEI PKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQ FELIDNEFTEVERQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIAMGLVFICVKNGNMRCTICI
2330	AGQ81043 A/chicken/Rizhao/515/2013 2013// HA 523788976	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEEMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2331	AGR33894 A/chicken/Rizhao/719b/2013 2013// HA 524845213	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDRSKYREEAMQNRXXXXXXXXXXXXXKDVILWFSFGA SCFILLAIVMGLVFICVKNGNMRCTICI
2332	AGR49399 A/chicken/Jiangxi/SD001/2013 2013/05/03 HA 525338528	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2333	AGR49495 A/chicken/Shanghai/S1358/2013 2013/04/03 HA 525338689	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIKNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI

2334	AGR49506 A/chicken/Shanghai/S1410/2013 2013/04/03 HA 525338708	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2335	AGR49554 A/chicken/Zhejiang/SD033/2013 2013/04/11 HA 525338789	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVRRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2336	AGR49566 A/duck/Anhui/SC702/2013 2013/04/16 HA 525338809	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDNRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2337	AGR49722 A/homing pigeon/Jiangsu/SD184/2013 2013/04/20 HA 525339071	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS EIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2338	AGR49734 A/pigeon/Shanghai/S1069/2013 2013/04/02 HA 525339091	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTITFSFNGAFIAPDRASF LRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPE IPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQ FELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLREN AEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASC FILLAIVMGLVFICVKNGNMRCTICI
2339	AGR49770 A/wild pigeon/Jiangsu/SD001/2013 2013/04/17 HA 525339151	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI

2340	AGY41893 A/Huizhou/01/2013 2013/08/08 HA 552049496	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2341	AGY42258 A/mallard/Sweden/91/2002 2002/12/12 HA 552052155	FALVAIIPINADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSRGKRTVDLGQCGLL GTIXGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYSGIRTNGA XSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRNDPALIIWGIHHSGSTTEQTKLYGSGNK LITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDRASFLRGKSMGI QSGVQIDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLLLATGMKNVPEIPKGRGLFG AIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFT EVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLRENAEEDGTG CFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMG LVFMCVKNGNMRCTICI
2342	AHA11441 A/guinea fowl/Nebraska/17096/2011 2011/04/10 HA 557478572	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRPTDL GQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRNKPALIVWGVHHSGSATEQT KLYGSGSKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRAS FFRGESLGVQSDVPLDSGCEGDCFHKGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNVP ENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEIWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQLR ENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESLQNRIQIDPVKLSSGYKDIILWFSFGAS CFLLLAIAMGLVFICIKNGNMRCTICI
2343	AHA11452 A/turkey/Minnesota/32710/2011 2011/07/12 HA 557478591	MNTQILALIACMLVGTKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRPTD LGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEEPLRQILRGSGGIDKESMGFTY SGIRTNGATSTCRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRNKPALIVWGVHHSGSATEQ TKLYGSGSKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRA SFFRGESLGVQSDVPLDSGCEGDCFHKGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNV PENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QQFEMIDNEFSEIEQQIGNVINWTRDSMTEIWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQLR ENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESLQNRIQIDPVKLSSGYKDIILWFSFGAS CFLLLAIAMGLVFICIKNGNMRCTICI
2344	AHA11461 A/turkey/Minnesota/31900/2011 2011/07/05 HA 557478606	MNTQILALIACMLVGTKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRPTD LGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEEPLRQILRGSGGIDKESMGFTY SGIRTNGATSTCRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRNKPALIVWGVHHSGSATEQ TKLYGSGSKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRA SFFRGESLGVQSDVPLDSGCEGDCFHKGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNV PENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEIWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQLR ENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESLQNRIQIDPVKLSSGYKDIILWFSFGAS CFLLLAIAMGLVFICIKNGNMRCTICI
2345	AHK10585 A/chicken/Guangdong/G1/2013 2013/05/05 HA 587680636	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI

2346	AGG53366 A/wild duck/Korea/CSM42-34/2011 2011/03/ HA 459252887	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVD LGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGLTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGIHHSGSSTEQ TKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVP ELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVRLSSGYKDVILWFSFGA SCFILLAIAMGLVFICVKNGNMRCTICI
2347	AGG53377 A/wild duck/Korea/CSM42-1/2011 2011/03/ HA 459252925	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVD LGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGLTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGIHHSGSSTEQ TKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVP ELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVRLSSGYKDVILWFSFGA SCFILLAIAMGLVFICVKNGNMRCT
2348	AGG53399 A/wild duck/Korea/MHC39-26/2011 2011/03/ HA 459253005	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVD LGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGIHHSGSTTEQ TKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVP EPPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTN QQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQL RENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFG ASCFILLAIAMGLVFICVKNGNMRCTICI
2349	AGG53432 A/wild duck/Korea/MHC35-41/2011 2011/03/ HA 459253136	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVD LGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGIHHSGSTTEQ TKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVP EPPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTN QQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQL RENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFG ASCFILLAIAMGLVFICVKNGNMRCT
2350	AGG53476 A/wild duck/Korea/SH19-27/2010 2010/12/ HA 459253257	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVD LGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGIHHSGSTTEQ TKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVP ELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGA SCFILLAIAMGLVFICVKNGNMRCTI
2351	AGG53487 A/wild duck/Korea/SH19-50/2010 2010/01/ HA 459253278	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVD LGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGIHHSGSTTEQ TKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVP ELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGA SCFILLAIAMGLVFICVKNGNMRCTICI

2352	AGG53520 A/wild duck/Korea/SH20-27/2008 2008/12/ HA 459253409	QILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVDLGQ CGLLGTITGPPQCDQLLEFSADLIIERREGTDVCYPGKFVNEEALRQILRESGGIEKETMGFTYSGIR TNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHHSGSTTEQTKLYG SGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRG KSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLMLATGMKNVPELPK GRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFE LIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVKRQLRENA EEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQINPVKLSSGYKDVILWFSFGASCFI LLAIAMGLVFICVKNGNMR
2353	AGL43637 A/Taiwan/1/2013 2013// HA 496297389	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGPSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIINNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2354	AGL97639 A/mallard/Minnesota/AI09-3770/2009 2009/09/12 HA 505555371	IACMLVGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKLCTQGKRPTDLGQCGLL GTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYSGIRTNGA TSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHSGSATEQTKLYGSGN KLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRASFFRGESL GVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNVPENPKTR GLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTNQQFELID NEFSEIEQQIGNVINWTRDSMTELWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQLRENAEED GTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQNRIQIDPVKLS
2355	AGO02477 A/Xuzhou/1/2013 2013/04/25 HA 512403688	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGSKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKSRNMRCTICI
2356	AGR84942 A/Suzhou/5/2013 2013/04/12 HA 526304561	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGSKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2357	AGR84954 A/Nanjing/6/2013 2013/04/11 HA 526304594	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNRNMRCTICI

2358	AGR84978 A/Wuxi/4/2013 2013/04/07 HA 526304656	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKSRNMRCTICI
2359	AGR84990 A/Wuxi/3/2013 2013/04/07 HA 526304688	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKSRNMRCTICI
2360	AGR85002 A/Zhenjiang/1/2013 2013/04/07 HA 526304708	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKSRNKRCTICI
2361	AGR85026 A/Nanjing/2/2013 2013/04/05 HA 526304762	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKSRNMRCTICI
2362	AGU02230 A/Zhejiang/DTID-ZJU05/2013 2013/04/ HA 532808765	LVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGGEVVNATETVERTNIPRICSKGKRTVDLGQCGL RGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGIRTN GATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLYGS GNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRG KSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEIPK GRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFE LIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAE EDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFIL LAIVMGLVFICVKNGNMRCT
2363	AGU02233 A/Zhejiang/DTID-ZJU08/2013 2013/04/ HA 532808788	FALIAIIPTNADKICLGHHAVSNGTKVNTLTERGGEVVNATETVERTNFPRICSKGKRTVDLGQCGLR GTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGIRTNGA TSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLYGSGN KLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRGKS MGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEIPKGR GLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELID NEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEED GTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAI VMGLVFICVKNGNMRCT

2364	AGW82588 A/tree sparrow/Shanghai/01/2013 2013/05/09 HA 546235348	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTIGI
2365	AGW82600 A/Shanghai/CN01/2013 2013/04/11 HA 546235368	ALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDLGQCGLLGT ITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGIRTNGATS ACRRSRSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLYGSGNKL VTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRGKSMGI QSGVQVDANCEGDCYHSGGTIMSNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEIPKGRGL FGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNE FNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGT GCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVM GLVFICVKNGNMRCTICI
2366	AGW82612 A/Shanghai/JS01/2013 2013/04/03 HA 546235388	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKNPALIVWGIHHSGSTAEQT KLYGSGNKLVTVGSSNYQQSFAPSPGARTQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRA SFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNV PEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTN QQFELIDNEFTEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGA SCFILLAIAMGLVFICVKNGNMRCTICI
2367	AHA11472 A/turkey/Minnesota/31676/2009 2009/12/08 HA 557478625	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANVKKICTQGKRPTD LGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTY SGIRTNGETSACRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRDKPALIIWGVHHSGSATEQ TKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRA SFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNV PEKPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITNKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQL RENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRKESLQNRIQIDPVKLSSGYKDIILWFSFGA SCFLLLAIAMGLVFICIKNGNMRCTICI
2368	AHA11483 A/turkey/Minnesota/14135-2/2009 2009/08/07 HA 557478644	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANVKKICTQGKRPTD LGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRDKPALIIWGVHHSGSATEQ TKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRA SFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNV PEKPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITSKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQL RENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRKESLQNRIQIDPVKLSSGYKDIILWFSFGA SCFLLLAIAMGLVFICIKNGNMRCTICI
2369	AHA11500 A/Zhejiang/DTID-ZJU10/2013 2013/10/14 HA 557478676	TQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDLGQ CGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGIR TNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLY GSGNKLVTVGSSNYQQSFVPSPGARPPVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFL RGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEI PKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQ FELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLREN AEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASC FILLAIVMGLVFICVKN

2370	AHA57050 A/turkey/Minnesota/14659/2009 2009/08/12 HA 558484427	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANVKKICTQGKRPTD LGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRDKPALIIWGVHHSGSATEQ TKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRA SFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNV PEKPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITSKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQL RENAEEDGTGCFEIFHNCDDQCMESIRNNTYDHTQYRKESLQNRIQIDPVKLSSGYKDIILWFSFGA SCFLLLAIAMGLVFICIKNGNMRCTICI
2371	AHA57072 A/turkey/Minnesota/18421/2009 2009/09/09 HA 558484465	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANVKKICTQGKRPTD LGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNSNDAAFPQMTKSYRNPRDKPALIIWGVHHSGSATEQ TKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFIAPDRA SFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLLLATGMRNV PEKPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIDKTN QQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRKQL RENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRKESLQNRIQIDPVKLSSGYKDIILWFSFGA SCFLLLAIAMGLVFICIKNGNMRCTICI
2372	AHD25003 A/Guangdong/02/2013 2013/10/ HA 568260567	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNM
2373	AHF20528 A/Hong Kong/470129/2013 2013/11/30 HA 570933555	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISSLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2374	AHF20568 A/Shanghai/CN02/2013 2013/04/02 HA 570933626	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIMSNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNV PEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTN QQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLR ENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGA SCFILLAIVMGLVFICVKNGNMRCTICI
2375	AHH25185 A/Guangdong/04/2013 2013/12/16 HA 576106234	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIEKEAMGFTYS GIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI

2376	AHJ57411 A/Shanghai/PD-01/2014 2014/01/17 HA 585478041	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVSSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCKGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRIIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2377	AHJ57418 A/Shanghai/PD-02/2014 2014/01/17 HA 585478256	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDICYPGKFVNEEALRQILRESGGIDKEAMGFTYSG IRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKL YGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASF LKGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPE IPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRIIEKTNQQ FELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLREN AEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASC FILLAIVMGLVFICVKNGNMRCTICI
2378	AHK10800 A/Shanghai/01/2014 2014/01/03 HA 587681014	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRIIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2379	AHM24224 A/Beijing/3/2013 2013/04/16 HA 594704802	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVKEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2380	AHN96472 A/chicken/Shanghai/PD-CN-02/2014 2014/01/21 HA 602701641	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQKSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2381	AHZ39686 A/Anhui/DEWH72-01/2013 2013// HA 632807036	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDDAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI

2382	AHZ39710 A/Anhui/DEWH72-03/2013 2013// HA 632807076	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTDGATSACRRSGSSFYAEMKWLLSNTDDAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2383	AHZ39746 A/Anhui/DEWH72-06/2013 2013// HA 632807136	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTK LYGSGNKLVTVGSSNYQQSFVPSPGERPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRAS FLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVP EIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQ QFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRE NAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGAS CFILLAIVMGLVFICVKNGNMRCTICI
2384	AHZ41929 A/mallard/Sweden/1621/2002 2002/12/12 HA 632810949	MNTQILVFALVAIIPINADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSRGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRNDPALIIWGIHHSGSTTEQTK LYGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIAPDRASFL RGKSMGIQSGVQIDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLLLATGMKNVPEIP KGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQF ELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSEMNKLYERVRRQLREN AEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASC FILLAIAMGLVFMCVKNGNMRCTICI
2385	AHZ42537 A/mallard/Minnesota/AI09-3770/2009 2009/09/12 HA 632811964	MNTQILAFIACMLVGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKLCTQGK RPTDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDK ESMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWG VHHSGSATEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTV TFTFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCP RYVKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADY KSTQSAIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTELWSYNAELLVAME NQHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTE SLQNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2386	AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000 2000/05/20 HA 632811984	MNTQILAFIACMLVGVRGDKICLGHHAVANGTKVNTLTEKGIEVVNATETVESANIKKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFDSDLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKE SMGFTYSGIRTNGATSACRRLGSSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWG VHHSGSANEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTV TFTFNGAFIAPDRASFFRGESLGIQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGKCPR YVKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIDKTNQQFELMDNEFNEIEQQIGNVINWTRDSMTEVWSYNAELLVAME NQHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTE SLQNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLIFICIKNGNMRCTICI
2387	AID70634 A/Shanghai/Mix1/2014 2014/01/03 HA 660304650	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRIIEKTNQQFELIDNEFNEVEKQISNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2388	AIN76383 A/Zhejiang/LS01/2014 2014/02/08 HA 684694637	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGTTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2368	AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013 2013/12/ HA 699978931	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVEVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2369	AIU47013 A/chicken/Suzhou/040201H/2013 2013/04/ HA 699979673	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDMILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2370	AJJ90490 A/chicken/Shenzhen/742/2013 2013/12/10 HA 755178094	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2371	AJJ90526 A/chicken/Shenzhen/898/2013 2013/12/09 HA 755178154	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDICYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRANGATSACKRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISSLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQIDPVKLSRGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2372	AJJ90538 A/silkie chicken/Shenzhen/918/2013 2013/12/09 HA 755178174	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2373	AJJ90576 A/chicken/Shenzhen/1665/2013 2013/12/12 HA 755178238	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDICYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRANGATSACKRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQIDPVKLSRGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2374	AJJ90588 A/chicken/Shenzhen/2110/2013 2013/12/13 HA 755178258	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSIGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2375	AJJ90661 A/chicken/Dongguan/2912/2013 2013/12/18 HA 755178380	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2376	AJJ90673 A/silkie chicken/Dongguan/3049/2013 2013/12/18 HA 755178400	MNTQILVFALTAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2377	AJJ90795 A/silkie chicken/Dongguan/3281/2013 2013/12/18 HA 755178604	MNTQILVFALIAIIPTNADKICLGHHAVPNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2378	AJJ90891 A/silkie chicken/Dongguan/3520/2013 2013/12/19 HA 755178764	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKXPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2379	AJJ90951 A/chicken/Dongguan/3544/2013 2013/12/19 HA 755178864	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYRNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2380	AJJ91035 A/chicken/Shenzhen/3780/2013 2013/12/19 HA 755179004	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDNRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2381	AJJ91155 A/chicken/Dongguan/4037/2013 2013/12/19 HA 755179204	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2382	AJJ92005 A/chicken/Shenzhen/801/2013 2013/12/09 HA 755180629	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSRGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2383	AJJ94254 A/chicken/Dongguan/1374/2014 2014/02/21 HA 755184382	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2384	AJJ94606 A/chicken/Dongguan/191/2014 2014/02/20 HA 755184968	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2385	AJJ96552 A/chicken/Jiangxi/12206 /2014 2014/03/16 HA 755188219	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTI DLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHNKYREEAMQN RIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2386	AJJ96684 A/chicken/Jiangxi/13207 /2014 2014/03/30 HA 755188439	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKINTLTERGVEVVNATETVERTNIPRICSKGKKTV DLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2387	AJJ96732 A/chicken/Jiangxi/13223 /2014 2014/03/30 HA 755188519	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2388	AJK00354 A/duck/Zhejiang/LS02/2 014 2014/01/12 HA 755194469	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIVERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHH SGSTTEQTKLYGSGNKLITVGSSNYQQSFVPSPGARPLVNGQSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYV KQESLLLATGMKNVPEVPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQVTGKLNRLIEKTNQQFELIDHEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQ HTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2389	AJJ91264 A/silkie chicken/Dongguan/4129 /2013 2013/12/19 HA 755179386	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLMEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2390	AJJ91314 A/chicken/Shaoxing/241 7/2013 2013/10/20 HA 755179470	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPPVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2391	AJJ91402 A/chicken/Huzhou/4045/ 2013 2013/10/24 HA 755179618	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKEVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2392	AJJ91476 A/chicken/Huzhou/4076/ 2013 2013/10/24 HA 755179743	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSRGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2393	AJJ91725 A/chicken/Shaoxing/520 1/2013 2013/10/28 HA 755180161	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2394	AJJ91885 A/Shenzhen/SP4/2014 2014/01/16 HA 755180429	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGVTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSRGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2395	AJJ91909 A/Shenzhen/SP26/2014 2014/01/20 HA 755180469	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDICYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRANGATSACKRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISSLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDGCMASIRNNTYDHSKYREEAMQN RIQIDPVKLSRGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2396	AJJ91945 A/Shenzhen/SP38/2014 2014/01/22 HA 755180529	MNTQILAFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIGGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2397	AJJ91957 A/Shenzhen/SP44/2014 2014/01/23 HA 755180549	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGTTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISSLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2398	AJJ91969 A/Shenzhen/SP48/2014 2014/01/23 HA 755180569	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2399	AJJ91993 A/chicken/Dongguan/4119/2013 2013/12/19 HA 755180609	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLLGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFTLLAIVMGLVFICVKNGNMRCTICI
2400	AJJ92031 A/chicken/Dongguan/4064/2013 2013/12/19 HA 755180672	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVESSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2401	AJJ92967 A/silkie chicken/Jiangxi/9469/2014 2014/02/16 HA 755182232	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGVTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2402	AJJ93027 A/chicken/Jiangxi/9558/2014 2014/02/16 HA 755182332	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVKEEALRQILRESGGIDKEA MGFTYSGIRTNGVTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2403	AJJ93051 A/chicken/Jiangxi/10573/2014 2014/02/18 HA 755182372	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGVTSACRRSGSSFYAEMKWLLSNTDDAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2404	AJJ93845 A/silkie chicken/Dongguan/157/2014 2014/02/20 HA 755183695	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2405	AJJ93857 A/chicken/Dongguan/169/2014 2014/02/20 HA 755183715	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACMRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2406	AJJ93869 A/chicken/Dongguan/173/2014 2014/02/20 HA 755183735	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTVTGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2407	AJJ93881 A/chicken/Dongguan/189/2014 2014/02/20 HA 755183755	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTVTGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPKY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2408	AJJ93907 A/chicken/Dongguan/449/2014 2014/02/20 HA 755183799	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2409	AJJ93931 A/chicken/Dongguan/536/2014 2014/02/20 HA 755183839	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISKLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2410	AJJ93943 A/chicken/Dongguan/568/2014 2014/02/20 HA 755183859	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIEKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSGGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2411	AJJ93979 A/silkie chicken/Dongguan/656/2014 2014/02/20 HA 755183919	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTVTGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFGLIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2412	AJJ94134 A/chicken/Dongguan/1051/2014 2014/02/21 HA 755184182	MNTQILVLALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVXLSXGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2413	AJJ94158 A/chicken/Dongguan/1075/2014 2014/02/21 HA 755184222	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYRGEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2414	AJJ94182 A/chicken/Dongguan/1177/2014 2014/02/21 HA 755184262	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACKRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSIAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2415	AJJ94194 A/silkie chicken/Dongguan/1264/2014 2014/02/21 HA 755184282	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTI DLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKSTQ SAIDQVTGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYRGEAMQ
2416	AJJ94206 A/silkie chicken/Dongguan/1268/2014 2014/02/21 HA 755184302	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISDLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM
2417	AJJ94344 A/silkie chicken/Dongguan/1451/2014 2014/02/21 HA 755184532	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNSTETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRTVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ
2418	AJJ94356 A/chicken/Dongguan/1456/2014 2014/02/21 HA 755184552	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM
2419	AJJ94396 A/chicken/Dongguan/1494/2014 2014/02/21 HA 755184618	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPETPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM
2420	AJJ94754 A/chicken/Dongguan/748/2014 2014/02/20 HA 755185215	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIEKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSNAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM

2421	AJJ94838 A/chicken/Dongguan/835/2014 2014/02/20 HA 755185356	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSASTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFGFGASCFILLAIVMGLVFICVKNGNMRCTICI
2422	AJJ94862 A/chicken/Dongguan/843/2014 2014/02/20 HA 755185396	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIEKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSGGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2423	AJJ94886 A/chicken/Dongguan/851/2014 2014/02/20 HA 755185436	MNTQILAFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2424	AJJ94910 A/chicken/Dongguan/874/2014 2014/02/20 HA 755185476	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSASTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2425	AJJ94959 A/silkie chicken/Dongguan/967/2014 2014/02/21 HA 755185558	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACXRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2426	AJJ95048 A/chicken/Dongguan/1009/2014 2014/02/21 HA 755185708	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPETPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2427	AJJ95171 A/chicken/Dongguan/1314/2014 2014/02/21 HA 755185913	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF NFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2428	AJJ95227 A/chicken/Dongguan/1382/2014 2014/02/21 HA 755186006	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDICYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2429	AJJ95251 A/chicken/Dongguan/1401/2014 2014/02/21 HA 755186046	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYKRVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2430	AJJ95346 A/chicken/Dongguan/1548/2014 2014/02/21 HA 755186206	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYKRVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHNKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2431	AJJ95382 A/chicken/Dongguan/1690/2014 2014/02/21 HA 755186266	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSIGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2432	AJJ95464 A/chicken/Shenzhen/138/2014 2014/02/19 HA 755186404	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYRGEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFMLLAIVMGLVFICVKNGNMRCTICI

2433	AJJ95572 A/chicken/Dongguan/1100/2014 2014/02/21 HA 755186584	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIEKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSGGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2434	AJJ95584 A/silkie chicken/Dongguan/1519/2014 2014/02/21 HA 755186604	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYRGEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFMLLAIVMGLVFICVKNGNMRCTICI
2435	AJJ95596 A/Shenzhen/SP58/2014 2014/01/25 HA 755186624	MNTQILAFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2436	AJJ95620 A/Shenzhen/SP75/2014 2014/02/15 HA 755186664	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGSTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAVVMGLVFICVKNGNMRCTICI
2437	AJJ95632 A/Shenzhen/SP62/2014 2014/02/05 HA 755186684	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNATFPQMTKSYKNTRKSPALIIWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2438	AJJ96720 A/chicken/Jiangxi/13220/2014 2014/03/30 HA 755188499	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTTIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSRGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2439	AJJ96817 A/chicken/Jiangxi/9513/2014 2014/02/16 HA 755188661	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATEIVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGVTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2440	AJJ96841 A/Shenzhen/SP139/2014 2014/04/02 HA 755188701	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSTCRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRACFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVERQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2441	AJJ96889 A/chicken/Jiangxi/13496/2014 2014/04/11 HA 755188781	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTXIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKXA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSXGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2442	AJJ96901 A/chicken/Jiangxi/13502/2014 2014/04/11 HA 755188801	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSXGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2443	AJJ96925 A/chicken/Jiangxi/13513/2014 2014/04/11 HA 755188841	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYNGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH TVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDLHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2444	AJJ97267 A/chicken/Jiangxi/13252/2014 2014/03/30 HA 755189411	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2445	AJJ97291 A/chicken/Jiangxi/13493 /2014 2014/04/06 HA 755189451	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYNGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ
2446	AJJ97331 A/chicken/Jiangxi/13512 /2014 2014/04/06 HA 755189517	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYNGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSIGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQN RIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2447	AJJ97373 A/chicken/Jiangxi/13521 /2014 2014/04/06 HA 755189587	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYNGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPXRASFLRGKSXGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2448	AJJ97443 A/chicken/Jiangxi/13530 /2014 2014/04/06 HA 755189702	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTTIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSRGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2449	AJJ97582 A/chicken/Jiangxi/14023 /2014 2014/04/13 HA 755189933	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2450	AJJ97697 A/chicken/Jiangxi/14517 /2014 2014/04/20 HA 755190125	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCDGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2451	AJJ97709 A/chicken/Jiangxi/14518 /2014 2014/04/20 HA 755190145	MNTQILVFALIAIIPANADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYNGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGNCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ
2452	AJJ97745 A/chicken/Jiangxi/14554 /2014 2014/04/20 HA 755190205	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELMDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2453	AJJ97757 A/chicken/Shantou/2537 /2014 2014/04/16 HA 755190225	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2454	AJJ97841 A/duck/Jiangxi/15044/2 014 2014/04/27 HA 755190365	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVRLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2455	AJJ97899 A/chicken/Jiangxi/15524 /2014 2014/05/05 HA 755190462	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFMCVKNGNMRCTICI
2456	AJJ97925 A/silkie chicken/Shantou/2050/2 014 2014/03/25 HA 755190506	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEVPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2457	AJJ97973 A/chicken/Shantou/4325/2014 2014/07/01 HA 755190586	MNTQILVFALISIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGVTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEVPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2458	AJJ97998 A/chicken/Shantou/4816/2014 2014/07/22 HA 755190628	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELVDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

Table 14: H10 Hemagglutinin Amino Acid Sequences

SEQ ID NO	Accession No / Strain / Protein	Amino Acid Sequence
2459	AAM19228 A/turkey/Minnesota/38429/1988 1988// HA 20335017	ACVLVEAKGDKICLGHHAVVNGTKVNTLTEKGIEVVNATETVETANIGKICTQGKRPTDLGQCG LLGTLIGPPQCDQFLEFESDLIIERREGNDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYSGI ITNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIVWGIHHSGSTTEQ TKLYGSGNKLITVESSKYQQSFTPSPGARPQVNGESGRIDFHWMLLDPNDTVTFTFNGAFIAP DRASFFKGESLGVQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQPSLLLA TGMRNVPENPKTRGLFGAIAGFIEKDGGSHYG
2460	AAY46211 A/mallard/Sweden/91/2002 2002// HA 66394828	MNTQILVFALVAIIPINADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSRGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKE TMGFTYSGIRTNGAPSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRNDPALIIWGIH HSGSTTEQTKLYGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQIDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVK QESLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQH TIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFMCVKNGNMRCTICI
2461	ABI84694 A/turkey/Minnesota/1/1988 1988/07/13 HA 115278573	MNTQILVFIACVLVEAKGDKICLGHHAVVNGTKVNTLTEKGIEVVNATETVETANIGKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFESDLIIERREGNDVCYPGKFTNEESLRQILRGSGGIDKE SMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIVWGI HHSGSTTEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWMLLDPNDTVT FTFNGAFIAPDRASFFKGESLGVQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGKCPR YVKQPSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYK STQSAIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHAQYRAES LQNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2462	ABS89409 A/blue-winged teal/Ohio/566/2006 2006// HA 155016324	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRP TDLGQCGLLGTLIGPPQCDQFLEFDTDLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKES MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVH HSGSATEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTF TFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRY VKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESL QNRIQIDPVRLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI

2463	ACD03594 A/ruddy turnstone/DE/1538/2000 2000// HA 187384848	MNTQILAFIACMLVGVRGDKICLGHHAVANGTKVNTLTEKGIEVVNATETVESANIKKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFDSDLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKE SMGFTYSGIRTNGATSACRRLGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGV HHSGSANEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVT FTFNGAFIAPDRASFFRGESLGIQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGKCPRY VKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIDKTNQQFELMDNEFNEIEQQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESL QNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLIFICIKNGNMRCTICI
2464	BAH22785 A/duck/Mongolia/119/2008 2008// HA 223717820	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIGKET MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHH SGSTTEQTKLYGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHNGGTIISNLPFQNINSRTVGKCPRYV KQESLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIERTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQ HTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSNGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2465	CAY39406 A/Anas crecca/Spain/1460/2008 2008/01/26 HA 254674376	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKE TMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIH HSGSTTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRY VKQESLMLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2466	ACX53683 A/goose/Czech Republic/1848-K9/2009 2009/02/04 HA 260907763	MNIQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERRGGSDVCYPGKFVNEEALRQILRESGGIDKE TMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIH HSGSTTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLKGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRY VKQESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQINPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2467	ACZ48625 A/turkey/Minnesota/38429/1988 1988// HA 269826341	MNTQILVFIACVLVEAKGDKICLGHHAVVNGTKVNTLTEKGIEVVNATETVETANIGKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFESDLIIERREGNDVCYPGKFTNEESLRQILRGSGGIDKE SMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIVWGI HHSGSTTEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWMLLDPNDTVT FTFNGAFIAPDRASFFKGESLGVQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGKCPR YVKQPSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYK STQSAIDQITGKLNRLIDKTNQQFEL
2468	ADC29485 A/mallard/Spain/08.00991.3/2005 2005/11/ HA 284927336	STQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAME NQHTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREE AMQNRIQIDPVKLSSGYKDVILWFSFGASCFILL
2469	ADK71137 A/blue-winged teal/Guatemala/CIP049-01/2008 2008/02/07 HA 301333785	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRP TDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKES MGFTYSGIRTNGATSACRRSGSSSYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVH HSGSATEQTKLYGSGNKLITVGSSKYQQSFTPSPGTRPQVNGQSGRIDFHWLLLDPNDTVTFT FNGAFIAPDRASFLRGKSLGIQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVK QTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIDKTNQHFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHT IDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQN RIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI

2470	ADK71148 A/blue-winged teal/Guatemala/CIP049-02/2008 2008/03/05 HA 301333804	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNXTETVETANIKKICTHGKRP TDLGQCGLLGTLIGPPQCDRFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKES MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVH HSGSATEQTKLYGSGNKLITVGSSKYQQSFTPSPGTRPQVNGQSGRIDFHWLLLDPNDTVTFT FNGAFIAPDRASFLRGKSLGIQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVK QTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHT IDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQN RIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2471	ADN34727 A/goose/Czech Republic/1848-T14/2009 2009/02/04 HA 307141869	MNIQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERRGGSDVCYPGKFVNEEALRQILRESGGIDKE TMGFTYSGIRTNGXTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIH HSGSTTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLKGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRY VKQESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQINPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2472	AEK84760 A/wild bird/Korea/A14/2011 2011/02/ HA 341610308	PAFIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQ ESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQS AIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTI DLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQN RIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2473	AEK84761 A/wild bird/Korea/A3/2011 2011/02/ HA 341610310	ILVFALVAIIPTNANKIGLGHHAVSNGTKVNTLTERGVEVFNATETVERTNVPRICSKGKKTVDL GQCGLRGTITGPPQCDQFLKFSPDLIIERQKGSDVCYPGKFVNEKPLRQILRESGGIDKETMGF AYNGIKTNGPPIACRKSGSSFYAKMKWLLSNTDKAAFPQMTKSYKNTRRNPALIVWGIHHSGS TTKQTKLYGIGSNLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGA FIPPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQES LMLATGMKNVPELPKGKGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAI DQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTID LADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRI QIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2474	AEK84763 A/wild bird/Korea/A9/2011 2011/02/ HA 341610314	ILVFALVAIIPTNANKIGLGHHAVSNGTKVNTLTERGVEFFNATETVEPTNVPRICSKGKKTVDLG QCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEKALRQILRESGGIDKETMGFA YSGIKTNGPPIACRKSGSSFYAKMKWLLSNTDKAAFPQMTKSYKNIRRDPALIVWGIHHSGSTT KQTNLYGIGSNLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFIFNGAFIA PDRASFLIGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLML ATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQIT GKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADS EMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDP VKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2475	AEK84765 A/spot-billed duck/Korea/447/2011 2011/04/ HA 341610318	LVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVDLG QCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFT YSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGIHHSGS TTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGA FIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQES LMLATGMKNVPEPPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAI DQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTID LADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMARIRNNTYDHSKYREEAMQNRI QIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI

2476	AEM98291 A/wild duck/Mongolia/1-241/2008 2008/04/ HA 344196120	SILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVD LGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMG FTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHHSG STTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFN GAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGSIISNLPFQNINSRAVGKCPRYVKQ ESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQS AIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTI DLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQINPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTI
2477	AFM09439 A/emperor goose/Alaska/44063-061/2006 2006/05/23 HA 390535062	QILAFIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETVNIKKICTQGKRPTDL GQCGLLGTLIGPPQCDQFLEFDADLIIERRKGTDVCYPGKFTNEESLRQILRGSGGIDKESMGF TYSGIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHSG SATEQTKLYGSGNKLITVGSSKYQQSFVPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFN GAFIAPERASFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQ TSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQS AIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQHTI DLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQNR IQINPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2478	AFV33945 A/guinea fowl/Nebraska/17096-1/2011 2011/04/05 HA 409676820	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERRIEVVNATETVETANIKKICTQGKRP TDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKES MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRNKPALIVWGV HHSGSATEQTKLYGSGSKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVT FTFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHKGGTIVSSLPFQNINPRTVGKCPR YVKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEIWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESL QNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2479	AFV33947 A/goose/Nebraska/17097-4/2011 2011/04/05 HA 409676827	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRP TDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKES MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIVWGV HHSASATEQTKLYGSGSKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTF TFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHKGGTIVSSLPFQNINPRTVGKCPRY VKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEIWSYNAELLVAMENQ HTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESLQ NRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2480	AFX85260 A/ruddy turnstone/Delaware Bay/220/1995 1995/05/21 HA 423514912	MNTQILAFIACMLIGINGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKRICTQGKRP IDLGQCGLLGTLIGPPQCDQFLEFDSDLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESM GFTYSGIRTNGATSACIRLGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHS GSANEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTF NGAFIAPDRASFFRGESLGVQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGRCPRYVK QTSLLLATGMKNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIDKTNQQFELIDNEFNEIEQQIGNVINWTRDSMTEVWSYNAELLVAMENQH TIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQN RIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2481	AGE08098 A/northern shoverl/Mississippi/11OS145/2011 2011/01/08 HA 444344488	MNTQILTLIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRP TDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKES MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVH HSGSATEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTF TFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHNGGTIVSSLPFQNINPRTVGKCPRY VKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESL QNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI

2482	AGI60301 A/Hangzhou/1/2013 2013/03/24 HA 475662454	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGISGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2483	AGI60292 A/Shanghai/4664T/2013 2013/03/05 HA 476403560	MNTQILVFALIAIIPANADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCHHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2484	AGJ72861 A/chicken/Zhejiang/DTI D-ZJU01/2013 2013/04/ HA 479280294	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGGEVVNATETVERTNIPRICSKGKKT VDLGQGGPRGTITGPPQCDQFLEFSADLIMERREGSDVCYPGKFVNEEALRQILRESGGIDKE AMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGI HHSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTV TFSFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCP RYVKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADY KSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAME NQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREE AMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2485	AGJ73503 A/Nanjing/1/2013 2013/03/28 HA 479285761	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2486	BAN16711 A/duck/Gunma/466/201 1 2011// HA 482661571	MNTQVLVFALMAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGK RTVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDK ETMGFTYSGIRTNGTTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIAWG IHHSGSTTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVT FSFNGAFIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPR YVKQESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAME NQHTIDLADSEMNKLYERVRRQLRENAEEDDTGCFEIFHKCDDDCMASIRNNTYDHSKYREE AMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2487	AGK84857 A/Hangzhou/2/2013 2013/04/01 HA 485649824	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQITKSYKNTRKSPALIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2488	AGL44438 A/Shanghai/02/2013 2013/03/05 HA 496493389	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2489	AGL33692 A/Shanghai/4655T/2013 2013/02/26 HA 491874175	GMIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGN VINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFH KCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLV FICVKNGNMRCTICI
2490	AGL33693 A/Shanghai/4659T/2013 2013/02/27 HA 491874186	GMIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIG NVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIF HKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGL VFICVKNGNMRCTICI
2491	AGL95088 A/Taiwan/S02076/2013 2013/04/22 HA 501485301	VFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDLGQ CGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTA EQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAF IAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSL LLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQ ITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLAD SEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQID PVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMR
2492	AGL95098 A/Taiwan/T02081/2013 2013/04/22 HA 501485319	LVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDLGQ CGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTA EQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAF IAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSL LLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQ ITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLAD SEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQID PVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCT
2493	AGM53883 A/Shanghai/5083T/2013 2013/04/20 HA 507593986	GFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRD SITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCM ASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGN MRCT
2494	AGM53884 A/Shanghai/5180T/2013 2013/04/23 HA 507593988	AQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWS YNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNT YDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2495	AGM53885 A/Shanghai/5240T/2013 2013/04/25 HA 507593990	QNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEV WSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIR NNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRC T
2496	AGM53886 A/Shanghai/4842T/2013 2013/04/13 HA 507593992	NAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVW SYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNN TYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCT
2497	AGM53887 A/Shanghai/4701T/2013 2013/04/06 HA 507593994	NAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVW SYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNN TYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTIC

2498	AGN69462 A/Wuxi/2/2013 2013/03/31 HA 511105778	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGSTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGSKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2499	AGN69474 A/Wuxi/1/2013 2013/03/31 HA 511105798	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLINGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2500	AGO51387 A/Jiangsu/2/2013 2013/04/20 HA 514390990	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYRXEAM XBXIQIDPVKLSSGYKDVXJWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2501	BAN59726 A/duck/Mongolia/147/2008 2008/08/29 HA 519661951	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIGKET MGFTYSGIRTNGATSACRRSRSSFYAEMKWLLSNTDNAAFPQMTRSYKNTRKDPALIIWGIHH SGSTTEQTKLYGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHNGGTIISNLPFQNINSRTVGKCPRYV KQESLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIERTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQ HTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSNGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2502	BAN59727 A/duck/Mongolia/129/2010 2010// HA 519661954	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERINVPRICSKGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKE TMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIH HSGSTTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRY VKQESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQINPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2503	AGQ80952 A/duck/Jiangxi/3096/2009 2009// HA 523788794	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTSIPRICSKGKRA VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKET MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQTTKSYKNTRKDPALIIWGIHH SGSTTEQTKLYGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHNGGTIISNLPFQNINSRAVGKCPRYV KQESLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFTEVERQIGNVINWTRDSMTEVWSYNAELLVAMENQ HTIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI

2504	AGQ80989 A/duck/Jiangxi/3257/2009 2009// HA 523788868	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTSIPRICSKGKRA VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKET MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQTTKSYKNTRKDPALIIWGIHH SGSTTEQTKLYGSGNKLITVGXSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHNGGTIISNLPFQNINSRAVGKCPRYV KQESLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFTEVERQIGNVINWTRDSMTEVWSYNAELLVAMENQ HTIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2505	AGQ81043 A/chicken/Rizhao/515/2013 2013// HA 523788976	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEE MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2506	AGR33894 A/chicken/Rizhao/719b/2013 2013// HA 524845213	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDRSKYREEAM QNRXXXXXXXXXXXXKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2507	AGR49399 A/chicken/Jiangxi/SD001/2013 2013/05/03 HA 525338528	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2508	AGR49495 A/chicken/Shanghai/S1358/2013 2013/04/03 HA 525338689	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIKNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2509	AGR49506 A/chicken/Shanghai/S1410/2013 2013/04/03 HA 525338708	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVT FSFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPR YVKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2510	AGR49554 A/chicken/Zhejiang/SD033/2013 2013/04/11 HA 525338789	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2511	AGR49566 A/duck/Anhui/SC702/2013 2013/04/16 HA 525338809	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDNRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2512	AGR49722 A/homing pigeon/Jiangsu/SD184/2013 2013/04/20 HA 525339071	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSEIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVT FSFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPR YVKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2513	AGR49734 A/pigeon/Shanghai/S1069/2013 2013/04/02 HA 525339091	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTITF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2514	AGR49770 A/wild pigeon/Jiangsu/SD001/2013 2013/04/17 HA 525339151	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVT FSFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPR YVKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2515	AGY41893 A/Huizhou/01/2013 2013/08/08 HA 552049496	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2516	AGY42258 A/mallard/Sweden/91/2002 2002/12/12 HA 552052155	FALVAIIPINADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSRGKRTVDLGQC GLLGTIXGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKETMGFTYS GIRTNGAXSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRNDPALIIWGIHHSGSTTE QTKLYGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFSFNGAFIA PDRASFLRGKSMGIQSGVQIDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQESLLLA TGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITG KLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTIDLADSE MNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPV KLSSGYKDVILWFSFGASCFILLAIAMGLVFMCVKNGNMRCTICI
2517	AHA11441 A/guinea fowl/Nebraska/17096/2011 2011/04/10 HA 557478572	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKRP TDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKES MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRNKPALIVWGV HHSGSATEQTKLYGSGSKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVT FTFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHKGGTIVSSLPFQNINPRTVGKCPR YVKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIDKTNQQFELINNEFSEIEQQIGNVINWTRDSMTEIWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESL QNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2518	AHA11452 A/turkey/Minnesota/32710/2011 2011/07/12 HA 557478591	MNTQILALIACMLVGTKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEEPLRQILRGSGGIDKE SMGFTYSGIRTNGATSTCRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRNKPALIVWGV HHSGSATEQTKLYGSGSKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVT FTFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHKGGTIVSSLPFQNINPRTVGKCPR YVKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIDKTNQQFEMIDNEFSEIEQQIGNVINWTRDSMTEIWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESL QNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2519	AHA11461 A/turkey/Minnesota/31900/2011 2011/07/05 HA 557478606	MNTQILALIACMLVGTKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEEPLRQILRGSGGIDKE SMGFTYSGIRTNGATSTCRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRNKPALIVWGV HHSGSATEQTKLYGSGSKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVT FTFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHKGGTIVSSLPFQNINPRTVGKCPR YVKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEIWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRAESL QNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2520	AHK10585 A/chicken/Guangdong/G1/2013 2013/05/05 HA 587680636	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2521	AGG53366 A/wild duck/Korea/CSM42-34/2011 2011/03/ HA 459252887	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKE TMGLTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGI HHSGSSTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRY VKQESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVRLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI

2522	AGG53377 A/wild duck/Korea/CSM42-1/2011 2011/03/ HA 459252925	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKE TMGLTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGI HHSGSSTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRY VKQESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVRLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCT
2523	AGG53399 A/wild duck/Korea/MHC39-26/2011 2011/03/ HA 459253005	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKE TMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGI HHSGSTTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRY VKQESLMLATGMKNVPEPPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAME NQHTIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREE AMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2524	AGG53432 A/wild duck/Korea/MHC35-41/2011 2011/03/ HA 459253136	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKE TMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGI HHSGSTTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRY VKQESLMLATGMKNVPEPPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAME NQHTIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREE AMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCT
2525	AGG53476 A/wild duck/Korea/SH19-27/2010 2010/12/ HA 459253257	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKE TMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGI HHSGSTTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRY VKQESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTI
2526	AGG53487 A/wild duck/Korea/SH19-50/2010 2010/01/ HA 459253278	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKE TMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRDPALIVWGI HHSGSTTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDASCEGDCYHSGGTIISNLPFQNINSRAVGKCPRY VKQESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2527	AGG53520 A/wild duck/Korea/SH20-27/2008 2008/12/ HA 459253409	QILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSKGKRTVD LGQCGLLGTITGPPQCDQLLEFSADLIIERREGTDVCYPGKFVNEEALRQILRESGGIEKETMG FTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHHSG STTEQTKLYGSGSKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFN GAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVKQ ESLMLATGMKNVPELPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQS AIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQHTI DLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQINPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMR

2528	AGL43637 A/Taiwan/1/2013 2013// HA 496297389	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGPSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIINNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2529	AGL97639 A/mallard/Minnesota/AI 09-3770/2009 2009/09/12 HA 505555371	IACMLVGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKLCTQGKRPTDLGQC GLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKESMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWGVHHSGSAT EQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVTFTFNGAFI APDRASFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPRYVKQTSLL LATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQI TGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTELWSYNAELLVAMENQHTIDLAD SEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTESLQNRIQIDP VKLS
2530	AGO02477 A/Xuzhou/1/2013 2013/04/25 HA 512403688	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGSKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKSRNMRCTICI
2531	AGR84942 A/Suzhou/5/2013 2013/04/12 HA 526304561	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGSKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2532	AGR84954 A/Nanjing/6/2013 2013/04/11 HA 526304594	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNRNMRCTICI
2533	AGR84978 A/Wuxi/4/2013 2013/04/07 HA 526304656	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKSRNMRCTICI

2534	AGR84990 A/Wuxi/3/2013 2013/04/07 HA 526304688	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKSRNMRCTICI
2535	AGR85002 A/Zhenjiang/1/2013 2013/04/07 HA 526304708	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKSRNKRCTICI
2536	AGR85026 A/Nanjing/2/2013 2013/04/05 HA 526304762	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKSRNMRCTICI
2537	AGU02230 A/Zhejiang/DTID-ZJU05/2013 2013/04/ HA 532808765	LVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGGEVVNATETVERTNIPRICSKGKRTVDLGQ CGLRGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTY SGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTA EQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAF IAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSL LLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQ ITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLAD SEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQID PVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCT
2538	AGU02233 A/Zhejiang/DTID-ZJU08/2013 2013/04/ HA 532808788	FALIAIIPTNADKICLGHHAVSNGTKVNTLTERGGEVVNATETVERTNFPRICSKGKRTVDLGQC GLRGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYS GIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAE QTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFI APDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLL LATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQI TGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADS EMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDP VKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCT
2539	AGW82588 A/tree sparrow/Shanghai/01/2013 2013/05/09 HA 546235348	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTIGI

2540	AGW82600 A/Shanghai/CN01/2013 2013/04/11 HA 546235368	ALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDLGQCG LLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGI RTNGATSACRRSRSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQ TKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIA PDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIMSNLPFQNIDSRAVGKCPRYVKQRSLL LATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQI TGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADS EMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDP VKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2541	AGW82612 A/Shanghai/JS01/2013 2013/04/03 HA 546235388	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKNPALIVWGIH HSGSTAEQTKLYGSGNKLVTVGSSNYQQSFAPSPGARTQVNGQSGRIDFHWLMLNPNDTVT FSFNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPR YVKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSITEVWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFICVKNGNMRCTICI
2542	AHA11472 A/turkey/Minnesota/316 76/2009 2009/12/08 HA 557478625	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANVKKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKE SMGFTYSGIRTNGETSACRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRDKPALIIWGV HHSGSATEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVT FTFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPR YVKQTSLLLATGMRNVPEKPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITNKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRKESL QNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2543	AHA11483 A/turkey/Minnesota/141 35-2/2009 2009/08/07 HA 557478644	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANVKKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKE SMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRDKPALIIWGV HHSGSATEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVT FTFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPR YVKQTSLLLATGMRNVPEKPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITSKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRKESL QNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2544	AHA11500 A/Zhejiang/DTID-ZJU10/2013 2013/10/14 HA 557478676	TQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRTVDL GQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGF TYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVS TAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPPVNGLSGRIDFHWLMLNPNDTVTFSFNG AFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQR SLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAI DQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDL ADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQ IDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKN
2545	AHA57050 A/turkey/Minnesota/146 59/2009 2009/08/12 HA 558484427	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANVKKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKE SMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSNNAAFPQMTKSYRNPRDKPALIIWGV HHSGSATEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVT FTFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPR YVKQTSLLLATGMRNVPEKPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITSKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHNCDDQCMESIRNNTYDHTQYRKES LQNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI

2546	AHA57072 A/turkey/Minnesota/18421/2009 2009/09/09 HA 558484465	MNTQILALIACMLIGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANVKKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKE SMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSNDAAFPQMTKSYRNPRDKPALIIWGV HHSGSATEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTVT FTFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCPR YVKQTSLLLATGMRNVPEKPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTEVWSYNAELLVAMEN QHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRKESL QNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2547	AHD25003 A/Guangdong/02/2013 2013/10/ HA 568260567	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNM
2548	AHF20528 A/Hong Kong/470129/2013 2013/11/30 HA 570933555	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISSLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2549	AHF20568 A/Shanghai/CN02/2013 2013/04/02 HA 570933626	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIMSNLPFQNIDSRAVGKCPR YVKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2550	AHH25185 A/Guangdong/04/2013 2013/12/16 HA 576106234	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIEKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2551	AHJ57411 A/Shanghai/PD-01/2014 2014/01/17 HA 585478041	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVSSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCKGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRIIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2552	AHJ57418 A/Shanghai/PD-02/2014 2014/01/17 HA 585478256	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDICYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLKGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRIIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2553	AHK10800 A/Shanghai/01/2014 2014/01/03 HA 587681014	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRIIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2554	AHM24224 A/Beijing/3/2013 2013/04/16 HA 594704802	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVKEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2555	AHN96472 A/chicken/Shanghai/PD-CN-02/2014 2014/01/21 HA 602701641	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2556	AHZ39686 A/Anhui/DEWH72-01/2013 2013// HA 632807036	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDDAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2557	AHZ39710 A/Anhui/DEWH72-03/2013 2013// HA 632807076	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTDGATSACRRSGSSFYAEMKWLLSNTDDAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2558	AHZ39746 A/Anhui/DEWH72-06/2013 2013// HA 632807136	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGERPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM
2559	AHZ41929 A/mallard/Sweden/1621/2002 2002/12/12 HA 632810949	MNTQILVFALVAIIPINADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNVPRICSRGKR TVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKE TMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRNDPALIIWGIH HSGSTTEQTKLYGSGNKLITVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLILNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQIDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYVK QESLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQH TIDLADSEMNKLYERVRRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIAMGLVFMCVKNGNMRCTICI
2560	AHZ42537 A/mallard/Minnesota/AI09-3770/2009 2009/09/12 HA 632811964	MNTQILAFIACMLVGAKGDKICLGHHAVANGTKVNTLTERGIEVVNATETVETANIKKLCTQGK RPTDLGQCGLLGTLIGPPQCDQFLEFDADLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDK ESMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWG VHHSGSATEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTV TFTFNGAFIAPDRASFFRGESLGVQSDVPLDSGCEGDCFHSGGTIVSSLPFQNINPRTVGKCP RYVKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADY KSTQSAIDQITGKLNRLIDKTNQQFELIDNEFSEIEQQIGNVINWTRDSMTELWSYNAELLVAME NQHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTE SLQNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLVFICIKNGNMRCTICI
2561	AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000 2000/05/20 HA 632811984	MNTQILAFIACMLVGVRGDKICLGHHAVANGTKVNTLTEKGIEVVNATETVESANIKKICTQGKR PTDLGQCGLLGTLIGPPQCDQFLEFDSDLIIERREGTDVCYPGKFTNEESLRQILRGSGGIDKE SMGFTYSGIRTNGATSACRRLGSSSFYAEMKWLLSNSDNAAFPQMTKSYRNPRNKPALIIWG VHHSGSANEQTKLYGSGNKLITVGSSKYQQSFTPSPGARPQVNGQSGRIDFHWLLLDPNDTV TFTFNGAFIAPDRASFFRGESLGIQSDVPLDSSCGGDCFHSGGTIVSSLPFQNINPRTVGKCPR YVKQTSLLLATGMRNVPENPKTRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYK STQSAIDQITGKLNRLIDKTNQQFELMDNEFNEIEQQIGNVINWTRDSMTEVWSYNAELLVAME NQHTIDLADSEMNKLYERVRKQLRENAEEDGTGCFEIFHKCDDQCMESIRNNTYDHTQYRTE SLQNRIQIDPVKLSSGYKDIILWFSFGASCFLLLAIAMGLIFICIKNGNMRCTICI
2562	AID70634 A/Shanghai/Mix1/2014 2014/01/03 HA 660304650	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRIIEKTNQQFELIDNEFNEVEKQISNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2563	AIN76383 A/Zhejiang/LS01/2014 2014/02/08 HA 684694637	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGTTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2564	AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013 2013/12/ HA 699978931	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVEVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2565	AIU47013 A/chicken/Suzhou/040201H/2013 2013/04/ HA 699979673	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDMILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2566	AJJ90490 A/chicken/Shenzhen/742/2013 2013/12/10 HA 755178094	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2567	AJJ90526 A/chicken/Shenzhen/898/2013 2013/12/09 HA 755178154	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDICYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRANGATSACKRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISSLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQIDPVKLSRGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2568	AJJ90538 A/silkie chicken/Shenzhen/918/2013 2013/12/09 HA 755178174	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2569	AJJ90576 A/chicken/Shenzhen/1665/2013 2013/12/12 HA 755178238	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDICYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRANGATSACKRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQIDPVKLSRGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2570	AJJ90588 A/chicken/Shenzhen/2110/2013 2013/12/13 HA 755178258	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSIGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2571	AJJ90661 A/chicken/Dongguan/2912/2013 2013/12/18 HA 755178380	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2572	AJJ90673 A/silkie chicken/Dongguan/3049/2013 2013/12/18 HA 755178400	MNTQILVFALTAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2573	AJJ90795 A/silkie chicken/Dongguan/3281/2013 2013/12/18 HA 755178604	MNTQILVFALIAIIPTNADKICLGHHAVPNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2574	AJJ90891 A/silkie chicken/Dongguan/3520/2013 2013/12/19 HA 755178764	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKXPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2575	AJJ90951 A/chicken/Dongguan/3544/2013 2013/12/19 HA 755178864	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYRNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2576	AJJ91035 A/chicken/Shenzhen/3780/2013 2013/12/19 HA 755179004	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRRSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDNRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2577	AJJ91155 A/chicken/Dongguan/4037/2013 2013/12/19 HA 755179204	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2578	AJJ92005 A/chicken/Shenzhen/801/2013 2013/12/09 HA 755180629	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSRGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2579	AJJ94254 A/chicken/Dongguan/1374/2014 2014/02/21 HA 755184382	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2580	AJJ94606 A/chicken/Dongguan/191/2014 2014/02/20 HA 755184968	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2581	AJJ96552 A/chicken/Jiangxi/12206/2014 2014/03/16 HA 755188219	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTI DLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHNKYREEAMQN RIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2582	AJJ96684 A/chicken/Jiangxi/13207 /2014 2014/03/30 HA 755188439	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKINTLTERGVEVVNATETVERTNIPRICSKGKKTV DLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2583	AJJ96732 A/chicken/Jiangxi/13223 /2014 2014/03/30 HA 755188519	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2551	AJK00354 A/duck/Zhejiang/LS02/2 014 2014/01/12 HA 755194469	MNTQILVFALVAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIVERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKDPALIIWGIHH SGSTTEQTKLYGSGNKLITVGSSNYQQSFVPSPGARPLVNGQSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNINSRAVGKCPRYV KQESLLLATGMKNVPEVPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQVTGKLNRLIEKTNQQFELIDHEFTEVEKQIGNVINWTRDSMTEVWSYNAELLVAMENQ HTIDLADSEMNKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2552	AJJ91264 A/silkie chicken/Dongguan/4129 /2013 2013/12/19 HA 755179386	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLMEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2553	AJJ91314 A/chicken/Shaoxing/241 7/2013 2013/10/20 HA 755179470	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPPVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2554	AJJ91402 A/chicken/Huzhou/4045/ 2013 2013/10/24 HA 755179618	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKEVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2555	AJJ91476 A/chicken/Huzhou/4076/2013 2013/10/24 HA 755179743	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSRGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2556	AJJ91725 A/chicken/Shaoxing/5201/2013 2013/10/28 HA 755180161	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2557	AJJ91885 A/Shenzhen/SP4/2014 2014/01/16 HA 755180429	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGVTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSRGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2558	AJJ91909 A/Shenzhen/SP26/2014 2014/01/20 HA 755180469	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDICYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRANGATSACKRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISSLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDGCMASIRNNTYDHSKYREEAMQN RIQIDPVKLSRGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2559	AJJ91945 A/Shenzhen/SP38/2014 2014/01/22 HA 755180529	MNTQILAFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIGGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2560	AJJ91957 A/Shenzhen/SP44/2014 2014/01/23 HA 755180549	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGTTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISSLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2561	AJJ91969 A/Shenzhen/SP48/2014 2014/01/23 HA 755180569	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2562	AJJ91993 A/chicken/Dongguan/4119/2013 2013/12/19 HA 755180609	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLLGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFTLLAIVMGLVFICVKNGNMRCTICI
2563	AJJ92031 A/chicken/Dongguan/4064/2013 2013/12/19 HA 755180672	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVESSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2564	AJJ92967 A/silkie chicken/Jiangxi/9469/2014 2014/02/16 HA 755182232	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGVTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2565	AJJ93027 A/chicken/Jiangxi/9558/2014 2014/02/16 HA 755182332	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVKEEALRQILRESGGIDKEA MGFTYSGIRTNGVTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2566	AJJ93051 A/chicken/Jiangxi/10573/2014 2014/02/18 HA 755182372	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGVTSACRRSGSSFYAEMKWLLSNTDDAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2567	AJJ93845 A/silkie chicken/Dongguan/157/2014 2014/02/20 HA 755183695	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2568	AJJ93857 A/chicken/Dongguan/169/2014 2014/02/20 HA 755183715	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACMRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2569	AJJ93869 A/chicken/Dongguan/173/2014 2014/02/20 HA 755183735	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTVTGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2570	AJJ93881 A/chicken/Dongguan/189/2014 2014/02/20 HA 755183755	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTVTGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPKY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2571	AJJ93907 A/chicken/Dongguan/449/2014 2014/02/20 HA 755183799	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2572	AJJ93931 A/chicken/Dongguan/536/2014 2014/02/20 HA 755183839	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISKLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2573	AJJ93943 A/chicken/Dongguan/568/2014 2014/02/20 HA 755183859	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIEKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM
2574	AJJ93979 A/silkie chicken/Dongguan/656/2014 2014/02/20 HA 755183919	QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTVTGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFGLIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM
2575	AJJ94134 A/chicken/Dongguan/1051/2014 2014/02/21 HA 755184182	QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI MNTQILVLALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ
2576	AJJ94158 A/chicken/Dongguan/1075/2014 2014/02/21 HA 755184222	NRIQIDPVXLSXGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYRGEAM
2577	AJJ94182 A/chicken/Dongguan/1177/2014 2014/02/21 HA 755184262	QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACKRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSIAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM
2578	AJJ94194 A/silkie chicken/Dongguan/1264/2014 2014/02/21 HA 755184282	QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTI DLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHS VSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSF NGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKSTQ SAIDQVTGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYRGEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFMLLAIVMGLVFICVKNGNMRCTICI

2579	AJJ94206 A/silkie chicken/Dongguan/1268/2014 2014/02/21 HA 755184302	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISDLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2580	AJJ94344 A/silkie chicken/Dongguan/1451/2014 2014/02/21 HA 755184532	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNSTETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRTVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2581	AJJ94356 A/chicken/Dongguan/1456/2014 2014/02/21 HA 755184552	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2582	AJJ94396 A/chicken/Dongguan/1494/2014 2014/02/21 HA 755184618	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPETPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2583	AJJ94754 A/chicken/Dongguan/748/2014 2014/02/20 HA 755185215	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIEKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSNAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSGGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2584	AJJ94838 A/chicken/Dongguan/835/2014 2014/02/20 HA 755185356	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSASTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFGFGASCFILLAIVMGLVFICVKNGNMRCTICI

2585	AJJ94862 A/chicken/Dongguan/843/2014 2014/02/20 HA 755185396	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIEKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSGGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2586	AJJ94886 A/chicken/Dongguan/851/2014 2014/02/20 HA 755185436	MNTQILAFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2587	AJJ94910 A/chicken/Dongguan/874/2014 2014/02/20 HA 755185476	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSASTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2588	AJJ94959 A/silkie chicken/Dongguan/967/2014 2014/02/21 HA 755185558	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACXRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2589	AJJ95048 A/chicken/Dongguan/1009/2014 2014/02/21 HA 755185708	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPETPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDNDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2590	AJJ95171 A/chicken/Dongguan/1314/2014 2014/02/21 HA 755185913	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF NFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2591	AJJ95227 A/chicken/Dongguan/1382/2014 2014/02/21 HA 755186006	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDICYPGKFVNEEALRQILRESGGIDKEAM GFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2592	AJJ95251 A/chicken/Dongguan/1401/2014 2014/02/21 HA 755186046	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYKRVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2593	AJJ95346 A/chicken/Dongguan/1548/2014 2014/02/21 HA 755186206	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYKRVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHNKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2594	AJJ95382 A/chicken/Dongguan/1690/2014 2014/02/21 HA 755186266	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSIGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNR IQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2595	AJJ95464 A/chicken/Shenzhen/138/2014 2014/02/19 HA 755186404	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYRGEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFMLLAIVMGLVFICVKNGNMRCTICI
2596	AJJ95572 A/chicken/Dongguan/1100/2014 2014/02/21 HA 755186584	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIEKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSGGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2597	AJJ95584 A/silkie chicken/Dongguan/1519/2014 2014/02/21 HA 755186604	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPERASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYRGEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFMLLAIVMGLVFICVKNGNMRCTICI
2598	AJJ95596 A/Shenzhen/SP58/2014 2014/01/25 HA 755186624	MNTQILAFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRANGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2599	AJJ95620 A/Shenzhen/SP75/2014 2014/02/15 HA 755186664	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGSTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAVVMGLVFICVKNGNMRCTICI
2600	AJJ95632 A/Shenzhen/SP62/2014 2014/02/05 HA 755186684	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNATFPQMTKSYKNTRKSPALIIWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVETQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2601	AJJ96720 A/chicken/Jiangxi/13220/2014 2014/03/30 HA 755188499	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTTIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSRGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2602	AJJ96817 A/chicken/Jiangxi/9513/2014 2014/02/16 HA 755188661	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATEIVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGVTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2603	AJJ96841 A/Shenzhen/SP139/2014 2014/04/02 HA 755188701	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSTCRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRACFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVERQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2604	AJJ96889 A/chicken/Jiangxi/13496/2014 2014/04/11 HA 755188781	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTXIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKXA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSXGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2605	AJJ96901 A/chicken/Jiangxi/13502/2014 2014/04/11 HA 755188801	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSXGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2606	AJJ96925 A/chicken/Jiangxi/13513/2014 2014/04/11 HA 755188841	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYNGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH TVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDLHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2607	AJJ97267 A/chicken/Jiangxi/13252/2014 2014/03/30 HA 755189411	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2608	AJJ97291 A/chicken/Jiangxi/13493/2014 2014/04/06 HA 755189451	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYNGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2609	AJJ97331 A/chicken/Jiangxi/13512 /2014 2014/04/06 HA 755189517	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYNGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSIGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVK QRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQ SAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTI DLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQN
2610	AJJ97373 A/chicken/Jiangxi/13521 /2014 2014/04/06 HA 755189587	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYNGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPXRASFLRGKSXGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ
2611	AJJ97443 A/chicken/Jiangxi/13530 /2014 2014/04/06 HA 755189702	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTTIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSRGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2612	AJJ97582 A/chicken/Jiangxi/14023 /2014 2014/04/13 HA 755189933	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2613	AJJ97697 A/chicken/Jiangxi/14517 /2014 2014/04/20 HA 755190125	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCDGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMENQ HTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAM QNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2614	AJJ97709 A/chicken/Jiangxi/14518 /2014 2014/04/20 HA 755190145	MNTQILVFALIAIIPANADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYNGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGNCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2615	AJJ97745 A/chicken/Jiangxi/14554 /2014 2014/04/20 HA 755190205	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELMDNEFNEVEKQIGNVINWTRDSITELWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2616	AJJ97757 A/chicken/Shantou/2537 /2014 2014/04/16 HA 755190225	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFKHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2617	AJJ97841 A/duck/Jiangxi/15044/2 014 2014/04/27 HA 755190365	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVRLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2618	AJJ97899 A/chicken/Jiangxi/15524 /2014 2014/05/05 HA 755190462	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKRT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIAKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHRKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFMCVKNGNMRCTICI
2619	AJJ97925 A/silkie chicken/Shantou/2050/2 014 2014/03/25 HA 755190506	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEVPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI
2620	AJJ97973 A/chicken/Shantou/4325 /2014 2014/07/01 HA 755190586	MNTQILVFALISIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRKSGGIDKEA MGFTYSGIRTNGVTSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPAIIVWGIHH SVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFS FNGAFIAPDRASFLRGKSMGIQSGVQVDADCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYV KQRSLLLATGMKNVPEVPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKST QSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQH TIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQ NRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

2621	AJJ97998 A/chicken/Shantou/4816 /2014 2014/07/22 HA 755190628	MNTQILVFALIAIVPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKT VDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEA MGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIH HSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTF SFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRY VKQKSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKS TQSAIDQITGKLNRLIEKTNQQFELVDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMEN QHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEA MQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI

Table 15: Examples of Wild type Hemagglutinin Antigens

SEQ ID NO	Protein / Strain	Nucleic Acid Sequence
2622	H1	AGCAAAAGCAGGGGAAAATAAAAACAACCAAAATGAAGGCAAACCTACTGGTCCTGTTATGTGCACTTGC AGCTGCAGATGCAGACACAATATGTATAGGCTACCATGCGAACAATTCAACCGACACTGTTGACACAGTG CTCGAGAAGAATGTGACAGTGACACACTCTGTTAACCTGCTCGAAGACAGCCACAACGGAAAACTATGTA GATTAAAAGGAATAGCCCCACTACAATTGGGGAAATGTAACATCGCCGGATGGCTCTTGGGAAACCCAGA ATGCGACCCACTGCTTCCAGTGAGATCATGGTCCTACATTGTAGAAACACCAAACTCTGAGAATGGAATAT GTTATCCAGGAGATTTCATCGACTATGAGGAGCTGAGGGAGCAATTGAGCTCAGTGTCATCATTCGAAAG ATTCGAAATATTTCCCAAAGAAAGCTCATGGCCCAACCACAACACAACCAAAGGAGTAACGGCAGCATGC TCCCATGCGGGGAAAAGCAGTTTTTACAGAAATTTGCTATGGCTGACGGAGAAGGAGGGCTCATACCCAA AGCTGAAAAATTCTTATGTGAACAAGAAAGGGAAAGAAGTCCTTGTACTGTGGGGTATTCATCACCCGTCT AACAGTAAGGATCAACAGAATATCTATCAGAATGAAAATGCTTATGTCTCTGTAGTGACTTCAAATTATAAC AGGAGATTTACCCCGGAAATAGCAGAAAGACCCAAAGTAAGAGATCAAGCTGGGAGGATGAACTATTACT GGACCTTGCTAAAACCCGGAGACACAATAATATTTGAGGCAAATGGAAATCTAATAGCACCAAGGT ATGCTTTCGCACTGAGTAGAGGCTTTGGGTCCGGCATCATCACCTCAAACGCATCAATGCATGAGTGTAA CACGAAGTGTCAAACACCCCTGGGAGCTATAAACAGCAGTCTCCCTTTCCAGAATATACACCCAGTCACA ATAGGAGAGTGCCCAAAATACGTCAGGAGTGCCAAATTGAGGATGGTTACAGGACTAAGGAACATTCCGT CCATTCAATCCAGAGGTCTATTTGGAGCCATTGCCGGTTTTATTGAAGGGGGATGGACTGGAATGATAGA TGGATGGTACGGTTATCATCATCAGAATGAACAGGGATCAGGCTATGCAGCGGATCAAAAAAGCACACAA AATGCCATTAACGGGATTACAAACAAGGTGAACTCTGTTATCGAGAAAATGAACATTCAATTCACAGCTGT GGGTAAAGAATTCAACAAATTAGAAAAAAGGATGGAAAATTTAAATAAAAAAGTTGATGATGGATTTCTGG ACATTTGGACATATAATGCAGAATTGTTAGTTCTACTGGAAAATGAAAGGACTCTGGATTTCCATGACTCA AATGTGAAGAATCTGTATGAGAAAGTAAAAAGCCAATTAAAGAATAATGCCAAAGAAATCGGAAATGGATG TTTTGAGTTCTACCACAAGTGTGACAATGAATGCATGGAAAGTGTAAGAAATGGGACTTATGATTATCCCA AATATTCAGAAGAGTCAAAGTTGAACAGGGAAAAGGTAGATGGAGTGAAATTGGAATCAATGGGGATCTA TCAGATTCTGGCGATCTACTCAACTGTCGCCAGTTCACTGGTGCTTTTGGTCTCCCTGGGGGCAATCAGT TTCTGGATGTGTTCTAATGGATCTTTGCAGTGCAGAATATGCATCTGAGATTAGAATTTCAGAAATATGAG GAAAAACACCCTTGTTTCTACT

2623	H7	AGCGAAAGCAGGGGATACAAAATGAACACTCAAATCCTGGTATTCGCTCTGATTGCGATCATTCCAACAA ATGCAGACAAAATCTGCCTCGGACATCATGCCGTGTCAAACGGAACCAAAGTAAACACATTAACTGAAAG AGGAGTGGAAGTCGTCAATGCAACTGAAACAGTGGAACGAACAAACATCCCCAGGATCTGCTCAAAAGG GAAAAGGACAGTTGACCTCGGTCAATGTGGACTCCTGGGGACAATCACTGGACCACCTCAATGTGACCA ATTCCTAGAATTTTCAGCCGATTTAATTATTGAGAGGCGAGAAGGAAGTGATGTCTGTTATCCTGGGAAAT TCGTGAATGAAGAAGCTCTGAGGCAAATTCTCAGAGAATCAGGCGGAATTGACAAGGAAGCAATGGGATT CACATACAGTGGAATAAGAACTAATGGAGCAACCAGTGCATGTAGGAGATCAGGATCTTCATTCTATGCA GAAATGAAATGGCTCCTGTCAAACACAGATGATGCTGCATTCCCGCAGATGACTAAGTCATATAAAAATAC AAGAAAAAGCCCAGCTCTAATAGTATGGGGGATCCATCATTCCGTATCAACTGCAGAGCAAACCAAGCTA TATGGGAGTGGAAACAAACTGGTGACAGTTGGGAGTTCTAATTATCAACAATCTTTTGTACCGAGTCCAG GAGCGAGACCACAAGTTAATGGTCTATCTGGAAGAATTGACTTTCATTGGCTAATGCTAAATCCCAATGAT ACAGTCACTTTCAGTTTCAATGGGGCTTTCATAGCTCCAGACCGTGCAAGCTTCCTGAGAGGAAAATCTAT GGGAATCCAGAGTGGAGTACAGGTTGATGCCAATTGTGAAGGGGACTGCTATCATAGTGGAGGGACAAT AATAAGTAACTTGCCATTTCAGAACATAGATAGCAGGGCAGTTGGAAAATGTCCGAGATATGTTAAGCAAA GGAGTCTGCTGCTAGCAACAGGGATGAAGAATGTTCCTGAGATTCCAAAGGGAAGAGGCCTATTTGGTG CTATAGCGGGTTTCATTGAAAATGGATGGGAAGGCCTAATTGATGGTTGGTATGGTTTCAGACACCAGAA TGCACAGGGAGAGGGAACTGCTGCAGATTACAAAAGCACTCAATCGGCAATTGATCAAATAACAGGAAAA TTAAACCGGCTTATAGAAAAAACCAACCAACAATTTGAGTTGATAGACAATGAATTCAATGAGGTAGAGAA GCAAATCGGTAATGTGATAAATTGGACCAGAGATTCTATAACAGAAGTGTGGTCATACAATGCTGAACTCT TGGTAGCAATGGAGAACCAGCATACAATTGATCTGGCTGATTCAGAAATGGACAAACTGTACGAACGAGT GAAAAGACAGCTGAGAGAGAATGCTGAAGAAGATGGCACTGGTTGCTTTGAAATATTTCACAAGTGTGAT GATGACTGTATGGCCAGTATTAGAAATAACACCTATGATCACAGCAAATACAGGGAAGAGGCAATGCAAA ATAGAATACAGATTGACCCAGTCAAACTAAGCAGCGGCTACAAAGATGTGATACTTTGGTTTAGCTTCGG GGCATCATGTTTCATACTTCTAGCCATTGTAATGGGCCTTGTCTTCATATGTGTAAAGAATGGAAACATGC GGTGCACTATTTGTATATAAGTTTGGAAAAAAACACCCTTGTTTCTAC
2624	H10	ATGTACAAAATAGTAGTGATAATCGCGCTCCTTGGAGCTGTGAAAGGTCTTGATAAAATCTGTCTAGGACA TCATGCAGTGGCTAATGGGACCATCGTAAAGACTCTCACAAACGAACAGGAAGAGGTAACCAACGCTACT GAAACAGTGGAGAGTACAGGCATAAACAGATTATGTATGAAAGGAAGAAAACATAAAGACCTGGGCAACT GCCATCCAATAGGGATGCTAATAGGGACTCCAGCTTGTGATCTGCACCTTACAGGGATGTGGGACACTCT CATTGAACGAGAGAATGCTATTGCTTACTGCTACCCTGGAGCTACTGTAAATGTAGAAGCACTAAGGCAG AAGATAATGGAGAGTGGAGGGATCAACAAGATAAGCACTGGCTTCACTTATGGATCTTCCATAAACTCGG CCGGGACCACTAGAGCGTGCATGAGGAATGGAGGGAATAGCTTTTATGCAGAGCTTAAGTGGCTGGTAT CAAAGAGCAAAGGACAAAACTTCCCTCAGACCACGAACACTTACAGAAATACAGACACGGCTGAACACCT CATAATGTGGGGAATTCATCACCCTTCTAGCACTCAAGAGAAGAATGATCTATATGGAACACAATCACTGT CCATATCAGTCGGGAGTTCCACTTACCGGAACAATTTTGTTCCGGTTGTTGGAGCAAGACCTCAGGTCAA TGGACAAAGTGGCAGAATTGATTTTCACTGGACACTAGTACAGCCAGGTGACAACATCACCTTCTCACAC AATGGGGGCCTGATAGCACCGAGCCGAGTTAGCAAATTAATTGGGAGGGGATTGGGAATCCAATCAGAC GCACCAATAGACAATAATTGTGAGTCCAAATGTTTTTGGAGAGGGGGTTCTATAAATACAAGGCTTCCCTT TCAAAATTTGTCACCAAGAACAGTGGGTCAGTGTCCTAAATATGTGAACAGAAGAAGCTTGATGCTTGCAA CAGGAATGAGAAACGTACCAGAACTAATACAAGGGAGAGGTCTATTTGGTGCAATAGCAGGGTTTTTAGA GAATGGGTGGGAAGGAATGGTAGATGGCTGGTATGGTTTCAGACATCAAAATGCTCAGGGCACAGGCCA GGCCGCTGATTACAAGAGTACTCAGGCAGCTATTGATCAAATCACTGGGAAACTGAATAGACTTGTTGAA AAAACCAATACTGAGTTCGAGTCAATAGAATCTGAGTTCAGTGAGATCGAACACCAAATCGGTAACGTCAT CAATTGGACTAAGGATTCAATAACCGACATTTGGACTTATCAGGCTGAGCTGTTGGTGGCAATGGAGAAC CAGCATACAATCGACATGGCTGACTCAGAGATGTTGAATCTATATGAAAGAGTGAGGAAACAACTAAGGC AGAATGCAGAAGAAGATGGGAAAGGATGTTTTGAGATATATCATGCTTGTGATGATTCATGCATGGAGAG CATAAGAAACAACACCTATGACCATTCACAGTACAGAGAGGAAGCTCTTTTGAACAGATTGAATATCAACC CAGTGACACTCTCTTCTGGATATAAAGACATCATTCTCTGGTTTAGCTTCGGGGCATCATGTTTTGTTCTTC TAGCCGTTGTCATGGGTCTTTTCTTTTTCTGTCTGAAGAATGGAAACATGCGATGCACAATCTGTATTTAG

Table 16: Codon Optimized Sequences Encoding H1 Hemagglutinin

Seq ID No.	Nucleotide sequence
2625	ATGAAGGCAAACCTCTTAGTCCTTCTCTGCGCACTGGCCGCTGCCGACGCAGATACCATATGTATCGGTTATCACGCCAATAACTC TACCGATACAGTCGATACGGTGCTTGAGAAAAACGTCACCGTGACTCATAGCGTGAACCTGCTCGAGGACTCCCATAACGGTAAG CTATGCCGACTAAAGGGCATCGCCCCCCTGCAGCTGGGGAAATGCAACATCGCTGGCTGGTTGTTAGGTAATCCCGAATGCGATC CACTGCTGCCCGTGAGATCCTGGTCTTACATCGTCGAAACCCCTAACTCTGAAAATGGCATCTGCTATCCGGGAGACTTTATTGAC TATGAGGAGCTCCGAGAACAGCTGAGCTCAGTATCCAGCTTTGAACGCTTCGAAATTTTTCCAAAGGAATCCAGCTGGCCAAACCA TAACACGAATGGCGTCACAGCCGCATGCAGCCACGAGGGAAAGAGCAGCTTTTATAGGAACCTTCTGTGGTTGACGGAAAAAGAG GGAAGTTATCCCAAACTTAAAAACAGCTATGTGAATAAAAAGGGCAAGGAGGTCTTGGTACTTTGGGGAATCCATCACCCTCCAAA CAGTAAGGAACAACAAAATCTGTACCAGAATGAAAATGCATACGTGTCCGTGGTTACATCTAATTACAACCGACGGTTTACTCCCG AGATCGCCGAGCGACCGAAAGTTCGCGACCAGGCCGGGAGGATGAACTATTATTGGACGTTGCTGAAACCCGGTGACACAATTAT CTTTGAGGCAAATGGTAATCTTATCGCCCCGATGTACGCATTCGCACTGTCGCGCGGATTTGGCAGTGGGATTATAACCAGTAACG CATCTATGCATGAATGCAACACAAAGTGCCAGACTCCTTTAGGCGCTATAAACTCTTCTTTGCCCTATCAGAATATACACCCAGTGA CCATCGGCGAGTGTCCAAAGTACGTACGCAGCGCCAAGCTCCGAATGGTTACAGGCCTCAGAAATAACCCCAGTATCCAGAGCC GCGGCTTATTCGGGGCGATCGCTGGATTTATTGAGGGGGGCTGGACGGGAATGATAGATGGGTGGTACGGCTACCATCATCAGA ATGAGCAGGGTAGCGGCTATGCGGCCGACCAGAAAAGCACCCAGAACGCTATCAACGGTATAACCAATAAAGTGAACACTGTGAT CGAAAAAATGAACATACAGTTCACTGCCGTGGGAAAGGAATTCAACAAACTGGAAAAAAGGATGGAAAATCTTAACAAAAAAGTGG ATGACGGGTTTTTGGACATTTGGACATACAATGCCGAACTGCTGGTTCTGCTGGAGAACGAAAGGACACTGGACTTTCACGATTCC AACGTCAAGAATCTCTATGAAAAAGTGAAGTCCCAATTGAAAAACAACGCCAAGGAGATCGGTAATGGGTGCTTTGAGTTCTATCA CAAGTGTGACAACGAGTGCATGGAGAGCGTTAGGAACGGCACATACGATTACCCAAAATATTCTGAGGAGTCTAAGTTGAACAGA GAGAAGGTCGATGGCGTTAAACTGGAAAGCATGGGAATTTACCAGATCCTTGCCATCTACTCCACCGTGGCCTCCTCCCTTGTGC TGCTGGTGTCGCTGGGGGCCATCTCCTTTTGGATGTGCTCCAATGGTTCACTGCAATGTCGAATCTGCATC

2626	ATGAAGGCTAACCTCCTAGTGCTGCTCTGTGCCTTGGCGGCTGCGGATGCTGACACCATTTGCATTGGGTACCACGCCAATAATT CAACCGATACTGTCGACACCGTGTTGGAGAAGAATGTGACCGTGACTCATTCCGTGAACCTTCTTGAAGATTCGCACAACGGGAA GCTGTGTCGACTCAAAGGCATCGCGCCTCTACAGTTGGGCAAATGTAACATCGCTGGGTGGCTGCTCGGGAACCCAGAATGTGAT CCTTTGCTGCCAGTGAGATCGTGGTCATACATTGTGGAGACCCCCAATTCAGAGAACGGGATATGCTACCCAGGTGACTTTATCGA TTATGAGGAACTTCGGGAACAACTGTCTTCAGTGTCGTCATTTGAGCGTTTCGAGATCTTTCCCAAAGAGTCCTCGTGGCCAAACC ACAACACTAACGGCGTTACTGCCGCCTGCTCACATGAAGGAAAGAGCTCATTCTACAGAAACCTGTTGTGGCTCACCGAGAAGGA GGGGAGTTATCCCAAGTTGAAAAACAGTTATGTTAACAAGAAGGGCAAAGAAGTGTTAGTCCTGTGGGGAATCCACCACCCACCG AATTCTAAAGAGCAGCAGAATCTATATCAGAACGAGAACGCCTATGTGTCAGTTGTAACTTCAAATTACAACCGCAGGTTTACTCCT GAAATCGCGGAGCGCCCGAAAGTGCGTGATCAGGCTGGACGTATGAATTACTACTGGACTTTGCTAAAACCTGGCGACACCATCA TTTTCGAAGCTAACGGCAATCTCATTGCGCCGATGTACGCCTTTGCTCTCTCTAGGGGGTTCGGGTCTGGGATCATTACCAGCAAT GCTTCCATGCACGAGTGTAATACAAAGTGCCAGACTCCCCTCGGCGCGATTAATAGCTCGCTCCCCTACCAAAACATACACCCAGT GACAATTGGCGAGTGCCCTAAATATGTGCGCTCTGCTAAGTTACGTATGGTGACGGGCCTAAGAAACAACCCCTCAATTCAGAGC AGGGGGCTGTTCGGAGCTATTGCAGGTTTCATCGAAGGGGGCTGGACAGGGATGATCGACGGCTGGTATGGGTATCACCACCAG AACGAACAGGGATCCGGCTATGCTGCCGATCAGAAATCAACACAAAATGCAATCAATGGCATCACTAACAAAGTGAATACCGTAAT TGAGAAGATGAACATACAGTTTACCGCTGTCGGTAAAGAATTCAACAAGCTGGAGAAACGGATGGAGAATCTGAACAAGAAAGTC GATGATGGCTTCTTGGATATCTGGACCTATAATGCTGAGTTACTTGTGCTCTTGGAAAACGAGCGTACTCTCGACTTTCATGATTCA AATGTCAAGAATCTGTATGAAAAGGTAAAGTCCCAGCTAAAAAACAATGCCAAGGAGATCGGGAATGGCTGCTTTGAATTCTACCA CAAATGCGATAATGAGTGCATGGAGTCGGTGAGGAACGGAACCTACGATTACCCAAAATATAGCGAGGAATCAAAACTGAACAGA GAGAAGGTGGATGGTGTGAAGCTGGAATCCATGGGTATCTACCAGATCCTCGCCATCTATTCAACTGTGGCTAGTTCCCTGGTGC TGCTCGTGAGCCTTGGTGCAATCTCCTTCTGGATGTGTTCCAACGGGTCATTGCAGTGTAGGATCTGCATA
2627	ATGAAGGCTAACTTGCTGGTCTTGCTTTGCGCGCTGGCCGCTGCTGACGCCGACACCATCTGTATTGGGTATCATGCTAACAACT CCACGGATACCGTGGATACAGTGTTGGAAAAAAATGTTACAGTTACTCATTCGGTGAACCTGCTGGAGGACTCTCACAATGGGAAG CTGTGCAGGCTCAAAGGTATCGCTCCGCTCCAACTAGGCAAATGTAATATTGCCGGCTGGCTGTTAGGAAACCCAGAGTGTGACC CTCTCTTACCCGTGAGATCTTGGTCCTACATCGTCGAGACCCCCAACAGCGAAAACGGAATTTGTTATCCCGGAGACTTCATAGAT TATGAGGAGCTGCGCGAACAGCTCTCTTCTGTTAGCTCTTTTGAGCGTTTTGAGATCTTTCCTAAAGAGAGTTCATGGCCCAACCA CAATACCAACGGAGTGACAGCCGCATGCTCCCATGAAGGGAAGTCTAGCTTCTATAGAAATCTTCTGTGGCTGACTGAAAAAGAAG GATCTTACCCCAAGTTGAAGAATTCTTATGTGAACAAAAAGGGCAAAGAGGTGTTGGTACTCTGGGGCATTCATCACCCACCTAAT TCAAAGGAGCAGCAAAACCTTTATCAAAACGAGAATGCCTATGTTAGCGTCGTCACTTCTAATTATAACCGGAGATTCACCCCGGA GATAGCGGAACGCCCCAAGGTTCGGGATCAGGCTGGCCGCATGAATTACTACTGGACCCTCCTAAAACCTGGCGACACCATAATC TTCGAAGCAAATGGCAACCTGATCGCACCTATGTACGCCTTTGCGCTGTCTAGGGGATTCGGCTCGGGCATTATTACCAGTAATGC CTCAATGCACGAGTGTAATACAAAATGCCAGACCCCTCTGGGAGCCATTAATTCTAGTCTGCCTTATCAGAATATTCATCCTGTGAC CATTGGCGAATGTCCGAAATATGTTAGATCCGCAAAGCTGCGAATGGTTACCGGACTTCGCAACAACCCCTCTATACAAAGCCGC GGACTCTTTGGAGCAATCGCTGGCTTCATTGAGGGTGGCTGGACGGGCATGATCGATGGGTGGTACGGATACCATCATCAGAAC GAGCAGGGAAGCGGATATGCCGCAGACCAAAAATCTACTCAGAACGCTATTAACGGAATTACCAACAAAGTCAACACAGTTATCGA GAAAATGAATATTCAGTTCACCGCAGTTGGTAAGGAATTTAACAAACTGGAGAAACGAATGGAGAACTTGAACAAGAAGGTCGACG ATGGCTTCTTAGACATCTGGACATATAATGCCGAACTGTTGGTCCTCCTGGAGAACGAGAGAACCTTGGACTTTCACGACTCTAAC GTGAAGAATCTTTACGAGAAGGTGAAGTCCCAACTGAAGAATAATGCAAAAGAGATCGGAAACGGCTGTTTTGAGTTCTACCACAA GTGCGACAATGAGTGCATGGAGAGCGTGCGGAATGGGACATATGATTATCCCAAGTATAGTGAAGAAAGCAAGCTGAATCGAGAA AAGGTTGACGGAGTGAAGCTGGAGAGCATGGGGATCTACCAGATCTTAGCAATTTATAGCACAGTGGCTTCTTCCCTTGTGTTGCT GGTCTCGCTGGGAGCTATCAGCTTTTGGATGTGCAGCAACGGAAGTCTGCAGTGCCGCATTTGCATT

2628	ATGAAAGCCAACCTGTTGGTGCTGTTATGCGCGCTGGCAGCCGCAGATGCTGATACTATCTGTATCGGATACCATGCCAATAACA GCACCGATACCGTGGATACTGTGTTAGAGAAGAATGTGACAGTGACGCATTCAGTAAACCTCCTCGAGGACAGTCATAATGGAAAA CTGTGTAGGCTCAAGGGTATAGCTCCGCTGCAGCTGGGCAAGTGTAACATCGCTGGGTGGCTGCTGGGCAACCCTGAATGCGAT CCCCTGTTGCCCGTTCGGAGTTGGTCTTACATCGTGGAGACCCCAAACTCTGAGAACGGAATTTGTTACCCCGGTGACTTTATCGA TTACGAGGAATTGCGGGAGCAGCTTAGCTCAGTGTCTTCATTTGAACGGTTTGAAATCTTTCCCAAGGAAAGCAGCTGGCCCAATC ATAATACTAACGGCGTGACTGCCGCTTGTTCCCACGAAGGCAAGAGCTCCTTCTACAGAAATCTGCTTTGGTTGACAGAGAAAGAG GGCAGCTATCCAAAGCTGAAGAACAGTTATGTGAATAAGAAGGGCAAGGAGGTCCTGGTTCTTTGGGGGATCCATCATCCCCCTA ATAGTAAGGAGCAGCAGAACTTATACCAGAATGAGAATGCCTACGTCAGCGTTGTCACCTCAAACTATAACCGGCGATTTACACCC GAAATTGCCGAACGTCCCAAGGTCCGCGACCAAGCCGGGCGTATGAACTACTATTGGACACTTCTGAAACCTGGGGACACAATAA TTTTCGAAGCCAACGGTAACTTAATCGCTCCCATGTACGCTTTCGCTCTCAGCAGGGGCTTCGGCAGCGGCATTATCACTTCCAAT GCGTCCATGCATGAATGTAATACGAAGTGCCAGACCCCTCTCGGTGCCATAAACTCTTCTCTCCCCTACCAGAACATTCACCCTGT CACCATAGGAGAATGCCCAAAGTATGTCCGGTCGGCAAAGTTGAGGATGGTTACGGGCCTGCGTAACAACCCTTCTATACAGTCT AGGGGCCTGTTCGGAGCAATCGCCGGCTTCATCGAAGGAGGCTGGACTGGGATGATCGATGGCTGGTACGGCTATCATCACCAG AATGAGCAGGGCTCAGGTTATGCAGCCGATCAGAAGTCCACACAGAACGCTATTAACGGGATAACGAACAAGGTGAACACCGTCA TTGAAAAGATGAACATTCAGTTTACAGCCGTTGGAAAAGAGTTCAACAAACTGGAAAAGCGGATGGAGAACCTGAACAAAAAGGTG GACGATGGATTTCTGGATATTTGGACATATAACGCCGAGCTGCTGGTACTCCTTGAGAACGAGCGTACGCTTGACTTTCATGACAG TAACGTGAAGAATCTGTACGAGAAGGTCAAAAGCCAGCTGAAGAACAATGCCAAAGAAATTGGGAACGGGTGTTTCGAATTCTACC ACAAGTGTGACAATGAATGCATGGAGAGTGTTCGGAACGGAACCTACGACTATCCTAAATATTCTGAGGAGTCCAAGCTGAATCGG GAAAAGGTTGACGGGGTGAAACTGGAGAGTATGGGAATCTACCAGATCCTGGCTATTTACTCCACAGTCGCTTCCTCTCTCGTCTT GTTGGTGTCCCTTGGTGCCATCTCATTTTGGATGTGTTCTAATGGTTCTCTTCAGTGCCGAATCTGTATA
2629	ATGAAAGCTAACCTTCTCGTATTGCTCTGCGCTTTGGCTGCAGCCGATGCCGATACGATTTGCATCGGATACCACGCAAATAACAG CACTGACACTGTGGACACGGTACTGGAAAAGAATGTTACCGTGACACACTCCGTCAACCTGCTCGAGGACTCCCACAACGGAAAG TTGTGTAGGCTTAAAGGTATCGCCCCTCTACAGTTAGGGAAGTGCAACATCGCAGGGTGGTTGTTAGGAAACCCCGAATGTGACC CACTCCTGCCTGTGCGCTCTTGGAGCTATATTGTGGAAACCCCGAATAGCGAAAACGGAATTTGTTACCCTGGCGACTTTATCGAT TATGAAGAGCTCCGAGAGCAACTGAGCAGCGTGTCCAGCTTTGAGCGATTTGAAATATTCCCCAAAGAGAGTTCCTGGCCCAACC ATAACACTAACGGAGTGACCGCTGCATGTTCGCACGAAGGCAAATCGAGTTTTTACCGGAACCTGCTTTGGCTCACCGAGAAGGA AGGAAGTTATCCCAAACTCAAAAATTCATACGTGAATAAAAAGGGTAAGGAGGTGCTGGTGCTGTGGGGAATCCACCATCCTCCAA ACAGCAAGGAACAACAGAATCTCTACCAGAATGAGAACGCCTATGTGAGCGTGGTAACCTCCAACTACAATCGGCGGTTCACTCC TGAGATCGCCGAGCGCCCGAAGGTCCGGGATCAAGCAGGTCGAATGAACTACTACTGGACCCTGCTGAAGCCAGGTGATACTAT CATTTTCGAGGCTAATGGAAACCTGATTGCGCCTATGTACGCTTTCGCCCTGTCTAGGGGGTTTGGGTCCGGTATCATCACGTCTA ACGCTTCAATGCATGAATGCAACACGAAGTGTCAGACCCCGCTTGGCGCCATAAACAGCTCCCTCCCCTACCAGAACATTCACCC CGTCACCATAGGCGAGTGTCCCAAGTACGTGCGGTCAGCCAAGTTGCGGATGGTAACGGGCCTCCGGAATAATCCAAGTATACAA TCCCGAGGACTTTTCGGGGCAATCGCGGGCTTCATTGAGGGCGGATGGACCGGCATGATTGATGGATGGTACGGTTACCACCAC CAGAACGAGCAGGGCAGTGGATATGCCGCTGACCAGAAGAGCACTCAGAACGCAATTAATGGCATCACCAATAAGGTGAACACC GTCATTGAGAAGATGAACATCCAGTTTACGGCAGTAGGCAAGGAGTTTAACAAGCTGGAGAAACGTATGGAAAACCTGAATAAGAA AGTGGACGACGGGTTTTTGGATATTTGGACATATAACGCCGAACTCCTCGTGCTGCTCGAGAATGAAAGAACATTAGACTTTCACG ATAGTAACGTGAAAAACCTGTATGAAAAGGTCAAATCCCAGCTAAAGAACAACGCCAAAGAGATCGGCAATGGCTGTTTTGAGTTT TACCACAAGTGTGATAATGAATGCATGGAAAGTGTTCGGAACGGAACCTACGATTATCCAAAATACTCGGAGGAGTCCAAACTGAA TAGAGAGAAGGTGGACGGCGTCAAACTAGAGAGTATGGGTATCTATCAGATACTTGCGATCTACTCAACCGTCGCCAGCTCCTTA GTACTTCTCGTCAGCCTCGGTGCTATCTCCTTTTGGATGTGTAGTAACGGCAGTCTGCAGTGTCGGATATGCATC

2630	ATGAAGGCTAACCTGTTGGTGCTCCTATGTGCCCTGGCCGCTGCAGATGCTGACACAATCTGCATTGGGTATCATGCAAACAACTC CACCGACACTGTTGACACAGTTTTGGAAAAGAACGTGACAGTCACTCACAGCGTAAACCTCCTGGAAGACTCACACAATGGGAAG TTATGTCGGCTGAAGGGCATAGCCCCATTGCAACTGGGCAAGTGCAATATAGCGGGGTGGCTTCTCGGCAACCCGGAATGCGAC CCCCTGTTGCCCGTACGCAGCTGGAGTTACATTGTGGAGACGCCCAATTCCGAGAATGGCATCTGTTACCCGGGCGATTTCATAG ATTACGAAGAACTGCGGGAGCAGCTCTCCTCCGTTTCCTCGTTCGAACGGTTCGAGATTTTCCCTAAAGAATCATCATGGCCTAAT CACAATACAAATGGCGTGACGGCCGCATGCTCACACGAGGGCAAGTCCAGCTTTTACAGAAATCTCCTGTGGCTGACGGAGAAGG AAGGAAGCTACCCTAAGCTTAAGAATAGCTATGTAAACAAAAAGGGGAAGGAGGTGCTGGTTCTCTGGGGGATTCATCATCCGCC AAATAGTAAGGAACAGCAGAACCTTTACCAGAATGAGAATGCATACGTCTCTGTGGTCACGTCTAACTACAATCGGCGCTTCACAC CCGAGATCGCTGAACGGCCCAAAGTCCGGGACCAAGCCGGACGAATGAATTACTACTGGACCCTTTTGAAGCCCGGCGATACCA TTATCTTCGAAGCGAACGGGAACCTGATAGCTCCCATGTACGCCTTTGCTCTCAGTAGGGGATTCGGCTCCGGCATTATTACCTCT AATGCTTCTATGCACGAGTGCAACACAAAATGTCAGACTCCGCTCGGTGCCATCAACAGTTCGTTGCCATACCAGAATATACACCC CGTCACTATCGGGGAATGTCCTAAATACGTGAGGAGCGCCAAACTCAGGATGGTCACTGGACTCAGGAATAATCCAAGCATCCAG TCGAGGGGCTTATTCGGAGCAATTGCTGGCTTCATCGAAGGCGGGTGGACTGGCATGATCGATGGATGGTATGGGTACCACCAC CAGAACGAACAAGGTAGCGGATATGCCGCTGACCAGAAATCAACACAAAACGCCATCAACGGTATTACCAACAAGGTTAACACTGT TATTGAGAAGATGAATATCCAATTTACCGCCGTAGGGAAGGAGTTTAATAAGTTGGAGAAACGTATGGAGAATCTGAACAAGAAAG TTGACGACGGATTTCTGGACATCTGGACCTACAACGCCGAGCTGTTAGTGCTGTTGGAAAATGAGAGAACCCTTGACTTCCATGAT AGCAACGTGAAAAACCTATATGAAAAGGTCAAGTCCCAGCTGAAGAACAATGCAAAGGAAATAGGCAATGGCTGCTTCGAGTTTTA TCATAAATGCGACAACGAATGTATGGAAAGCGTCCGAAACGGCACTTATGACTATCCCAAATATAGCGAAGAGTCAAAGCTGAATC GGGAAAAGGTGGACGGGGTCAAGCTCGAGTCAATGGGGATCTACCAGATTCTTGCAATCTATTCCACAGTGGCGTCTTCCCTCGT GCTGCTGGTTAGTCTCGGCGCCATATCATTTTGGATGTGCAGCAATGGCTCCCTGCAGTGTCGGATCTGTATC
2631	ATGAAGGCGAACTTGCTGGTCCTGTTGTGCGCTTTAGCTGCTGCGGATGCCGATACAATCTGTATCGGGTATCACGCAAACAACA GTACAGATACCGTGGATACAGTTCTCGAGAAGAATGTCACTGTTACGCACAGTGTAAACCTGCTGGAGGACAGTCACAATGGGAA GCTGTGCCGGCTAAAGGGCATTGCGCCACTACAGCTCGGCAAGTGCAATATTGCGGGGTGGTTATTGGGTAACCCCGAATGCGA CCCACTATTGCCAGTGCGCTCTTGGTCGTACATCGTCGAAACCCCCAATTCAGAAAATGGCATTTGCTACCCAGGGGATTTTATAG ACTACGAAGAGTTACGAGAACAATTAAGTAGTGTTTCTTCCTTTGAGCGATTCGAGATTTTCCCCAAGGAGAGTTCTTGGCCCAAC CATAACACAAATGGAGTGACTGCCGCATGCTCACACGAGGGCAAATCATCCTTTTATAGAAACCTGCTTTGGCTGACTGAAAAGGA GGGGAGTTACCCGAAGCTTAAAAACTCCTATGTGAACAAGAAGGGCAAGGAAGTGCTGGTCCTGTGGGGGATCCACCACCCACC CAACTCCAAAGAGCAACAGAACCTCTACCAGAACGAGAATGCGTATGTGAGCGTGGTGACAAGTAACTATAATCGAAGGTTCACTC CCGAGATAGCAGAGCGTCCAAAAGTGCGAGATCAAGCCGGCCGAATGAACTACTATTGGACCTTACTGAAACCCGGCGATACCAT CATATTCGAAGCTAACGGCAATCTCATAGCTCCTATGTACGCTTTCGCTTTGAGTAGAGGTTTCGGATCTGGAATTATAACATCTAA TGCATCAATGCACGAGTGCAATACTAAATGCCAGACCCCTCTTGGTGCTATCAACTCTAGTTTACCCTATCAGAACATCCACCCTGT GACGATAGGAGAGTGTCCCAAGTATGTGCGCTCCGCTAAGCTTAGAATGGTCACCGGGCTTCGGAACAACCCTTCAATCCAAAGT CGAGGGCTTTTTGGCGCTATCGCTGGCTTCATCGAAGGCGGTTGGACAGGTATGATCGATGGTTGGTACGGCTACCATCATCAGA ACGAGCAGGGGTCTGGTTATGCGGCCGACCAAAAATCTACTCAGAATGCCATCAACGGGATCACAAATAAGGTAAACACTGTGAT TGAGAAGATGAACATCCAGTTTACTGCAGTGGGAAAGGAGTTTAACAAACTGGAGAAACGTATGGAAAACCTAAACAAAAAAGTGG ACGACGGATTCTTAGACATTTGGACATACAACGCCGAGCTGCTCGTGCTGCTTGAGAATGAGCGGACACTCGACTTCCACGACAG CAACGTGAAGAACCTATACGAAAAGGTCAAGAGCCAGCTCAAGAATAACGCGAAGGAGATTGGAAACGGCTGCTTCGAGTTCTAC CATAAGTGCGATAACGAATGTATGGAGTCTGTTCGGAATGGCACTTACGACTACCCAAAGTATAGCGAGGAATCTAAGCTCAACCG AGAGAAGGTGGATGGAGTGAAATTAGAGTCAATGGGAAATTTATCAGATTCTGGCCATCTACTCAACAGTTGCGAGCTCGTTAGTGC TGCTTGTCTCACTGGGCGCGATTAGTTTTTGGATGTGCAGTAATGGGTCCCTCCAGTGTCGCATCTGCATT

2632	ATGAAGGCCAACTTGCTGGTGTTACTATGTGCTCTCGCCGCCGCCGATGCCGATACTATTTGCATTGGCTATCACGCCAACAATAG TACGGACACCGTGGACACTGTCCTGGAAAAAAATGTCACCGTCACCCATTCTGTGAATCTCCTAGAGGATTCTCATAACGGCAAGC TGTGCCGGCTTAAAGGAATCGCTCCCTTGCAGCTTGGGAAGTGCAATATCGCCGGCTGGCTCCTTGGAAACCCGGAGTGTGATCC ATTGCTCCCTGTCAGAAGCTGGAGCTATATCGTTGAGACCCCGAACAGTGAGAATGGCATCTGTTACCCCGGAGATTTCATTGATT ACGAGGAGCTGCGCGAACAGCTCAGCTCAGTCTCTAGTTTCGAGCGATTTGAGATCTTCCCCAAGGAGTCTAGCTGGCCAAATCA TAACACCAACGGAGTGACAGCCGCCTGTAGCCATGAAGGGAAAAGCAGCTTCTACCGAAATTTGCTGTGGCTAACTGAGAAAGAG GGGAGTTATCCTAAGCTCAAAAACAGCTATGTGAACAAGAAGGGGAAAGAAGTCCTGGTGTTGTGGGGCATCCACCATCCCCCAA ATAGCAAGGAACAGCAGAACCTTTACCAGAACGAGAATGCATACGTGAGCGTCGTAACATCCAATTATAACAGACGGTTCACACCC GAGATAGCCGAACGGCCAAAAGTGAGGGACCAGGCCGGAAGGATGAATTACTATTGGACTCTCTTAAAGCCAGGCGACACCATTA TATTTGAGGCAAACGGAAACTTGATTGCGCCAATGTACGCATTCGCACTGTCACGGGGATTCGGATCCGGGATTATCACGTCCAAT GCATCGATGCATGAGTGCAACACCAAGTGTCAGACCCCACTGGGAGCCATCAATTCCAGCTTGCCCTATCAGAACATTCACCCCG TGACTATAGGAGAGTGCCCTAAATATGTCCGCAGCGCCAAACTGCGAATGGTTACCGGCCTGCGCAACAATCCTTCGATCCAGTC ACGGGGCCTGTTCGGGGCCATAGCTGGCTTCATTGAGGGGGGATGGACCGGTATGATTGATGGGTGGTACGGCTACCACCATCA GAACGAGCAGGGTTCAGGCTACGCGGCTGACCAGAAATCCACTCAGAACGCTATCAATGGCATCACAAACAAGGTCAATACTGTT ATAGAAAAAATGAATATCCAATTCACTGCTGTGGGTAAAGAGTTTAACAAGCTGGAGAAAAGAATGGAAAATCTGAACAAAAAAGTG GATGACGGCTTCTTGGATATCTGGACATACAACGCTGAACTTCTGGTGCTGTTGGAAAACGAAAGGACTCTGGATTTCCACGATTC CAACGTGAAAAATCTCTACGAAAAAGTTAAGTCCCAACTGAAGAATAACGCGAAGGAGATCGGAAACGGATGCTTTGAGTTCTATC ATAAGTGTGATAATGAGTGTATGGAATCAGTTAGAAACGGTACGTACGATTACCCAAAATACTCCGAAGAGTCTAAGCTGAATCGC GAGAAAGTCGATGGGGTCAAACTCGAATCCATGGGAATTTACCAGATCTTAGCTATTTACAGTACAGTGGCCAGCTCTCTTGTCTT ACTGGTGTCACTTGGCGCGATTAGTTTCTGGATGTGCTCCAATGGCAGCCTCCAGTGTCGGATTTGCATA
2633	ATGAAGGCCAACCTGCTTGTACTGCTATGCGCCTTGGCTGCTGCAGACGCCGACACAATTTGCATCGGCTATCATGCCAACAATTC TACAGATACAGTGGACACTGTGCTGGAGAAAAACGTTACTGTCACTCATAGCGTTAATCTCCTCGAGGACTCTCACAATGGAAAAC TCTGTCGTCTGAAAGGCATCGCGCCACTGCAGTTGGGTAAATGTAATATCGCCGGCTGGTTGTTGGGCAACCCCGAATGTGACCC CCTCCTCCCAGTGAGGTCCTGGTCTTATATCGTAGAGACACCTAATTCGGAGAATGGAATTTGTTACCCTGGCGACTTCATAGACT ATGAGGAACTGAGGGAGCAGCTTTCTTCTGTGAGTTCTTTTGAAAGATTTGAGATCTTCCCTAAAGAATCATCTTGGCCCAACCATA ATACCAACGGCGTTACAGCAGCCTGTTCCCACGAGGGCAAGTCGAGTTTCTACAGGAATTTGTTATGGCTGACTGAGAAAGAGGG GTCTTATCCAAAGCTGAAAAACAGCTATGTCAACAAGAAAGGCAAAGAGGTTTTGGTGCTGTGGGGAATCCATCACCCACCAAACT CGAAGGAACAGCAAAATCTTTACCAAAACGAAAATGCATACGTATCAGTGGTTACTAGCAATTATAATCGGCGCTTTACGCCAGAG ATTGCCGAAAGACCTAAGGTGAGAGACCAAGCAGGAAGGATGAATTACTACTGGACACTACTGAAACCAGGCGATACTATCATTTT TGAGGCCAACGGAAACCTGATCGCACCCATGTACGCCTTCGCACTGTCGCGGGGTTTTGGGTCTGGCATCATCACCTCAAATGCC AGCATGCATGAGTGCAATACTAAATGCCAAACACCTTTAGGTGCCATTAATAGCTCACTCCCCTACCAGAACATCCATCCAGTAACT ATCGGGGAGTGTCCCAAGTACGTGCGATCTGCAAAACTGAGGATGGTCACTGGTTTGCGCAACAATCCCTCTATCCAGTCAAGGG GCCTATTTGGCGCAATTGCGGGCTTTATCGAGGGAGGGTGGACGGGTATGATTGACGGATGGTACGGCTACCACCACCAGAATG AACAGGGTTCGGGGTACGCGGCAGATCAGAAGAGTACTCAGAACGCAATTAATGGTATAACTAATAAGGTCAACACCGTAATTGAA AAAATGAATATCCAATTCACAGCTGTTGGGAAAGAATTTAACAAACTGGAGAAGAGGATGGAGAATCTCAATAAGAAGGTTGATGA CGGCTTTCTCGATATCTGGACTTACAATGCTGAGCTGCTAGTTCTGCTCGAGAACGAACGGACACTCGATTTTCACGATTCCAACG TTAAAAACCTGTACGAAAAGGTCAAATCTCAGCTGAAGAATAATGCTAAGGAAATTGGTAATGGCTGTTTCGAATTTTATCATAAAT GTGATAATGAGTGCATGGAAAGTGTCCGTAACGGCACCTACGATTATCCAAAGTATAGCGAGGAAAGCAAGCTTAATCGTGAAAAA GTAGATGGCGTGAAGCTGGAAAGTATGGGCATATATCAGATCCTGGCCATATACAGTACTGTTGCCTCCAGTCTAGTTTTACTGGT ATCCCTCGGAGCTATCTCATTCTGGATGTGTAGCAATGGCAGCCTTCAGTGCCGTATCTGTATA

2634	ATGAAAGCTAACCTGCTAGTTCTACTTTGTGCCCTTGCCGCTGCCGACGCTGACACCATCTGCATAGGGTACCACGCTAACAACAG TACTGACACAGTGGATACCGTTCTTGAAAAGAACGTCACCGTAACTCACTCTGTGAATTTGCTCGAGGATTCACACAATGGGAAAC TCTGCAGACTGAAAGGAATTGCTCCACTTCAGTTGGGCAAGTGTAACATCGCTGGCTGGCTGCTTGGGAATCCGGAGTGCGACCC ATTGCTGCCTGTGCGCAGTTGGAGCTACATTGTGGAGACCCCCAACAGCGAGAATGGCATCTGTTATCCCGGAGATTTCATCGAC TACGAGGAGCTCCGAGAACAATTATCCTCTGTGTCTAGTTTCGAACGGTTTGAGATCTTTCCCAAAGAATCAAGCTGGCCCAACCA CAACACAAACGGTGTGACAGCAGCATGCAGTCATGAGGGCAAGTCGTCCTTCTATAGGAATCTGCTGTGGCTTACTGAGAAGGAG GGTAGCTACCCGAAACTCAAGAACTCCTACGTCAATAAGAAGGGCAAAGAGGTGTTAGTCTTGTGGGGAATACACCACCCACCCA ACTCTAAAGAGCAACAAAACCTGTACCAGAACGAAAACGCCTACGTCTCCGTAGTAACCTCAAATTATAACAGACGCTTTACGCCC GAGATCGCAGAGCGCCCCAAGGTCAGGGACCAGGCAGGGAGAATGAACTACTATTGGACTTTGTTGAAGCCCGGAGACACAATT ATCTTCGAGGCGAATGGCAATCTGATTGCCCCCATGTACGCTTTTGCACTGTCCCGCGGTTTTGGTTCCGGCATCATAACTTCCAA CGCATCGATGCACGAATGTAACACCAAATGTCAGACACCATTGGGCGCCATAAACTCCAGTCTCCCATATCAAAATATCCACCCAG TCACAATCGGGGAATGTCCTAAATACGTGAGGTCAGCAAAGCTTAGGATGGTCACGGGACTGCGCAACAATCCCTCAATACAGTC CAGAGGGCTTTTCGGAGCCATAGCCGGCTTCATCGAAGGCGGCTGGACTGGGATGATCGATGGCTGGTATGGGTATCATCACCA GAATGAGCAGGGCTCCGGCTATGCTGCCGATCAGAAATCCACACAGAACGCTATCAATGGGATAACTAACAAAGTTAACACCGTG ATTGAGAAGATGAACATACAATTTACTGCAGTGGGAAAGGAGTTCAACAAACTGGAGAAGCGGATGGAGAACTTAAATAAAAAGGT GGATGACGGCTTCCTGGACATCTGGACTTATAACGCTGAACTGTTGGTACTGCTTGAGAATGAGCGGACCCTCGACTTTCACGAC TCAAACGTGAAGAATCTCTATGAGAAGGTAAAATCCCAGTTGAAAAACAACGCCAAGGAGATCGGGAATGGCTGCTTTGAATTTTA CCACAAGTGCGACAATGAGTGTATGGAAAGTGTGCGCAACGGAACATATGACTATCCTAAATATAGTGAAGAGTCCAAACTGAACC GTGAAAAGGTGGATGGGGTGAAACTCGAGTCCATGGGAATCTACCAAATCCTCGCCATTTATTCAACCGTGGCAAGCAGCCTCGT GCTACTGGTCAGTCTCGGCGCGATTTCTTTCTGGATGTGCTCCAACGGGTCACTGCAGTGTCGCATATGCATC
2635	ATGAAGGCGAACCTCCTTGTGCTTCTCTGCGCTCTCGCAGCTGCTGATGCAGACACGATTTGCATAGGGTATCATGCAAACAATAG TACAGACACTGTGGACACCGTGCTGGAGAAAAACGTGACTGTCACGCACTCAGTTAATCTTTTGGAGGACTCCCATAATGGAAAAC TGTGCAGACTTAAGGGGATCGCACCCCTGCAGTTAGGCAAGTGTAATATAGCCGGGTGGCTTCTGGGGAACCCAGAATGTGACC CACTGTTACCTGTGCGAAGCTGGTCTTACATAGTGGAGACGCCAAATTCTGAGAACGGCATCTGTTACCCTGGGGACTTTATTGAT TATGAGGAACTCCGGGAACAACTCTCGAGCGTTAGCAGCTTCGAAAGGTTTGAAATTTTTCCCAAGGAATCTTCATGGCCTAACCA CAACACTAACGGTGTGACGGCTGCTTGCTCACATGAGGGAAAGTCCAGTTTCTATAGGAACCTGTTGTGGCTGACTGAGAAAGAA GGCAGCTACCCAAAGCTGAAGAACTCTTACGTTAACAAAAAAGGCAAGGAAGTGTTAGTACTGTGGGGTATCCACCACCCACCAA ACAGCAAGGAGCAGCAGAACTTGTACCAGAACGAAAATGCCTACGTCAGTGTTGTCACATCAAACTACAATAGAAGATTTACGCCA GAAATTGCCGAAAGGCCAAAGGTGCGGGACCAAGCTGGGCGGATGAACTATTATTGGACCCTTCTGAAGCCGGGAGACACGATC ATATTCGAAGCAAACGGGAATCTGATCGCTCCGATGTACGCCTTCGCACTCAGCCGGGGTTTTGGATCAGGGATCATCACATCTAA CGCCTCTATGCATGAATGTAACACGAAGTGCCAGACTCCTCTAGGCGCTATCAACTCTAGCTTACCCTACCAGAATATACATCCCG TCACCATCGGCGAGTGTCCTAAGTATGTGCGCAGCGCCAAACTGCGGATGGTGACTGGCCTGAGAAATAATCCGTCCATCCAGTC AAGAGGGCTGTTTGGAGCAATCGCCGGCTTCATTGAAGGGGGCTGGACAGGTATGATTGATGGTTGGTACGGATACCACCATCAG AATGAACAGGGGAGTGGCTACGCAGCAGATCAAAAGTCGACACAAAACGCGATAAACGGAATCACAAACAAGGTCAATACCGTGA TTGAAAAGATGAATATTCAGTTCACCGCCGTCGGCAAAGAATTTAACAAGCTTGAGAAACGTATGGAAAATCTCAATAAGAAAGTTG ACGACGGTTTCTTAGACATCTGGACCTATAACGCAGAACTGCTGGTGCTACTGGAAAACGAGAGAACCCTTGATTTCCATGATAGC AACGTGAAAAACCTTTACGAGAAAGTTAAGAGTCAGCTCAAAAACAATGCTAAAGAAATCGGAAACGGCTGCTTCGAGTTTTATCAT AAGTGCGACAATGAGTGTATGGAGAGTGTTCGAAACGGGACATATGATTACCCGAAGTATTCAGAGGAGAGCAAACTCAATCGCG AAAAAGTTGATGGTGTGAAGCTGGAATCTATGGGCATCTACCAGATACTGGCCATCTACTCAACAGTGGCAAGTTCTCTGGTATTG CTCGTTTCCCTGGGAGCTATCTCCTTCTGGATGTGTTCGAACGGCTCACTTCAGTGTAGGATTTGCATA

2636	ATGAAGGCCAACCTGTTGGTTCTCTTGTGTGCTCTGGCAGCCGCAGACGCTGACACTATTTGTATCGGATACCACGCCAATAATAG TACTGACACCGTTGACACAGTACTTGAGAAGAACGTGACAGTAACCCACTCAGTGAACCTCCTTGAGGATAGCCACAATGGAAAAT TATGTAGACTTAAGGGTATCGCTCCCCTCCAGCTGGGCAAATGTAATATTGCCGGCTGGCTACTGGGGAATCCGGAATGTGATCC TCTCCTGCCAGTCCGATCATGGAGTTATATAGTAGAAACTCCAAATAGCGAGAATGGAATTTGTTACCCCGGCGATTTTATTGACTA CGAAGAGCTCCGAGAACAGCTCAGTTCAGTGTCTTCATTCGAACGGTTTGAAATCTTCCCAAAAGAATCCTCATGGCCCAATCACA ACACTAACGGGGTCACGGCAGCATGCAGCCACGAAGGGAAGTCGTCTTTCTACCGAAATCTCCTGTGGCTGACCGAGAAAGAGG GTTCGTACCCCAAGCTCAAGAATTCTTACGTTAACAAGAAGGGGAAGGAAGTACTTGTGCTGTGGGGTATACACCATCCACCTAAT TCCAAAGAGCAGCAGAACCTGTATCAAAACGAAAATGCATACGTCAGCGTGGTTACAAGTAACTACAACAGGCGCTTTACCCCAGA GATCGCCGAGAGGCCTAAGGTTCGGGATCAGGCCGGGAGAATGAACTACTATTGGACTCTCCTCAAGCCCGGTGACACAATAATC TTCGAGGCCAATGGAAATCTTATAGCACCGATGTATGCCTTCGCACTGTCACGCGGGTTTGGCTCAGGCATTATAACATCTAACGC TAGTATGCATGAGTGCAATACCAAGTGCCAGACCCCACTGGGTGCAATCAACAGCTCCTTGCCTTACCAGAACATACACCCAGTCA CTATTGGCGAGTGCCCGAAATACGTACGCTCTGCCAAGCTGCGTATGGTCACCGGACTCCGCAATAACCCCAGTATCCAAAGCCG AGGTTTGTTTGGCGCTATTGCTGGATTCATTGAAGGAGGCTGGACGGGAATGATTGACGGGTGGTACGGCTATCATCATCAGAAT GAGCAAGGGTCAGGTTACGCCGCCGACCAGAAGAGTACACAGAATGCAATCAACGGAATAACCAATAAGGTCAACACCGTCATCG AAAAGATGAATATCCAGTTCACGGCAGTGGGAAAGGAATTCAATAAGCTAGAAAAACGCATGGAGAATCTCAACAAGAAGGTTGAC GATGGGTTTCTGGATATCTGGACATACAACGCTGAGCTGCTTGTGCTGCTGGAGAACGAGAGGACACTGGACTTCCACGACTCTA ATGTCAAGAATCTTTATGAAAAAGTCAAATCCCAGCTCAAAAATAATGCAAAGGAGATCGGAAATGGCTGCTTCGAATTCTACCACA AGTGTGACAACGAGTGCATGGAGTCTGTGCGGAACGGAACCTACGACTATCCAAAGTATTCTGAGGAATCAAAACTCAATCGAGA GAAAGTTGATGGGGTAAAACTGGAATCTATGGGGATTTACCAGATCCTGGCCATCTACTCCACAGTGGCTAGTTCCTTGGTTCTGT TGGTGTCTCTAGGTGCGATCTCCTTCTGGATGTGTTCCAATGGGAGTCTTCAATGTCGAATCTGCATA
2637	ATGAAGGCAAATTTGCTTGTTTTACTGTGTGCTTTGGCTGCGGCTGATGCCGACACCATATGCATCGGTTACCACGCTAATAACAG CACCGACACAGTGGATACGGTGCTGGAGAAAAATGTGACCGTAACTCACAGTGTGAACCTGCTTGAGGACAGCCACAACGGTAAG CTCTGCCGTCTGAAAGGAATCGCCCCGCTGCAATTGGGAAAGTGCAACATTGCTGGCTGGCTCCTGGGTAATCCTGAGTGTGACC CGCTGCTGCCCGTGCGAAGCTGGTCCTACATCGTGGAGACTCCAAACTCAGAGAACGGGATCTGTTACCCTGGAGATTTTATAGA CTATGAAGAGCTAAGGGAACAATTGTCGTCTGTCAGTTCCTTTGAACGGTTCGAGATTTTCCCAAAGGAGTCCAGTTGGCCTAATC ATAATACCAATGGTGTCACTGCTGCATGTAGTCACGAAGGCAAGTCCTCATTTTATCGGAATTTGCTGTGGCTGACTGAGAAGGAG GGCAGCTATCCGAAGTTGAAGAATTCATACGTTAATAAAAAAGGTAAGGAAGTGCTTGTCCTTTGGGGCATTCATCATCCCCCTAA CAGTAAGGAACAGCAGAATCTCTACCAGAACGAGAACGCATACGTGTCCGTGGTTACTAGTAATTACAATAGGAGGTTCACACCAG AGATTGCGGAACGGCCTAAAGTCCGGGATCAGGCCGGTAGAATGAACTATTATTGGACTTTGCTGAAACCCGGAGACACAATCAT ATTCGAAGCTAATGGTAACCTGATTGCTCCTATGTACGCATTCGCTCTGTCCCGTGGCTTCGGCTCCGGGATTATCACCTCTAATG CTAGCATGCACGAGTGTAACACCAAATGCCAGACCCCACTCGGGGCGATTAACAGCAGCCTCCCATACCAGAACATTCATCCCGT TACTATAGGCGAGTGCCCAAAATATGTTCGGTCTGCAAAACTGCGGATGGTGACGGGGCTGCGAAATAACCCTTCAATACAGAGT AGGGGGCTGTTCGGAGCAATAGCTGGTTTTATTGAGGGCGGATGGACTGGCATGATCGATGGCTGGTATGGATACCATCACCAAA ACGAGCAAGGATCTGGTTATGCAGCGGACCAGAAGTCAACTCAGAATGCCATTAACGGCATTACTAATAAAGTTAATACCGTTATT GAAAAGATGAACATTCAGTTTACCGCCGTAGGGAAGGAGTTTAACAAGCTTGAAAAGCGCATGGAGAACCTCAACAAGAAAGTGG ATGATGGGTTCCTTGACATCTGGACATATAACGCCGAATTACTAGTGCTTTTGGAAAACGAAAGAACACTGGATTTCCATGACTCCA ATGTGAAGAATTTATATGAAAAGGTCAAAAGTCAGCTCAAGAATAATGCAAAGGAAATCGGCAATGGTTGCTTCGAATTTTACCATA AGTGTGACAACGAGTGTATGGAGAGTGTAAGGAATGGGACATACGATTACCCAAAGTATAGCGAGGAATCAAAGCTGAACAGAGA GAAAGTGGACGGCGTGAAATTGGAGTCCATGGGGATCTACCAGATTCTGGCGATCTATAGCACCGTCGCCAGCAGCTTGGTCTTG CTGGTATCTCTCGGGGCCATTTCCTTTTGGATGTGTTCAAATGGGAGCCTCCAATGTAGAATCTGTATC

2638	ATGAAAGCCAATCTACTGGTGTTGCTCTGTGCCCTGGCTGCTGCCGATGCTGACACGATATGTATCGGCTACCACGCAAACAACT CCACCGATACAGTTGATACCGTGCTCGAAAAGAATGTCACCGTAACACATAGCGTGAACCTTTTGGAGGATTCGCACAACGGTAAG CTGTGTCGTTTAAAGGGCATCGCCCCTCTTCAGCTGGGAAAATGCAATATTGCGGGCTGGTTGCTCGGTAACCCCGAGTGTGACC CTCTGCTCCCCGTGCGTAGTTGGTCATATATTGTGGAGACACCCAATAGCGAGAATGGAATTTGTTACCCCGGGGACTTTATCGAT TACGAAGAGCTGCGGGAACAACTGTCTAGTGTCAGCTCGTTCGAACGCTTTGAGATTTTCCCAAAAGAGAGCTCCTGGCCCAACC ATAACACTAACGGGGTGACTGCTGCCTGTTCCCATGAGGGGAAGTCTAGTTTTTACCGAAATCTGCTGTGGCTAACGGAGAAAGA GGGCTCTTATCCAAAGCTAAAGAATAGTTACGTAAACAAGAAAGGAAAAGAGGTGCTCGTTCTCTGGGGCATTCATCACCCACCAA ACTCCAAGGAGCAGCAGAATCTGTATCAGAATGAAAACGCCTACGTATCCGTGGTGACAAGCAACTACAACAGGCGATTCACGCC CGAGATCGCCGAGCGGCCTAAAGTGAGGGATCAGGCTGGCAGGATGAACTACTATTGGACGCTGCTGAAGCCAGGGGATACTAT TATTTTTGAGGCGAACGGAAACTTAATTGCCCCCATGTACGCATTTGCGCTCTCCAGAGGATTCGGCTCTGGAATCATCACGTCAA ACGCTAGCATGCACGAGTGTAATACAAAGTGTCAGACCCCACTGGGGGCAATCAATAGCAGCTTACCTTACCAAAACATCCATCCA GTGACTATCGGTGAATGCCCGAAGTACGTAAGGTCCGCAAAGCTGAGAATGGTGACGGGCCTCAGAAATAACCCTAGTATACAGA GTCGTGGCCTGTTTGGGGCCATTGCTGGATTTATCGAAGGAGGGTGGACCGGAATGATAGACGGCTGGTATGGCTACCACCACC AGAATGAGCAGGGATCTGGGTACGCTGCTGATCAGAAGAGCACGCAGAACGCCATTAACGGCATAACCAACAAGGTGAATACAGT CATCGAAAAAATGAACATACAGTTTACAGCAGTTGGGAAAGAATTCAACAAACTGGAGAAGAGGATGGAAAACTTGAACAAAAAAG TCGATGACGGGTTTCTCGACATCTGGACTTACAACGCCGAGCTGCTTGTGTTACTGGAGAACGAAAGGACACTCGACTTTCACGA CTCTAACGTGAAGAATCTTTATGAGAAAGTGAAAAGCCAACTGAAAAACAATGCCAAAGAAATTGGTAACGGCTGCTTCGAGTTTTA CCATAAGTGTGACAACGAGTGTATGGAATCAGTCCGCAACGGTACCTACGACTATCCAAAATACTCTGAAGAATCAAAGCTGAATA GGGAGAAGGTGGACGGCGTTAAATTGGAGAGCATGGGCATTTATCAAATCCTTGCCATTTATTCGACAGTGGCCTCCTCTTTAGTG CTCTTAGTGTCCCTGGGCGCCATCAGTTTCTGGATGTGTAGTAACGGCTCACTCCAATGTCGAATTTGCATT
2639	ATGAAAGCGAACCTCCTGGTTTTATTGTGCGCACTTGCCGCTGCTGATGCAGACACCATCTGTATTGGGTATCACGCGAACAATTC GACCGACACGGTCGATACAGTGCTTGAGAAAAATGTGACTGTAACACATTCCGTGAATCTCCTCGAAGACAGCCATAACGGGAAG CTTTGCAGGCTTAAGGGGATTGCACCGCTGCAGCTCGGGAAATGTAATATAGCCGGGTGGCTATTAGGGAACCCGGAGTGCGAC CCTTTGCTTCCAGTGAGGTCCTGGTCATATATTGTCGAGACCCCAAATTCAGAAAACGGGATCTGTTACCCCGGCGACTTCATTGA CTACGAAGAGCTGAGGGAACAATTAAGTTCTGTCTCCTCTTTTGAACGGTTTGAGATATTCCCAAAAGAGTCCTCCTGGCCCAATC ATAACACGAATGGCGTGACAGCTGCGTGTAGCCATGAAGGGAAGAGCAGTTTCTACAGGAATCTTCTTTGGTTAACAGAAAAAGAA GGTAGTTATCCGAAACTGAAAAACAGCTACGTAAATAAAAAAGGCAAGGAGGTCCTGGTGTTGTGGGGCATTCATCACCCTCCCAA TAGCAAGGAACAGCAGAATCTCTATCAGAATGAAAACGCCTATGTCAGCGTTGTTACTAGCAACTACAACCGGCGGTTTACCCCCG AAATTGCTGAGCGACCCAAAGTTCGCGATCAGGCCGGTCGGATGAATTACTACTGGACCCTTCTGAAGCCTGGCGACACCATTAT TTTTGAGGCCAACGGCAACCTAATTGCCCCAATGTATGCATTCGCCCTGAGCCGAGGTTTTGGAAGTGGCATAATCACGTCCAATG CCTCTATGCACGAATGCAACACCAAGTGCCAGACACCCCTTGGCGCTATCAACTCTTCTCTACCTTATCAGAACATCCATCCCGTT ACTATTGGGGAGTGCCCCAAATACGTCCGGTCTGCTAAACTTCGGATGGTGACAGGCCTCCGCAATAATCCTTCGATTCAAAGTC GGGGCCTGTTTGGTGCAATCGCCGGGTTCATTGAGGGAGGGTGGACTGGCATGATCGACGGTTGGTATGGGTACCACCATCAGA ACGAACAAGGCAGTGGGTACGCTGCCGATCAGAAAAGCACCCAGAATGCTATCAACGGCATTACGAACAAGGTGAACACTGTGAT AGAGAAGATGAACATCCAGTTCACTGCTGTCGGTAAAGAGTTTAACAAACTTGAAAAAAGAATGGAGAACCTCAATAAAAAGGTGG ACGATGGTTTCCTGGATATATGGACATATAACGCCGAACTACTTGTGCTCTTGGAAAATGAACGGACACTGGACTTCCACGACTCT AATGTGAAAAATTTATATGAGAAAGTGAAGTCACAGCTCAAGAATAACGCGAAGGAAATTGGCAACGGGTGTTTCGAGTTCTATCA TAAGTGCGACAATGAGTGCATGGAATCAGTTCGAAATGGAACCTATGACTATCCAAAGTACTCAGAAGAGAGTAAACTGAACCGGG AGAAAGTCGACGGAGTCAAGCTCGAATCTATGGGTATTTATCAGATTCTGGCCATCTACAGCACAGTCGCTTCTTCTCTTGTACTG CTGGTTTCTTTGGGCGCAATCTCTTTCTGGATGTGCTCAAACGGGAGCCTGCAGTGCCGGATCTGCATC

2640	ATGAAGGCTAATCTGTTAGTACTTTTATGTGCACTTGCCGCAGCTGACGCGGATACGATCTGCATTGGGTACCATGCCAACAACAG TACCGACACCGTCGATACCGTTTTGGAAAAGAATGTGACCGTGACCCACAGCGTCAACCTTCTCGAGGATAGTCACAATGGGAAA CTCTGTCGGCTGAAAGGCATAGCTCCTCTGCAGCTCGGAAAATGTAACATCGCCGGCTGGTTATTGGGGAACCCCGAATGCGATC CCCTCCTCCCAGTAAGGTCTTGGTCCTATATCGTGGAAACACCTAACAGCGAGAATGGCATCTGCTACCCCGGGGACTTCATTGAT TACGAAGAACTAAGAGAACAACTGTCGAGTGTGAGTAGTTTCGAAAGATTTGAAATATTCCCTAAGGAGTCAAGTTGGCCCAATCA CAACACGAACGGCGTTACGGCAGCGTGCTCCCATGAAGGGAAATCAAGCTTCTATAGAAATCTGCTCTGGCTAACAGAAAAGGAG GGCTCATACCCAAAGTTGAAGAACTCCTACGTCAATAAAAAAGGTAAGGAAGTATTGGTCCTGTGGGGGATCCATCACCCCCCAAA TTCGAAAGAGCAACAGAACCTTTATCAGAATGAGAATGCTTACGTTAGCGTAGTTACGAGCAACTATAATCGGCGATTCACACCCG AGATTGCTGAGAGACCTAAGGTGAGAGATCAGGCAGGTCGGATGAACTACTACTGGACTCTTCTAAAGCCCGGCGATACTATAAT CTTTGAGGCTAACGGCAATCTGATTGCCCCTATGTATGCTTTCGCATTGTCCAGAGGTTTTGGGAGTGGGATCATCACGAGCAATG CCAGTATGCATGAGTGTAACACTAAATGTCAGACACCCTTAGGGGCTATCAACAGTTCTCTTCCCTATCAGAATATCCACCCAGTAA CCATCGGAGAATGCCCTAAATACGTCAGATCTGCAAAACTCAGGATGGTGACTGGCCTCAGAAATAACCCCAGTATACAGAGCAG AGGTTTATTCGGGGCGATCGCTGGCTTTATAGAAGGAGGCTGGACGGGCATGATTGACGGCTGGTATGGCTATCATCATCAGAAT GAACAGGGCAGCGGTTACGCAGCCGACCAGAAGAGCACACAAAACGCTATTAACGGAATCACCAACAAGGTGAATACAGTCATTG AAAAAATGAATATTCAATTCACTGCAGTCGGAAAGGAATTCAACAAACTGGAAAAAAGGATGGAGAATCTCAACAAAAAAGTCGAC GACGGGTTCCTGGATATCTGGACATACAATGCCGAACTCCTCGTCCTGCTCGAGAATGAGCGGACCCTTGATTTCCACGATTCCA ATGTCAAGAACCTTTACGAGAAGGTGAAATCCCAGTTAAAGAATAACGCCAAAGAGATTGGCAATGGGTGCTTCGAATTTTATCAC AAGTGCGACAACGAGTGTATGGAAAGCGTCAGAAATGGCACCTACGACTATCCTAAATATTCAGAAGAGTCTAAACTGAATAGGGA AAAAGTCGACGGCGTCAAGCTTGAGTCTATGGGGATATATCAGATTCTTGCGATTTATTCAACTGTCGCATCCTCCCTCGTTCTGCT CGTGTCCCTGGGCGCAATCTCATTCTGGATGTGCTCTAATGGTTCATTGCAGTGTCGCATCTGTATC
2641	ATGAAGGCAAACCTGCTGGTTCTTCTGTGTGCATTAGCCGCGGCCGACGCCGATACAATATGTATCGGATACCACGCAAATAACA GCACCGACACCGTTGACACAGTGCTTGAGAAAAACGTGACCGTGACACACAGTGTGAATCTGCTAGAGGACTCCCATAACGGGAA GCTGTGCCGGTTGAAGGGAATCGCTCCGCTGCAGTTGGGCAAATGCAATATTGCTGGGTGGCTGTTAGGTAACCCTGAATGCGAT CCACTCCTCCCTGTACGTTCTTGGTCGTATATAGTGGAAACTCCCAATAGCGAAAATGGTATTTGTTATCCTGGGGATTTCATAGAT TACGAAGAGCTGCGCGAACAGCTCAGCTCTGTGAGCTCTTTCGAGCGCTTCGAAATATTCCCCAAGGAGTCCAGTTGGCCCAACC ACAATACCAACGGCGTTACCGCAGCCTGTAGCCACGAGGGCAAGTCCAGCTTTTACCGCAACCTCTTGTGGTTGACCGAAAAGGA GGGAAGCTATCCGAAACTCAAGAACAGCTACGTCAATAAAAAAGGCAAGGAGGTACTAGTGCTGTGGGGCATCCACCACCCTCCA AATAGTAAAGAGCAGCAGAATCTGTATCAGAACGAAAACGCGTATGTCTCAGTTGTGACAAGCAATTACAACCGCAGGTTCACTCC GGAGATAGCTGAAAGACCCAAAGTACGGGATCAGGCTGGGCGAATGAACTACTACTGGACACTACTCAAACCAGGTGACACCATT ATTTTCGAAGCAAATGGCAATTTGATCGCACCCATGTACGCATTTGCCTTGTCCCGGGGGTTCGGCTCCGGCATTATTACGTCTAA CGCATCTATGCACGAGTGCAACACGAAATGCCAGACTCCCCTGGGGGCCATAAACTCAAGCCTGCCATATCAGAACATCCACCCG GTCACCATCGGCGAGTGCCCCAAATACGTCCGCTCAGCTAAACTCAGGATGGTCACGGGCCTCAGAAATAATCCTAGCATTCAGT CTCGCGGTCTGTTTGGCGCCATCGCAGGCTTCATTGAAGGGGGCTGGACCGGAATGATCGATGGGTGGTACGGGTATCATCACC AGAACGAGCAAGGCAGTGGCTATGCTGCCGATCAGAAATCAACCCAGAATGCGATTAACGGAATAACTAACAAGGTGAACACCGT CATTGAGAAGATGAATATTCAGTTCACGGCCGTCGGAAAGGAGTTTAACAAGCTCGAAAAGCGCATGGAGAACCTGAACAAGAAG GTAGATGATGGCTTCCTGGACATTTGGACCTACAATGCTGAGCTCCTCGTCTTGCTGGAGAATGAACGGACTCTCGACTTCCACGA TTCCAACGTTAAGAATTTATATGAGAAAGTGAAGAGTCAACTTAAAAACAACGCGAAGGAAATCGGTAATGGCTGCTTTGAGTTCTA CCATAAGTGTGATAACGAGTGTATGGAAAGCGTGCGAAATGGGACTTATGACTACCCTAAGTATTCGGAAGAATCTAAACTGAATA GAGAGAAGGTGGATGGGGTGAAGCTCGAATCTATGGGGATTTACCAGATCTTGGCTATCTATTCGACAGTGGCTTCCAGTCTTGT CCTACTGGTGTCTCTTGGCGCCATTTCCTTCTGGATGTGCTCCAACGGATCTCTGCAGTGCCGCATCTGTATA

2642	ATGAAGGCAAATTTGTTAGTCTTGTTGTGCGCTCTGGCCGCTGCAGATGCAGACACCATCTGTATCGGGTATCACGCAAATAATTC CACAGATACAGTGGACACCGTTCTCGAGAAAAACGTCACCGTGACCCATAGCGTGAACCTTCTCGAGGACTCTCACAACGGTAAA TTATGCAGGCTTAAGGGAATAGCTCCCCTGCAACTTGGTAAGTGCAATATAGCAGGATGGCTCCTGGGCAACCCAGAATGTGACC CGCTACTGCCAGTGAGGAGCTGGTCTTACATCGTCGAAACGCCTAACTCCGAAAACGGGATCTGTTATCCTGGCGATTTCATTGAT TATGAAGAGTTACGGGAGCAGTTATCAAGCGTGTCGAGCTTCGAAAGGTTTGAGATCTTCCCTAAGGAGTCAAGTTGGCCGAACC ACAACACGAATGGCGTCACAGCCGCATGTTCCCATGAAGGGAAGTCCTCTTTCTATCGGAATCTTTTGTGGCTAACGGAGAAGGA GGGATCCTATCCTAAGCTTAAAAACTCCTACGTAAACAAAAAGGGCAAGGAGGTCTTAGTACTGTGGGGCATACACCACCCACCTA ACTCAAAGGAGCAGCAGAACCTTTATCAGAATGAAAATGCTTACGTCTCGGTGGTGACCAGCAACTACAACAGACGCTTCACGCC GGAAATTGCCGAGAGACCAAAGGTAAGGGATCAAGCCGGTCGAATGAATTACTATTGGACCTTATTGAAACCCGGGGACACCATC ATATTTGAAGCTAACGGCAATTTGATCGCTCCAATGTACGCGTTTGCCCTTTCCCGCGGGTTCGGTTCTGGGATCATAACCTCTAA CGCTTCCATGCATGAATGTAACACAAAGTGCCAGACTCCATTAGGGGCCATTAATAGCAGTTTACCTTATCAAAATATCCACCCAGT CACTATCGGAGAGTGCCCAAAATACGTGAGGAGCGCGAAATTGAGGATGGTGACAGGCCTTCGGAACAATCCCTCTATCCAGAGC AGAGGACTGTTTGGCGCAATTGCTGGGTTTATAGAGGGGGGATGGACCGGCATGATTGACGGATGGTACGGATACCATCATCAGA ATGAGCAGGGAAGTGGATATGCTGCAGACCAAAAGTCAACTCAGAATGCTATTAATGGTATCACCAACAAGGTGAACACAGTCATA GAAAAGATGAACATCCAGTTTACCGCCGTTGGGAAGGAATTTAACAAGCTGGAAAAAAGGATGGAGAATCTCAATAAAAAGGTAGA CGATGGGTTCCTTGATATTTGGACTTACAATGCTGAGCTGTTAGTCCTACTAGAAAATGAGAGAACCCTTGACTTTCACGACTCCAA CGTTAAAAACTTATACGAGAAAGTGAAATCCCAATTAAAGAATAACGCGAAAGAGATCGGCAACGGGTGTTTCGAATTCTATCACAA ATGCGACAACGAATGCATGGAGTCTGTGAGGAATGGGACTTATGATTATCCCAAGTATTCAGAGGAGTCTAAACTGAATCGAGAAA AAGTGGACGGAGTAAAGCTGGAGTCCATGGGCATATATCAAATCCTCGCCATCTATTCCACAGTCGCCTCCTCCCTTGTTTTGTTG GTTTCCCTTGGTGCGATCAGTTTTTGGATGTGCTCTAACGGGTCCCTACAGTGCCGAATTTGTATT
2643	ATGAAAGCTAATCTTCTCGTGCTACTGTGCGCCTTAGCAGCTGCCGACGCTGATACCATCTGTATAGGCTATCATGCCAATAACTC CACAGACACCGTTGATACTGTGCTTGAGAAGAATGTGACGGTAACCCACAGCGTTAATCTCCTTGAGGACAGCCATAATGGCAAAC TGTGTCGGCTGAAGGGAATCGCTCCCCTACAATTGGGAAAATGCAACATTGCAGGATGGTTGTTAGGAAACCCAGAATGCGACCC TCTACTGCCAGTGCGCAGCTGGAGTTACATTGTTGAAACACCTAACAGCGAAAATGGTATTTGCTACCCTGGGGATTTCATAGATT ACGAGGAGTTAAGAGAGCAGCTGTCCAGTGTGTCTTCCTTCGAAAGGTTTGAAATCTTCCCAAAGGAGTCCTCATGGCCCAACCA CAACACGAATGGCGTCACAGCTGCCTGTTCACACGAGGGTAAGTCAAGTTTTTATAGGAACTTGCTTTGGTTAACCGAAAAAGAGG GAAGCTACCCAAAGCTGAAAAATAGCTACGTGAATAAAAAGGGCAAAGAGGTGCTGGTGCTATGGGGTATCCATCATCCGCCAAA CAGCAAAGAGCAGCAAAACCTGTACCAGAACGAAAATGCCTACGTGTCTGTTGTGACGAGCAACTACAACCGTAGATTTACCCCG GAAATCGCAGAGCGGCCGAAGGTGCGCGATCAGGCGGGACGTATGAATTACTATTGGACATTGCTTAAGCCCGGAGACACAATTA TTTTCGAAGCTAACGGGAATCTCATTGCCCCAATGTATGCTTTCGCACTGAGCCGAGGCTTTGGGAGCGGCATCATCACAAGTAAT GCTAGCATGCATGAGTGCAATACCAAGTGTCAAACGCCTCTGGGCGCAATCAATTCCAGTTTACCATATCAGAACATTCATCCAGT CACCATCGGGGAGTGCCCTAAATATGTAAGGTCTGCTAAACTGCGGATGGTCACTGGTCTCCGCAACAATCCATCTATCCAGAGC CGGGGTTTGTTCGGAGCTATTGCTGGATTTATAGAGGGGGGCTGGACGGGGATGATTGATGGCTGGTACGGCTATCACCATCAG AACGAACAGGGCAGCGGCTACGCTGCAGACCAAAAAAGTACTCAGAACGCTATTAATGGCATTACCAACAAGGTTAACACTGTGAT TGAGAAGATGAACATCCAGTTCACTGCGGTAGGAAAGGAATTCAACAAACTTGAGAAGCGGATGGAAAATCTCAACAAGAAGGTA GACGATGGGTTCCTAGATATATGGACCTATAATGCCGAGCTCTTGGTCTTGTTAGAAAACGAGCGGACCTTAGACTTCCATGATAG CAACGTGAAAAACCTGTACGAGAAAGTGAAGAGCCAGTTAAAGAATAACGCTAAGGAAATCGGAAACGGGTGCTTTGAGTTCTACC ATAAGTGTGACAATGAGTGTATGGAATCTGTGCGTAACGGGACCTACGACTACCCCAAATATTCTGAGGAATCAAAACTAAATCGA GAGAAGGTAGACGGCGTGAAGTTAGAATCTATGGGTATCTATCAAATCCTGGCTATATACTCAACCGTGGCCTCATCCCTAGTTCT CCTGGTATCCCTCGGCGCCATCTCTTTCTGGATGTGCAGCAACGGCTCTCTCCAGTGTCGCATTTGCATC

2644	ATGAAGGCCAATTTGCTGGTGTTGTTGTGTGCTCTGGCAGCAGCCGATGCTGACACTATCTGCATCGGCTACCATGCCAACAATA GTACCGATACTGTGGACACTGTGTTAGAGAAGAACGTGACAGTCACTCACTCAGTAAACTTGCTGGAGGATAGCCATAATGGCAAA CTCTGTCGCCTCAAGGGCATCGCCCCTCTGCAGCTGGGAAAATGCAACATTGCTGGATGGCTGCTGGGGAACCCGGAATGCGAC CCGCTGCTCCCCGTGAGATCCTGGAGTTATATAGTTGAAACTCCCAACAGTGAGAACGGGATATGTTATCCTGGCGACTTCATTGA CTACGAGGAGCTGCGCGAACAGCTCTCTAGCGTATCTTCGTTTGAACGTTTCGAAATATTTCCCAAGGAATCTTCTTGGCCAAATC ACAATACCAACGGAGTCACGGCGGCCTGCTCGCACGAGGGGAAGAGTTCCTTTTATCGCAACCTACTGTGGCTCACCGAGAAGG AGGGTTCCTACCCCAAGCTGAAAAACAGCTATGTGAACAAAAAAGGCAAAGAAGTCTTGGTACTGTGGGGCATTCATCACCCACC CAACAGCAAGGAACAGCAGAACCTGTATCAAAATGAAAATGCTTACGTCTCCGTTGTGACATCAAACTACAACAGACGGTTCACCC CTGAAATTGCTGAGAGGCCTAAGGTCAGGGACCAGGCCGGGCGGATGAATTACTACTGGACACTGCTGAAGCCGGGAGACACGA TTATCTTCGAAGCAAACGGCAATCTGATCGCCCCAATGTACGCGTTTGCTCTTTCTCGGGGATTTGGCTCTGGGATCATTACCTCT AATGCATCCATGCACGAATGCAACACCAAATGTCAGACCCCACTAGGCGCTATTAATAGTTCCCTGCCCTATCAGAATATCCACCC GGTGACCATCGGCGAATGTCCTAAATATGTGCGGAGTGCCAAGCTGAGGATGGTCACTGGCTTACGTAACAATCCCTCAATTCAG AGCCGGGGCCTCTTTGGAGCAATCGCTGGATTCATCGAGGGGGGTTGGACCGGAATGATCGACGGCTGGTACGGATATCATCAC CAGAATGAACAAGGGTCTGGCTACGCCGCCGACCAAAAATCTACCCAAAACGCTATTAATGGGATTACCAATAAGGTGAACACTGT AATAGAAAAAATGAATATTCAGTTTACTGCCGTTGGAAAGGAATTCAATAAGCTGGAGAAACGTATGGAGAATCTGAACAAGAAGG TAGATGACGGATTCCTGGACATCTGGACCTATAATGCAGAATTGCTCGTTCTGCTTGAAAATGAGCGCACACTGGACTTTCATGAC TCTAACGTCAAAAATCTGTATGAAAAGGTCAAGTCCCAGCTGAAAAATAATGCGAAAGAGATTGGGAACGGATGTTTCGAATTCTAC CACAAGTGTGACAACGAATGCATGGAGTCCGTACGGAACGGCACGTACGACTACCCAAAATATTCTGAGGAAAGCAAGCTGAATA GAGAAAAGGTCGATGGGGTGAAACTGGAGAGTATGGGAATCTATCAGATTCTCGCCATCTACTCCACTGTCGCCAGCTCCTTGGT ACTTCTTGTTAGCTTGGGAGCCATATCATTCTGGATGTGCTCCAATGGATCCCTCCAATGCAGAATCTGCATC
2645	ATGAAAGCTAACCTTTTGGTGCTGCTCTGTGCCCTGGCAGCTGCCGATGCCGATACCATCTGTATCGGCTACCATGCTAATAATAG TACAGACACAGTGGACACCGTTCTGGAGAAAAATGTCACCGTGACACATAGCGTGAACTTGCTGGAGGATTCTCACAACGGGAAG CTGTGCCGTCTGAAAGGGATTGCCCCTCTTCAGCTCGGTAAATGCAACATCGCCGGGTGGCTGCTAGGCAATCCAGAGTGTGATC CCCTGCTGCCGGTGAGAAGCTGGAGCTACATCGTGGAGACACCCAACTCCGAAAATGGCATTTGCTACCCCGGCGATTTCATCGA CTACGAAGAACTGCGGGAGCAGCTGTCGTCTGTGAGCTCCTTCGAGAGATTCGAGATCTTTCCAAAAGAAAGCAGTTGGCCCAAT CACAATACCAACGGCGTGACCGCTGCATGTAGTCACGAGGGCAAGAGCTCATTCTATAGAAACCTGTTGTGGCTGACAGAGAAGG AGGGGAGCTATCCCAAACTGAAGAATTCCTATGTGAATAAAAAGGGGAAGGAGGTTCTGGTATTATGGGGCATCCATCATCCTCCT AACAGCAAAGAGCAGCAAAACCTTTATCAGAACGAGAATGCCTATGTATCGGTTGTTACCAGCAATTATAATCGGAGGTTTACCCC AGAGATAGCTGAGCGTCCAAAAGTGAGAGACCAAGCTGGCCGGATGAATTACTATTGGACCCTCCTCAAACCGGGCGATACCATC ATTTTCGAAGCCAATGGCAACTTAATAGCGCCCATGTATGCTTTTGCCCTTTCGAGGGGATTTGGATCAGGTATCATCACAAGCAA CGCATCAATGCACGAGTGTAACACTAAGTGTCAGACTCCACTCGGGGCAATTAACAGCAGCCTACCTTACCAGAACATCCACCCC GTGACGATTGGAGAGTGTCCCAAGTACGTAAGGTCAGCCAAGCTCAGGATGGTTACAGGGCTGCGAAATAACCCCTCTATTCAGT CAAGGGGCTTGTTCGGCGCCATCGCCGGGTTTATCGAGGGCGGCTGGACCGGAATGATCGACGGATGGTATGGCTACCATCACC AGAACGAACAAGGGTCTGGCTATGCTGCGGATCAGAAGTCAACACAGAATGCTATAAATGGTATCACGAATAAGGTAAATACAGTC ATTGAAAAGATGAATATCCAGTTTACGGCCGTTGGGAAGGAGTTCAACAAGCTTGAAAAGAGAATGGAAAATCTGAATAAAAAAGT GGACGACGGCTTTTTGGATATCTGGACATATAACGCGGAGCTGCTAGTGCTCCTGGAAAACGAGCGGACTCTTGACTTTCACGAT TCCAACGTCAAGAACCTCTATGAAAAGGTTAAATCACAATTGAAGAATAATGCTAAGGAGATTGGCAATGGGTGCTTTGAGTTCTAC CATAAGTGTGACAACGAATGTATGGAGTCAGTCCGGAACGGTACTTACGATTACCCCAAGTATTCGGAAGAGTCAAAGTTGAATAG AGAAAAAGTCGACGGCGTGAAACTCGAGTCTATGGGCATTTACCAGATCCTTGCGATTTATAGCACCGTCGCCAGTAGCTTGGTAC TCCTAGTATCTCTGGGTGCAATCTCCTTTTGGATGTGTTCTAACGGCAGCTTACAGTGCAGGATCTGCATT

2646	ATGAAAGCTAACTTGCTTGTGTTGTTGTGCGCTTTGGCCGCTGCCGACGCCGACACTATTTGCATCGGGTATCACGCCAATAATTC CACAGATACCGTCGATACCGTGCTCGAGAAGAACGTGACCGTCACTCACTCAGTGAACCTTCTGGAGGACTCCCATAACGGGAAG CTCTGTAGGTTGAAAGGAATCGCCCCACTGCAGCTTGGTAAATGCAATATAGCCGGTTGGCTGCTGGGAAACCCTGAGTGCGACC CACTCCTCCCCGTACGGTCTTGGAGCTACATCGTGGAGACGCCAAATTCCGAGAATGGAATCTGTTATCCCGGCGACTTTATCGAT TATGAGGAGTTGCGCGAGCAGCTGAGTTCAGTTTCATCCTTTGAACGGTTCGAGATTTTTCCAAAAGAGTCTAGCTGGCCTAATCA TAATACCAATGGGGTTACAGCCGCTTGCTCACATGAGGGGAAGTCCAGTTTCTATAGGAACCTGCTTTGGCTGACCGAGAAAGAG GGCTCCTATCCAAAGCTCAAGAACAGCTATGTAAATAAGAAAGGCAAGGAAGTGTTGGTCCTGTGGGGCATACACCACCCTCCCA ACTCGAAGGAGCAGCAAAACCTATACCAGAACGAGAATGCCTACGTCAGCGTGGTCACTAGCAATTACAATAGAAGGTTCACACC CGAAATTGCTGAGCGGCCGAAAGTGAGAGACCAGGCAGGACGAATGAACTACTATTGGACCCTGCTGAAGCCCGGGGATACAAT CATTTTCGAGGCGAACGGGAACCTCATTGCCCCCATGTACGCTTTTGCCCTCAGCAGAGGCTTTGGGTCTGGTATAATCACCAGTA ACGCTAGCATGCATGAATGTAATACCAAGTGCCAGACACCCTTAGGGGCCATTAACTCATCTCTCCCATATCAGAACATACATCCT GTGACCATCGGTGAATGCCCGAAATACGTAAGGAGCGCAAAATTAAGGATGGTGACCGGTCTTAGGAACAATCCATCCATTCAGT CTCGAGGGCTTTTCGGCGCCATTGCTGGATTCATAGAAGGCGGATGGACAGGCATGATCGATGGATGGTACGGTTATCACCACCA GAATGAGCAGGGGTCGGGCTACGCTGCCGATCAGAAGTCTACCCAAAATGCCATCAATGGCATAACCAACAAGGTGAACACAGTT ATCGAGAAAATGAACATACAGTTTACTGCAGTGGGAAAGGAGTTCAACAAACTGGAGAAACGAATGGAGAACTTGAATAAAAAAGT CGACGACGGCTTCCTTGATATTTGGACTTATAATGCCGAGCTTTTAGTCCTGTTAGAGAATGAACGTACATTGGACTTCCACGATAG CAACGTAAAGAACCTCTATGAGAAAGTGAAAAGTCAGCTAAAAAATAATGCCAAAGAGATCGGGAATGGTTGTTTCGAGTTTTATCA CAAGTGTGATAATGAGTGCATGGAGTCTGTGAGAAACGGCACGTACGATTACCCTAAGTATTCTGAGGAGAGTAAGTTAAATCGCG AGAAAGTTGATGGTGTGAAGCTTGAGTCAATGGGCATCTACCAGATCCTGGCCATCTACTCCACCGTGGCGTCTTCTCTGGTGCT GCTTGTTAGCCTGGGTGCTATCTCCTTCTGGATGTGTTCTAACGGCTCCCTGCAGTGTAGGATCTGCATC
2647	ATGAAAGCAAATCTGCTGGTGTTGCTGTGTGCCCTGGCTGCCGCAGACGCAGATACCATCTGTATCGGCTATCACGCTAATAACA GTACAGACACTGTGGATACTGTTCTCGAGAAGAACGTCACTGTGACCCACTCAGTAAATTTACTCGAAGACAGTCACAACGGCAAG TTGTGTAGACTCAAAGGCATTGCCCCTCTCCAGCTGGGAAAGTGCAACATTGCTGGATGGCTCCTGGGGAACCCCGAATGTGATC CGCTTTTGCCTGTGAGAAGCTGGTCCTACATAGTAGAAACCCCAAACTCCGAAAATGGTATTTGCTATCCGGGCGATTTTATAGATT ATGAAGAATTAAGGGAGCAGTTGTCTTCCGTGTCCTCCTTCGAACGCTTCGAAATCTTCCCCAAAGAGAGTTCCTGGCCAAACCAT AATACTAACGGGGTGACAGCGGCTTGCTCTCACGAAGGAAAATCTTCCTTCTACCGGAACCTGCTTTGGTTGACCGAGAAGGAGG GCAGTTACCCGAAATTGAAAAATAGTTACGTTAATAAGAAAGGGAAGGAAGTCCTGGTGTTATGGGGTATTCATCACCCCCCAAAC AGCAAGGAGCAGCAAAATTTGTACCAGAACGAGAATGCGTACGTGAGTGTGGTCACCAGTAATTACAACCGGAGGTTTACCCCCG AGATCGCAGAACGCCCGAAAGTACGGGACCAGGCAGGGAGGATGAATTACTATTGGACTCTTTTAAAGCCCGGGGATACCATCAT CTTTGAGGCAAACGGTAACCTCATAGCTCCCATGTACGCCTTCGCCCTGAGCCGCGGTTTCGGCAGTGGCATTATAACCTCTAAC GCCAGCATGCACGAATGTAACACTAAATGTCAGACCCCTCTCGGCGCCATTAATAGCAGTCTCCCTTACCAAAATATCCACCCCGT TACCATTGGCGAGTGTCCAAAATACGTAAGGTCCGCAAAACTGCGCATGGTGACTGGACTGAGAAACAATCCCTCTATTCAGTCGA GGGGACTATTCGGAGCCATCGCTGGGTTTATCGAAGGGGGATGGACTGGAATGATCGATGGTTGGTACGGGTACCACCACCAGA ACGAGCAAGGCTCAGGGTATGCCGCAGACCAGAAGAGTACCCAGAACGCCATCAACGGTATCACTAATAAGGTGAATACCGTCAT CGAGAAGATGAATATCCAGTTTACCGCTGTGGGCAAAGAGTTTAATAAGCTGGAAAAGCGCATGGAGAACCTGAATAAGAAGGTT GACGACGGCTTCCTCGATATTTGGACTTATAACGCCGAGCTGCTCGTTCTCCTCGAAAATGAACGCACCCTTGACTTTCACGATTC AAACGTCAAGAATCTGTATGAGAAGGTGAAGAGCCAGCTGAAGAATAACGCGAAGGAAATCGGGAACGGTTGTTTCGAATTTTACC ACAAGTGCGATAATGAGTGCATGGAGAGCGTGCGAAATGGCACATATGACTATCCTAAGTACAGCGAGGAAAGTAAACTCAACAG AGAAAAGGTTGATGGCGTCAAGCTCGAGTCAATGGGGATTTACCAGATTCTGGCCATCTATAGTACTGTGGCTTCCTCTCTGGTTT TGCTCGTTTCACTGGGAGCCATTTCTTTCTGGATGTGTTCGAACGGATCCCTCCAATGCAGGATTTGTATT

2648	ATGAAAGCAAATCTGCTCGTCCTCCTGTGCGCTCTTGCAGCCGCCGACGCCGACACCATATGCATTGGATATCATGCTAACAACTC AACAGACACCGTGGACACTGTCCTAGAAAAAAACGTGACCGTGACACACTCGGTCAATCTGTTAGAAGACAGCCATAACGGGAAG CTGTGCCGACTCAAGGGAATAGCCCCCCTGCAGCTGGGCAAGTGCAATATCGCCGGGTGGCTCCTCGGCAACCCTGAGTGCGAT CCCTTGCTTCCAGTTAGAAGTTGGTCCTATATCGTCGAAACTCCTAACTCAGAGAATGGAATTTGCTATCCCGGAGATTTCATCGAC TACGAAGAGCTCAGAGAGCAACTTTCCAGTGTGTCAAGTTTCGAACGCTTTGAAATCTTCCCAAAGGAATCTTCATGGCCCAACCA CAATACGAATGGAGTGACCGCCGCCTGTAGTCACGAGGGAAAAAGTAGCTTCTACCGCAATCTTTTGTGGCTGACCGAGAAGGAA GGAAGCTACCCAAAGCTGAAGAACTCCTACGTGAACAAAAAAGGTAAGGAGGTCCTGGTGCTGTGGGGGATTCACCATCCCCCTA ACAGTAAGGAGCAGCAAAATCTCTACCAGAACGAGAACGCTTACGTATCTGTTGTGACTTCTAATTATAACAGGAGGTTCACGCCA GAAATTGCTGAGAGGCCAAAGGTCAGGGACCAGGCTGGGCGCATGAACTACTACTGGACCCTGCTAAAGCCTGGTGACACTATC ATCTTTGAGGCCAATGGAAATTTAATTGCACCCATGTACGCATTTGCCCTATCGCGCGGGTTCGGGAGCGGGATAATTACCTCCAA CGCATCGATGCATGAGTGCAACACAAAATGTCAAACACCTCTCGGGGCCATTAACAGCTCCCTCCCCTATCAGAATATTCACCCGG TAACAATCGGTGAATGTCCTAAGTACGTACGCTCCGCCAAGCTGAGGATGGTAACCGGTCTGCGGAATAATCCTAGCATCCAGTC CCGGGGTCTTTTCGGCGCCATCGCAGGTTTTATAGAAGGGGGCTGGACTGGTATGATCGATGGATGGTACGGGTACCATCACCAA AATGAGCAAGGTTCGGGGTACGCAGCAGACCAGAAATCAACCCAAAATGCCATTAACGGAATTACAAATAAGGTTAACACCGTTAT CGAAAAGATGAATATCCAGTTTACAGCCGTAGGGAAAGAATTTAATAAGCTCGAGAAGCGGATGGAAAACCTGAACAAGAAAGTG GATGACGGGTTCCTTGACATCTGGACTTATAATGCAGAGCTATTAGTGCTGCTGGAGAACGAACGTACCCTGGATTTCCATGATTC AAACGTCAAAAATCTTTATGAGAAAGTTAAGTCCCAGCTCAAGAATAATGCGAAGGAAATCGGCAACGGGTGTTTCGAATTCTATCA CAAGTGCGATAACGAATGCATGGAGAGCGTTAGAAACGGCACCTATGACTACCCCAAGTACAGCGAAGAGTCTAAGCTGAACCGG GAAAAGGTCGACGGTGTGAAACTGGAATCCATGGGCATTTACCAGATCCTCGCTATTTACTCGACAGTCGCAAGTAGCCTGGTGC TTTTGGTTTCTCTCGGCGCAATCTCATTCTGGATGTGCTCCAACGGATCCCTCCAGTGTCGCATCTGTATC
2649	ATGAAAGCTAACCTGCTGGTCCTTCTGTGCGCTCTCGCAGCAGCAGATGCGGACACCATCTGCATAGGATACCATGCTAATAATTC CACAGACACTGTGGACACAGTGCTGGAAAAGAATGTAACGGTGACCCACAGCGTTAACTTGCTGGAGGATAGCCACAATGGTAAG CTTTGTCGTTTGAAAGGGATCGCACCCCTGCAGTTGGGCAAATGTAATATCGCCGGGTGGCTCCTGGGCAACCCGGAGTGCGAC CCTTTGCTGCCCGTGCGGAGTTGGTCATACATTGTGGAAACTCCTAACAGTGAGAATGGTATTTGCTATCCTGGGGACTTCATTGA CTATGAAGAACTTCGCGAACAGTTATCTAGCGTGTCTAGCTTTGAACGCTTTGAGATCTTCCCAAAGGAGAGTAGTTGGCCTAACC ATAACACCAACGGGGTCACAGCGGCCTGTTCGCACGAAGGCAAGTCCAGCTTTTATCGCAATCTGCTGTGGTTGACAGAGAAGGA GGGATCATACCCGAAGCTGAAAAATTCCTACGTGAACAAGAAGGGCAAAGAAGTACTGGTGTTATGGGGCATTCACCACCCACCA AATAGCAAAGAGCAGCAGAACCTTTATCAGAACGAGAACGCCTACGTGAGCGTTGTTACGAGTAATTACAACCGCCGATTTACTCC CGAGATTGCGGAAAGGCCGAAGGTGAGAGACCAGGCAGGCCGCATGAACTATTACTGGACACTCCTGAAGCCTGGTGACACAAT AATCTTCGAAGCTAACGGCAACCTGATAGCCCCAATGTACGCCTTTGCCCTAAGTAGAGGTTTCGGCTCTGGAATTATTACGTCTA ACGCCAGCATGCACGAGTGTAACACTAAATGCCAAACCCCACTAGGTGCTATTAACTCGTCTCTGCCGTACCAGAATATCCACCCA GTAACGATTGGAGAGTGCCCTAAATATGTGCGATCAGCAAAGCTTAGAATGGTGACTGGGTTACGCAACAATCCCTCTATTCAGTC CCGCGGCCTGTTTGGGGCGATAGCCGGATTCATCGAAGGGGGCTGGACGGGCATGATTGACGGCTGGTATGGGTATCATCATCA GAACGAACAGGGATCAGGATACGCAGCTGACCAAAAATCCACGCAGAACGCCATCAACGGGATCACGAATAAAGTGAATACAGTG ATCGAGAAAATGAACATTCAGTTCACGGCAGTTGGAAAAGAGTTCAACAAACTTGAGAAGAGGATGGAGAACCTGAATAAAAAGGT GGACGACGGATTCCTGGATATATGGACTTACAATGCCGAACTACTCGTCCTACTGGAAAATGAGCGAACGCTGGACTTCCATGATT CGAATGTCAAGAATCTGTATGAAAAGGTCAAGTCCCAGCTGAAAAATAATGCTAAAGAGATAGGCAACGGGTGTTTCGAGTTTTAC CATAAGTGCGATAATGAATGTATGGAGTCCGTGCGAAACGGTACTTACGACTACCCAAAATACTCAGAGGAGTCCAAACTCAACAG GGAGAAAGTGGACGGGGTTAAGCTGGAAAGTATGGGTATTTATCAGATCCTGGCCATCTATTCTACAGTTGCATCTTCTCTAGTGC TGCTGGTATCTCTGGGCGCAATATCTTTTTGGATGTGTTCAAATGGGAGCCTGCAGTGCCGGATTTGTATC

2650	ATGAAAGCAAATCTTCTGGTGTTACTGTGTGCTTTGGCAGCAGCCGATGCTGACACCATATGTATTGGCTATCACGCAAACAATTC AACCGACACAGTCGACACCGTCCTCGAAAAGAACGTTACCGTGACCCACTCAGTGAACCTCCTCGAGGACTCCCATAATGGCAAG CTGTGTAGGTTAAAGGGGATCGCCCCACTGCAGTTGGGAAAGTGTAATATTGCTGGGTGGCTTCTGGGTAACCCCGAATGCGATC CACTCCTCCCCGTTCGCAGCTGGAGTTACATTGTTGAAACCCCAAACAGCGAGAACGGTATCTGTTACCCAGGAGATTTTATTGAT TACGAGGAGTTACGAGAACAACTGTCCTCAGTGAGTTCTTTTGAAAGGTTTGAGATATTCCCTAAAGAGAGCAGCTGGCCGAATCA TAACACCAATGGGGTGACTGCAGCTTGCTCCCACGAGGGGAAGTCCTCCTTCTATCGCAATCTGTTGTGGCTCACCGAAAAGGAG GGATCATATCCGAAGTTGAAAAATAGCTACGTTAATAAGAAGGGAAAAGAGGTGCTGGTGCTGTGGGGAATCCATCATCCACCTAA TTCCAAGGAGCAACAGAATCTATACCAAAACGAAAATGCTTACGTTTCTGTGGTGACCTCAAATTATAACCGCCGCTTCACTCCAGA GATCGCTGAGCGACCAAAAGTGCGGGACCAAGCCGGCAGAATGAATTACTATTGGACCTTGCTGAAACCCGGAGACACCATAATT TTTGAAGCCAATGGGAACCTGATTGCCCCTATGTACGCCTTCGCCCTGTCTAGGGGTTTCGGCAGCGGGATCATAACATCTAACG CAAGCATGCACGAATGTAATACCAAGTGTCAGACCCCTCTGGGAGCCATTAATTCCAGTCTCCCATACCAGAACATCCACCCCGTT ACCATTGGGGAGTGTCCGAAATACGTTAGAAGTGCCAAATTACGAATGGTGACAGGATTGCGGAACAACCCCAGCATCCAGAGCA GAGGCCTATTTGGCGCCATCGCAGGATTCATCGAAGGCGGATGGACCGGAATGATCGACGGATGGTACGGGTATCACCACCAGA ATGAACAGGGGTCGGGTTACGCCGCCGACCAGAAATCTACCCAGAATGCCATTAATGGGATAACAAACAAGGTCAACACCGTGAT TGAGAAGATGAATATACAGTTTACTGCCGTCGGCAAAGAGTTTAATAAACTGGAAAAGCGAATGGAAAACCTGAACAAGAAAGTCG ACGATGGGTTCCTGGATATCTGGACATACAACGCTGAGTTGCTCGTGCTGCTCGAGAATGAGAGAACCTTGGACTTTCACGATTCA AACGTAAAAAACTTATACGAAAAAGTCAAGAGTCAGCTGAAAAATAATGCCAAGGAAATCGGGAACGGATGCTTTGAGTTCTATCA CAAGTGCGACAATGAGTGTATGGAATCCGTTAGGAACGGGACATACGATTACCCAAAATACTCCGAGGAGAGCAAGTTAAACAGA GAAAAAGTTGACGGAGTGAAACTAGAGTCCATGGGTATTTACCAGATCCTGGCCATTTATTCTACCGTCGCGTCATCACTGGTACT CCTTGTCTCCCTTGGAGCGATATCCTTCTGGATGTGTTCCAACGGCTCCCTTCAGTGCCGAATCTGCATT
2651	ATGAAGGCTAATCTTCTGGTGTTACTGTGCGCCCTGGCTGCCGCCGACGCCGACACCATCTGCATTGGTTATCACGCTAATAACTC GACGGATACAGTGGATACTGTACTGGAAAAGAACGTGACAGTCACCCACAGTGTTAACCTGCTCGAGGATAGTCACAATGGCAAG TTGTGCAGGCTGAAGGGGATCGCTCCCCTGCAACTCGGCAAGTGTAATATCGCCGGCTGGCTTTTAGGCAACCCCGAGTGCGAC CCCCTACTGCCCGTTAGGTCCTGGAGCTACATCGTGGAAACTCCTAACAGCGAGAACGGCATCTGCTATCCTGGGGACTTCATCG ATTACGAAGAGCTCCGGGAACAGCTCTCTTCCGTGAGCTCCTTTGAGAGATTTGAGATCTTCCCTAAGGAATCGTCCTGGCCTAAC CACAATACAAATGGCGTGACAGCTGCCTGTTCACACGAAGGGAAAAGTTCCTTCTATAGGAATCTGCTGTGGCTGACAGAGAAAG AGGGCTCTTATCCTAAATTAAAGAACTCCTACGTGAACAAGAAGGGTAAAGAGGTCTTAGTGCTGTGGGGCATTCACCATCCCCCC AATTCCAAGGAGCAGCAAAATCTGTACCAGAACGAGAACGCTTATGTGTCCGTCGTTACTTCTAACTATAATAGGCGTTTCACCCC GGAGATTGCTGAGAGGCCTAAGGTACGGGACCAGCCGGTAGAATGAATTACTACTGGACACTACTTAAGCCAGGTGACACAATC ATCTTTGAGGCAAACGGAAACCTAATCGCTCCAATGTATGCATTTGCCCTGAGCCGCGGATTCGGTAGCGGAATTATCACCAGCAA TGCCTCTATGCACGAGTGCAATACTAAGTGTCAGACTCCTTTGGGAGCGATCAACTCCAGCTTGCCATACCAAAACATTCACCCGG TGACTATTGGTGAATGCCCCAAGTACGTGCGGTCTGCAAAACTGAGAATGGTTACCGGACTCCGGAATAACCCTAGTATACAGTCT AGGGGGCTCTTTGGAGCCATTGCAGGTTTCATCGAAGGCGGCTGGACCGGCATGATCGACGGATGGTATGGTTATCACCACCAA AACGAGCAGGGATCAGGATATGCTGCAGATCAAAAATCTACTCAAAATGCAATAAACGGTATCACCAATAAGGTGAACACCGTAAT CGAAAAAATGAACATCCAGTTTACAGCCGTGGGCAAGGAGTTTAATAAGCTGGAGAAACGCATGGAGAACCTCAATAAGAAGGTT GACGATGGATTCCTGGACATCTGGACGTATAATGCCGAGCTCCTGGTACTGCTCGAGAATGAACGAACCCTCGACTTCCATGATA GCAACGTGAAGAACTTATATGAGAAGGTGAAGTCACAACTGAAAAACAATGCTAAGGAAATAGGCAACGGATGTTTTGAATTCTAC CACAAGTGTGACAACGAATGTATGGAATCGGTGAGGAATGGGACATATGACTACCCCAAGTACTCGGAAGAAAGTAAGCTTAATAG GGAGAAGGTGGACGGCGTGAAACTCGAGAGCATGGGGATTTACCAAATCCTAGCAATCTATTCCACAGTTGCTTCTTCTCTGGTG CTCCTTGTGTCACTTGGGGCGATTTCGTTCTGGATGTGCTCAAACGGGAGCCTCCAGTGCCGGATCTGCATC

2652	ATGAAGGCAAACCTGCTGGTCCTGCTCTGCGCACTGGCCGCGGCAGACGCCGACACAATCTGCATAGGCTATCACGCAAATAACA GTACTGACACCGTCGATACAGTCCTGGAGAAAAACGTGACTGTGACCCACAGCGTAAATCTGCTGGAAGACAGCCACAATGGAAA ATTGTGCAGATTGAAAGGAATAGCACCCCTGCAATTGGGGAAATGTAATATTGCCGGATGGCTGCTCGGCAACCCCGAATGCGAC CCGCTGCTGCCCGTCCGGTCTTGGAGTTATATCGTTGAGACACCGAACTCCGAGAATGGTATCTGCTATCCGGGCGACTTCATCG ACTATGAGGAACTCAGAGAGCAGCTGTCATCTGTGTCTTCATTCGAGCGTTTCGAGATTTTCCCAAAGGAGTCATCGTGGCCCAAC CATAACACAAACGGAGTCACGGCAGCATGCTCTCACGAAGGGAAAAGCTCTTTCTACCGCAACCTGCTCTGGTTGACTGAGAAAG AGGGGTCTTACCCCAAACTTAAGAACAGCTACGTGAACAAGAAGGGCAAGGAGGTCTTGGTACTGTGGGGCATCCATCACCCACC TAATAGTAAAGAGCAACAAAACCTATACCAGAACGAGAACGCATACGTGTCAGTAGTCACCTCTAATTACAACAGGCGGTTTACAC CAGAGATAGCTGAGCGGCCAAAGGTCCGGGACCAGGCAGGCAGGATGAACTACTATTGGACTTTGTTAAAGCCTGGAGACACCA TCATCTTTGAGGCTAACGGGAATCTGATTGCTCCCATGTATGCTTTTGCACTTTCCAGAGGCTTTGGATCCGGTATTATCACATCGA ATGCGTCCATGCATGAGTGTAACACTAAGTGTCAAACACCACTCGGCGCAATCAATTCTTCACTTCCTTATCAAAATATTCACCCCG TGACCATCGGCGAGTGCCCCAAGTATGTCAGGAGCGCTAAGCTGCGGATGGTGACAGGGTTGAGGAATAATCCTTCAATTCAGTC GAGGGGATTATTCGGTGCCATCGCGGGATTCATTGAGGGGGGGTGGACAGGAATGATAGACGGGTGGTATGGTTACCACCATCA GAACGAGCAGGGGTCAGGCTACGCCGCTGACCAAAAATCCACCCAAAATGCCATCAATGGAATCACAAATAAGGTAAATACAGTG ATCGAAAAAATGAACATTCAATTCACAGCTGTCGGAAAGGAGTTCAATAAGCTTGAGAAACGAATGGAGAACTTAAATAAAAAAGTT GACGACGGTTTCCTGGACATTTGGACATACAACGCAGAACTGCTCGTTTTACTAGAGAACGAACGGACGCTGGACTTTCATGATTC TAACGTAAAGAACTTGTACGAGAAGGTGAAAAGCCAGCTTAAAAATAACGCCAAAGAAATCGGAAACGGCTGCTTCGAATTCTACC ACAAATGTGATAACGAGTGCATGGAGAGCGTGAGGAATGGGACATATGACTATCCCAAATATTCTGAAGAGTCAAAACTGAATCGG GAAAAGGTCGATGGTGTCAAACTGGAGTCTATGGGCATCTATCAGATACTGGCAATCTACTCCACCGTGGCCAGCAGTCTCGTGC TGCTGGTGAGCCTGGGCGCAATAAGCTTCTGGATGTGCTCAAACGGAAGCCTGCAATGTAGGATTTGTATC
2653	ATGAAGGCCAATCTCCTGGTGCTGTTGTGCGCACTGGCAGCGGCGGATGCAGATACAATATGTATCGGATATCACGCAAATAACA GCACAGATACAGTGGACACCGTGCTGGAGAAGAATGTGACTGTGACGCATAGTGTTAACCTCTTGGAAGATAGCCACAATGGCAA GCTGTGTCGCTTGAAGGGAATCGCCCCCCTGCAGCTCGGGAAATGCAATATCGCGGGATGGCTTCTGGGTAACCCTGAATGCGA CCCGCTGCTGCCCGTGCGCTCATGGTCATACATTGTGGAGACACCTAATTCAGAGAATGGGATTTGCTATCCTGGGGACTTTATC GATTACGAGGAACTGAGAGAGCAGCTATCTTCAGTTAGTTCCTTCGAACGATTCGAGATCTTTCCAAAGGAGAGTAGCTGGCCCAA TCATAACACTAACGGAGTGACCGCCGCGTGCAGCCACGAGGGCAAATCCTCATTTTACCGAAACCTGTTATGGCTCACAGAAAAG GAGGGGTCTTACCCGAAGCTGAAAAACTCTTATGTGAACAAGAAGGGGAAAGAGGTGTTAGTTCTGTGGGGCATTCACCACCCTC CAAATTCCAAGGAACAACAAAACCTTTACCAAAACGAGAATGCATATGTCTCGGTCGTGACGTCGAACTATAATAGAAGATTTACCC CCGAGATTGCTGAACGACCCAAGGTGAGAGATCAGGCGGGAAGAATGAACTATTACTGGACCTTGCTCAAACCGGGAGATACAAT CATATTCGAGGCTAACGGAAACTTGATCGCCCCAATGTACGCCTTTGCCCTCTCCCGCGGTTTCGGGAGCGGTATCATCACCTCA AATGCTTCTATGCATGAATGCAATACCAAATGTCAAACACCTCTGGGCGCAATCAACTCGAGCCTGCCTTATCAGAACATTCACCCT GTCACCATTGGTGAGTGCCCCAAGTACGTAAGATCAGCTAAGCTGCGTATGGTGACCGGGCTGAGAAACAATCCCTCCATCCAAA GTCGCGGTTTATTCGGCGCCATTGCCGGGTTCATAGAGGGTGGGTGGACAGGAATGATCGATGGCTGGTATGGATATCATCATCA GAACGAACAAGGATCAGGGTATGCAGCAGACCAGAAGTCTACTCAAAACGCAATTAATGGCATCACCAATAAGGTTAACACTGTCA TAGAAAAGATGAACATCCAGTTCACTGCGGTGGGCAAAGAGTTTAACAAGCTCGAGAAGCGGATGGAGAACCTCAATAAGAAAGT CGATGACGGATTCCTGGATATATGGACATATAATGCCGAGCTGCTGGTGCTGCTGGAAAACGAAAGAACCCTCGATTTCCACGAC TCTAATGTGAAGAATCTGTACGAGAAGGTCAAGTCCCAGCTGAAAAATAACGCCAAGGAGATAGGGAACGGCTGCTTTGAATTCTA TCACAAGTGCGACAATGAGTGCATGGAGAGCGTTCGGAACGGGACTTACGATTATCCGAAGTATTCAGAGGAGAGCAAGCTTAAT CGTGAAAAGGTCGACGGAGTGAAATTAGAAAGTATGGGAATCTATCAGATTCTTGCAATCTATAGCACCGTTGCCAGCTCTCTGGT GTTACTGGTAAGCCTCGGGGCAATATCTTTTTGGATGTGTAGCAATGGGTCTCTCCAGTGTCGAATTTGTATT

2654	ATGAAGGCCAACCTGCTCGTGCTTCTGTGTGCGCTGGCCGCCGCCGACGCAGACACTATCTGTATCGGCTATCACGCCAATAACA GCACGGATACCGTGGATACCGTATTGGAGAAAAATGTCACGGTAACACACTCTGTGAATCTGCTTGAAGACTCCCATAACGGAAAA CTTTGTCGGCTCAAGGGCATTGCCCCTCTGCAACTGGGCAAGTGCAACATCGCCGGGTGGCTGCTGGGCAATCCGGAGTGCGAC CCTCTCCTGCCAGTCAGGTCCTGGTCCTATATCGTAGAAACCCCCAATAGTGAAAACGGCATATGCTACCCAGGTGATTTCATCGA TTATGAAGAGCTCAGAGAGCAACTGAGCTCCGTCAGCAGTTTTGAGCGGTTCGAGATTTTCCCAAAGGAATCATCATGGCCGAATC ACAACACCAATGGAGTGACCGCAGCGTGTTCTCATGAGGGGAAGAGCTCGTTCTATCGGAACCTGCTTTGGTTGACGGAGAAGGA GGGGTCATACCCCAAACTTAAGAACTCATACGTGAATAAAAAGGGCAAGGAGGTCCTCGTGCTGTGGGGGATTCACCACCCCCCC AATTCCAAGGAGCAGCAGAATCTGTACCAAAACGAGAATGCTTATGTGAGTGTGGTGACGTCTAATTATAATAGACGGTTTACACC TGAGATCGCAGAAAGACCCAAAGTGCGCGACCAGGCTGGGAGGATGAATTACTATTGGACTCTGTTAAAGCCCGGCGATACCATC ATTTTCGAGGCAAATGGGAATCTTATCGCTCCAATGTACGCCTTCGCCCTGTCACGGGGGTTCGGGAGCGGGATTATCACATCAA ATGCCAGCATGCATGAATGCAACACTAAGTGCCAGACCCCACTGGGCGCCATCAATTCCAGCCTCCCTTACCAGAACATACACCC TGTTACAATAGGAGAGTGCCCCAAGTACGTCCGATCCGCTAAACTGCGGATGGTTACAGGCCTACGTAATAATCCTAGCATTCAGT CACGGGGACTGTTTGGCGCAATAGCCGGGTTCATCGAGGGCGGGTGGACCGGAATGATTGATGGGTGGTACGGATACCACCATC AGAATGAGCAGGGATCTGGGTACGCCGCCGACCAGAAGAGCACACAGAATGCCATTAATGGCATAACCAACAAAGTGAACACCGT GATCGAGAAGATGAACATTCAATTTACGGCTGTAGGTAAGGAATTCAACAAACTGGAGAAACGGATGGAGAATTTGAACAAGAAGG TCGATGATGGGTTCCTCGATATTTGGACCTACAACGCTGAGCTTTTGGTGCTGCTGGAGAACGAGAGGACACTTGATTTTCATGAT TCGAATGTGAAAAACTTGTACGAAAAGGTCAAGTCACAGCTGAAGAACAACGCTAAAGAGATTGGAAATGGATGCTTCGAGTTTTA TCACAAATGCGACAATGAATGTATGGAGAGTGTGCGGAACGGAACATACGATTACCCCAAATATTCCGAGGAGTCTAAGCTTAACC GTGAGAAGGTCGATGGCGTTAAGTTAGAGTCCATGGGAATCTACCAGATCCTGGCTATATATAGCACCGTCGCAAGCTCTCTAGTC TTGCTCGTCTCACTGGGAGCAATCTCGTTTTGGATGTGCTCTAACGGGTCCCTACAGTGTCGGATTTGCATC
2655	ATGAAAGCTAATCTGCTGGTCCTGTTGTGTGCACTGGCGGCTGCAGATGCGGATACCATTTGTATCGGTTATCACGCTAATAACTC TACTGACACAGTCGACACGGTGTTGGAAAAGAACGTCACAGTCACCCACAGCGTCAATCTGCTCGAAGATTCACACAACGGAAAG CTATGTAGGTTAAAGGGTATTGCACCATTGCAACTGGGAAAGTGTAACATTGCTGGATGGCTGCTGGGGAACCCCGAGTGTGATC CTTTGCTCCCTGTCAGATCCTGGTCCTACATTGTGGAAACCCCTAACTCAGAGAACGGGATCTGCTATCCAGGAGATTTTATCGAC TATGAAGAGCTGAGGGAACAACTATCCAGCGTCTCCAGCTTTGAGAGGTTCGAGATATTCCCTAAGGAGTCAAGCTGGCCAAATC ACAATACAAACGGTGTCACAGCCGCCTGCAGTCACGAAGGCAAGTCATCGTTCTACCGAAATTTGTTGTGGTTGACGGAAAAAGA GGGATCATACCCAAAACTCAAGAATAGCTATGTCAACAAGAAAGGCAAGGAAGTGCTTGTCCTCTGGGGCATACATCATCCTCCTA ACTCTAAGGAGCAGCAAAACTTATACCAGAACGAGAACGCCTATGTGAGTGTGGTCACCAGCAATTATAATCGCAGGTTCACGCCT GAGATTGCCGAGAGGCCAAAAGTCAGAGACCAGGCCGGCAGGATGAATTACTATTGGACATTACTGAAACCAGGCGATACTATTA TCTTCGAGGCTAACGGAAATCTGATAGCACCAATGTACGCCTTTGCCTTGTCCCGTGGGTTCGGCTCTGGGATCATAACATCCAAC GCAAGCATGCATGAGTGCAACACAAAGTGCCAGACACCGCTTGGAGCCATTAATTCTTCTCTGCCATATCAGAACATTCATCCTGT AACAATCGGGGAATGCCCAAAGTACGTTAGAAGTGCCAAGCTGCGCATGGTGACCGGCCTGAGGAATAATCCCTCTATTCAGTCC AGAGGACTGTTCGGAGCCATCGCTGGGTTTATTGAGGGCGGCTGGACCGGAATGATAGATGGCTGGTACGGGTACCACCACCAG AATGAACAGGGCTCTGGCTATGCCGCGGACCAGAAATCTACCCAAAATGCCATCAATGGAATCACCAATAAAGTTAACACGGTGAT TGAGAAAATGAACATTCAATTTACCGCGGTAGGAAAGGAGTTTAATAAGCTTGAGAAGCGTATGGAGAATTTGAACAAGAAGGTGG ATGACGGTTTCCTCGATATTTGGACCTATAATGCCGAACTACTAGTTTTATTGGAGAACGAACGCACACTGGATTTCCACGACTCTA ATGTTAAAAATCTGTACGAGAAAGTGAAGTCACAGCTAAAGAATAATGCCAAGGAAATAGGAAATGGATGCTTTGAATTTTACCACA AATGTGACAACGAATGCATGGAATCTGTGCGGAATGGAACTTACGATTACCCTAAGTACTCTGAGGAGAGTAAACTGAACAGAGAG AAGGTAGACGGCGTAAAGCTCGAATCAATGGGAATCTACCAAATTCTGGCCATCTACTCTACCGTCGCCTCCTCTCTGGTTTTATT GGTCTCTCTTGGAGCCATTAGTTTCTGGATGTGCAGTAACGGCTCCCTGCAGTGCAGGATTTGTATC

2656	ATGAAGGCCAACCTCCTCGTGCTGCTCTGCGCCTTGGCAGCCGCAGACGCCGATACCATCTGTATAGGGTACCACGCCAACAATT CAACAGATACCGTCGATACCGTACTGGAGAAAAACGTCACCGTCACGCACAGCGTTAACCTGTTGGAAGATAGTCATAATGGAAA GCTCTGTCGCCTAAAAGGAATCGCGCCTCTGCAGCTGGGAAAGTGCAACATAGCCGGGTGGCTCCTCGGGAACCCGGAGTGTGA CCCTCTTTTGCCGGTGCGCTCATGGTCATATATTGTTGAGACCCCTAACAGCGAAAACGGAATATGTTACCCAGGGGACTTCATCG ATTATGAGGAATTACGAGAACAACTAAGTTCCGTCTCTTCCTTCGAGCGCTTTGAAATCTTTCCTAAAGAGAGTTCCTGGCCCAACC ACAACACTAATGGCGTGACAGCTGCTTGCAGTCATGAGGGCAAAAGTTCATTCTATAGAAATCTTCTCTGGCTAACTGAGAAAGAA GGATCCTACCCTAAACTCAAAAATTCCTATGTTAACAAGAAGGGCAAAGAGGTGCTCGTGCTGTGGGGGATCCACCACCCACCAA ATTCAAAAGAGCAACAAAACCTCTACCAGAATGAAAATGCCTACGTGAGCGTTGTGACTTCAAATTACAATAGACGGTTTACCCCAG AGATCGCAGAACGCCCTAAGGTACGTGATCAGGCTGGGCGAATGAACTACTACTGGACACTCCTCAAACCAGGCGACACTATAAT CTTCGAGGCTAACGGCAACCTGATTGCTCCAATGTACGCGTTCGCTCTTTCACGCGGATTCGGCTCCGGCATCATTACTAGTAATG CATCTATGCATGAATGCAATACTAAGTGCCAGACCCCACTCGGAGCCATAAACAGTTCTCTGCCATATCAGAACATACACCCTGTG ACTATAGGAGAATGCCCAAAATATGTTCGGTCCGCTAAGCTGCGCATGGTGACCGGACTCCGGAACAACCCATCCATACAATCAC GGGGGCTGTTTGGCGCCATTGCCGGCTTCATCGAGGGTGGATGGACTGGTATGATCGATGGTTGGTATGGATACCATCATCAGAA TGAACAGGGCTCCGGCTATGCCGCCGATCAGAAGTCCACCCAGAACGCCATTAACGGAATTACCAACAAGGTTAATACTGTGATC GAGAAGATGAATATTCAGTTCACAGCAGTCGGGAAGGAATTCAACAAACTGGAGAAACGTATGGAAAATCTCAACAAGAAGGTGG ACGATGGGTTTTTAGACATTTGGACTTATAATGCCGAACTGCTGGTGCTGCTGGAAAACGAGAGAACGTTGGACTTTCACGACAGC AACGTGAAAAACCTGTATGAAAAAGTGAAGTCCCAGCTCAAAAACAATGCAAAGGAAATCGGCAACGGTTGCTTTGAGTTCTACCA TAAGTGCGACAACGAATGTATGGAATCTGTGCGCAATGGAACCTACGACTATCCTAAGTATAGCGAAGAATCTAAATTAAATAGGG AAAAGGTTGATGGAGTGAAACTGGAGTCTATGGGGATCTACCAGATCCTTGCTATCTACTCAACCGTCGCTAGCTCCCTCGTACTG CTGGTCAGTTTAGGAGCGATATCTTTCTGGATGTGTTCTAATGGGTCACTGCAGTGTAGAATTTGCATC
2657	ATGAAGGCCAATCTGCTGGTGCTTCTCTGCGCCTTAGCCGCCGCCGATGCAGATACCATTTGTATTGGATATCACGCTAACAACAG TACAGATACTGTGGACACTGTCCTGGAAAAGAACGTCACCGTCACGCATAGCGTAAACCTGCTCGAGGACTCACACAACGGGAAG CTCTGTCGTCTGAAGGGCATCGCCCCCCTGCAGCTCGGCAAATGCAACATTGCGGGATGGCTTCTTGGAAACCCCGAGTGTGATC CCTTATTACCAGTGCGCTCTTGGTCCTATATAGTAGAAACCCCAAACAGCGAGAACGGGATTTGTTACCCCGGCGATTTCATAGAC TACGAGGAGTTGCGCGAACAGCTGTCTAGCGTCTCCTCATTCGAACGTTTCGAGATTTTCCCAAAGGAATCCTCGTGGCCCAATCA TAATACCAATGGAGTCACTGCCGCCTGCAGCCACGAGGGAAAGTCATCTTTCTATCGGAATCTGCTCTGGCTGACCGAAAAGGAG GGATCTTATCCTAAACTGAAAAATTCTTACGTCAATAAGAAAGGCAAGGAAGTCTTAGTATTATGGGGAATTCATCACCCTCCCAAC AGTAAGGAGCAGCAGAACCTGTACCAGAACGAGAACGCCTATGTTTCTGTCGTGACATCTAACTATAACAGACGTTTCACCCCAGA AATTGCTGAAAGGCCAAAGGTCCGTGATCAGGCCGGCAGGATGAATTACTACTGGACTCTTCTGAAGCCTGGTGACACAATCATTT TCGAAGCCAACGGCAACCTGATTGCTCCCATGTACGCGTTCGCTCTGAGTCGCGGCTTTGGATCGGGTATTATTACGTCCAATGC CTCTATGCATGAATGCAATACTAAATGTCAGACCCCTCTAGGCGCCATCAATTCTAGCCTCCCGTACCAGAACATTCATCCTGTGA CTATTGGTGAATGCCCAAAGTACGTGAGGTCCGCCAAGCTTCGCATGGTGACCGGCCTGCGTAACAACCCCTCCATCCAATCCAG AGGTCTCTTTGGGGCAATCGCCGGTTTTATTGAGGGTGGTTGGACCGGAATGATCGATGGTTGGTACGGGTACCACCATCAGAAT GAGCAGGGCAGTGGATATGCCGCTGACCAAAAATCTACACAAAACGCGATCAATGGAATCACCAATAAAGTGAACACAGTCATAG AAAAAATGAACATCCAGTTTACTGCGGTCGGGAAGGAATTTAATAAGCTCGAGAAAAGAATGGAAAATCTAAACAAAAAGGTCGAT GACGGTTTCCTCGATATCTGGACATACAATGCCGAATTACTGGTCCTGCTCGAGAATGAGCGGACATTAGACTTCCACGACAGCAA TGTAAAGAACCTTTACGAGAAGGTGAAGTCCCAGCTGAAGAACAATGCTAAAGAGATTGGGAATGGCTGTTTTGAATTTTACCATAA ATGCGACAACGAGTGTATGGAAAGCGTGCGGAACGGGACTTACGATTACCCAAAATATAGTGAGGAGAGCAAATTAAACCGTGAG AAGGTCGACGGGGTGAAGTTGGAATCCATGGGCATTTACCAAATCCTCGCAATTTACTCCACCGTGGCTTCTTCATTGGTTCTCCT GGTGTCCCTTGGGGCCATATCATTTTGGATGTGCTCTAATGGGAGTCTCCAGTGCCGTATATGCATT

2658	ATGAAGGCTAACCTACTGGTTCTGCTGTGTGCGCTTGCCGCCGCAGATGCCGACACCATCTGCATCGGGTATCACGCAAATAACT CCACCGACACAGTAGACACAGTATTGGAAAAAAACGTGACCGTCACCCACTCTGTGAACCTGCTCGAAGATAGCCATAATGGCAA ATTGTGTAGACTCAAAGGTATCGCTCCCTTGCAGCTAGGAAAGTGCAACATTGCAGGATGGCTCCTCGGCAACCCTGAGTGTGAC CCCCTCCTGCCCGTCCGATCCTGGTCCTACATCGTGGAGACTCCAAACTCGGAGAACGGAATATGTTATCCTGGGGATTTCATTG ACTATGAGGAGCTTAGGGAGCAACTATCCAGCGTATCTAGTTTCGAACGGTTCGAGATCTTTCCAAAAGAGTCCAGTTGGCCTAAC CATAACACGAACGGTGTGACCGCTGCTTGTAGCCATGAGGGGAAATCATCTTTCTATCGCAACTTGCTTTGGCTCACAGAGAAGGA GGGCTCCTACCCTAAGTTGAAAAACAGCTATGTCAATAAAAAAGGCAAAGAGGTTCTCGTGTTATGGGGTATTCACCACCCTCCTA ACTCTAAAGAGCAACAAAATCTGTATCAGAACGAGAATGCGTATGTGTCAGTCGTTACATCAAATTACAATAGGCGGTTTACACCAG AGATCGCTGAAAGACCTAAAGTTCGGGACCAGGCCGGCCGCATGAATTACTACTGGACTCTGCTGAAGCCAGGGGATACGATTAT CTTTGAAGCTAATGGTAACCTGATTGCTCCCATGTACGCCTTTGCCTTATCAAGAGGATTTGGATCCGGCATAATCACATCCAATGC TAGCATGCACGAGTGCAACACCAAATGCCAGACTCCTTTGGGAGCCATCAACTCTAGTCTGCCATATCAGAATATCCATCCTGTGA CCATCGGGGAGTGCCCGAAGTATGTGCGATCAGCCAAGCTGCGGATGGTCACCGGATTAAGAAATAACCCCTCCATTCAGTCTCG AGGACTGTTCGGTGCTATCGCTGGTTTTATTGAAGGTGGTTGGACAGGCATGATCGATGGCTGGTACGGTTATCACCACCAGAAC GAACAGGGGTCTGGGTACGCTGCCGATCAGAAGAGTACCCAGAACGCGATCAATGGCATTACCAACAAAGTGAACACAGTAATAG AAAAGATGAACATTCAGTTCACCGCTGTTGGTAAAGAGTTCAACAAGCTGGAAAAACGGATGGAGAATCTCAACAAAAAGGTTGAT GATGGGTTCCTGGATATCTGGACCTACAACGCGGAACTGCTGGTACTGCTGGAAAATGAACGGACACTTGACTTTCACGACTCTAA CGTGAAGAATCTGTACGAAAAGGTAAAATCACAGCTAAAGAACAATGCAAAGGAGATTGGGAATGGATGCTTCGAGTTCTACCATA AGTGCGACAACGAGTGTATGGAGTCTGTGCGGAATGGCACTTACGATTACCCTAAGTACTCCGAGGAATCTAAACTGAATCGAGA GAAAGTGGATGGTGTAAAGCTCGAGAGTATGGGCATTTATCAGATTCTGGCTATCTACTCGACTGTAGCCTCCTCACTGGTGTTAC TGGTGAGCCTCGGTGCAATCTCATTCTGGATGTGTAGTAACGGCTCCTTACAGTGTCGCATTTGTATC
2659	ATGAAGGCCAACTTGTTAGTCCTGCTGTGCGCTTTAGCGGCTGCCGACGCCGATACAATCTGTATTGGCTATCACGCTAACAATAG CACTGACACAGTCGATACAGTTCTGGAGAAGAATGTTACAGTCACGCATAGCGTGAACTTACTGGAAGACTCTCACAATGGCAAGT TGTGTCGGCTGAAGGGGATCGCTCCATTGCAACTCGGAAAATGCAATATAGCGGGCTGGCTGCTCGGAAATCCAGAATGTGATCC CCTTCTGCCTGTCCGCTCCTGGTCCTACATCGTGGAGACTCCTAATTCTGAGAACGGAATATGCTACCCTGGCGACTTCATCGACT ACGAAGAACTCCGGGAACAGCTCTCTAGCGTGAGTTCCTTTGAGAGATTCGAAATTTTTCCAAAAGAGTCCAGCTGGCCTAATCAT AATACAAACGGCGTGACTGCTGCATGCAGCCACGAAGGGAAATCGTCTTTTTATAGAAATCTGCTGTGGCTGACGGAGAAAGAAG GATCATACCCAAAGCTAAAGAACTCATATGTTAACAAGAAAGGGAAGGAGGTGCTGGTGCTGTGGGGGATACATCACCCACCCAA TTCTAAAGAGCAGCAGAACTTGTATCAGAACGAAAATGCCTACGTGAGCGTCGTGACTTCCAACTATAACCGAAGATTCACTCCCG AAATTGCCGAACGCCCCAAGGTCCGGGATCAGGCGGGACGCATGAATTACTATTGGACCCTGCTGAAACCTGGAGACACTATTAT CTTCGAGGCCAACGGCAACTTGATCGCTCCCATGTATGCCTTTGCCCTGAGTCGAGGTTTTGGAAGCGGGATTATCACTTCCAATG CATCAATGCATGAATGCAATACAAAGTGTCAGACACCCCTTGGCGCAATTAACTCATCTCTGCCCTACCAGAACATTCATCCAGTCA CTATCGGGGAGTGCCCTAAATATGTTAGATCTGCCAAGCTGCGGATGGTGACAGGGCTTAGAAATAACCCCAGCATTCAGAGCCG CGGCTTGTTTGGAGCCATTGCTGGGTTTATTGAGGGAGGGTGGACCGGAATGATAGACGGTTGGTACGGGTACCATCATCAGAAT GAGCAGGGCAGTGGGTACGCCGCTGACCAAAAGAGCACCCAGAACGCTATCAACGGAATTACCAATAAGGTCAATACAGTGATAG AAAAGATGAACATTCAGTTTACTGCTGTGGGGAAGGAGTTCAATAAACTGGAAAAACGGATGGAGAACCTCAACAAGAAAGTGGAC GACGGATTTCTGGACATTTGGACCTATAACGCCGAGTTACTGGTACTGCTTGAAAACGAAAGGACCTTAGACTTTCATGATTCTAAT GTTAAAAATCTGTATGAAAAGGTCAAGAGCCAACTTAAGAACAACGCCAAAGAAATCGGAAACGGTTGCTTCGAGTTTTATCATAAG TGTGACAATGAGTGCATGGAGTCTGTTCGAAATGGTACATACGACTATCCCAAGTATAGTGAAGAGAGCAAACTTAATCGGGAGAA GGTGGATGGCGTGAAACTGGAAAGCATGGGAATCTACCAGATCTTAGCCATATACAGTACAGTCGCTTCATCTCTCGTGCTTCTCG TCAGCCTCGGAGCAATCAGCTTTTGGATGTGTAGCAACGGCAGCCTCCAATGCCGAATCTGCATC

2660	ATGAAGGCAAACCTCCTTGTCCTCCTGTGTGCACTGGCAGCCGCCGACGCCGACACTATTTGTATCGGGTACCATGCAAATAACT CCACCGATACCGTGGACACAGTCCTGGAAAAAAACGTGACAGTGACGCATTCAGTGAATCTCCTTGAAGATAGCCACAATGGAAA GTTGTGCCGGCTGAAAGGAATCGCACCTTTACAGCTGGGAAAGTGTAACATTGCAGGGTGGCTGTTGGGGAACCCTGAGTGTGA CCCCCTCTTACCCGTCCGGTCTTGGAGTTACATAGTGGAAACACCAAATAGTGAGAACGGCATCTGCTACCCAGGAGATTTCATCG ATTATGAAGAGCTGCGTGAACAACTCTCTAGTGTTAGTTCTTTCGAAAGGTTCGAGATCTTTCCTAAAGAATCGTCATGGCCCAATC ACAATACAAACGGGGTCACAGCTGCTTGCTCACACGAAGGAAAGAGCAGTTTCTATAGAAATCTGCTGTGGCTGACCGAGAAGGA GGGTTCTTACCCTAAACTTAAAAATTCTTATGTTAACAAGAAGGGCAAAGAAGTGCTTGTCCTGTGGGGCATCCACCATCCCCCTA ACTCTAAAGAGCAGCAGAACTTGTACCAGAATGAGAACGCCTATGTGTCAGTGGTTACCTCAAATTATAACCGGAGATTTACGCCT GAGATCGCCGAGCGGCCCAAAGTGCGGGATCAAGCCGGACGTATGAACTACTATTGGACACTCCTTAAGCCAGGCGATACCATTA TCTTTGAGGCAAATGGCAACCTGATCGCACCCATGTACGCGTTCGCCTTGTCTCGAGGATTCGGCAGCGGCATCATCACGTCCAA CGCCAGTATGCATGAGTGTAATACAAAATGCCAAACTCCCCTCGGGGCAATTAACTCGTCTCTGCCATACCAGAACATACATCCAG TGACTATTGGTGAATGTCCCAAGTACGTCCGGTCTGCCAAGCTGCGGATGGTGACTGGCTTGCGCAATAACCCTAGCATTCAGTC CAGAGGGCTGTTCGGCGCAATCGCCGGGTTCATCGAGGGGGGCTGGACTGGCATGATCGATGGCTGGTATGGATATCACCATCA GAATGAACAGGGGAGCGGTTATGCGGCCGATCAGAAATCGACCCAGAACGCAATCAATGGAATTACCAACAAGGTTAACACAGTC ATAGAGAAAATGAACATCCAGTTCACGGCCGTTGGGAAGGAGTTTAATAAGTTGGAAAAGCGAATGGAAAACCTGAATAAAAAAGT CGACGACGGCTTCCTGGATATCTGGACCTATAACGCCGAGCTACTCGTTCTGTTGGAGAACGAGAGGACACTCGACTTCCATGAC AGCAATGTGAAAAACCTGTACGAAAAGGTGAAAAGTCAGCTCAAGAACAATGCAAAAGAGATCGGCAACGGATGTTTCGAGTTTTA CCACAAGTGCGATAATGAGTGTATGGAAAGCGTGCGCAATGGCACTTATGACTACCCCAAGTATTCTGAGGAGTCCAAATTGAATC GAGAAAAGGTGGATGGCGTCAAGCTAGAATCGATGGGGATATACCAGATATTGGCCATCTACTCCACCGTGGCCTCCTCCCTCGT CCTTCTTGTTTCGCTGGGCGCCATCTCTTTTTGGATGTGCTCAAACGGCTCGCTTCAGTGTCGCATTTGCATC
2661	ATGAAGGCAAACCTGTTAGTTCTCCTCTGCGCTTTGGCTGCGGCCGACGCCGACACAATCTGTATCGGTTATCATGCGAACAACTC AACTGACACAGTGGATACTGTGCTGGAAAAAAATGTGACAGTCACCCACTCTGTGAATCTGCTAGAGGACTCACACAACGGAAAG CTGTGCAGGCTCAAGGGCATTGCGCCACTGCAATTAGGAAAGTGTAACATCGCTGGATGGCTACTGGGAAACCCCGAGTGTGAC CCCCTGCTTCCTGTGCGTAGTTGGAGTTATATCGTCGAAACTCCAAACAGCGAAAACGGCATTTGTTATCCCGGCGATTTCATCGA TTATGAGGAATTGCGGGAGCAACTATCTAGCGTGTCTAGTTTTGAGAGATTCGAGATCTTCCCTAAAGAGTCTTCCTGGCCCAATC ATAACACAAATGGTGTCACCGCAGCCTGCAGTCACGAGGGGAAGAGTAGCTTTTATCGCAACCTGCTGTGGCTGACCGAAAAGGA AGGCTCCTATCCTAAACTTAAGAACAGCTACGTGAACAAAAAGGGTAAGGAGGTCCTTGTTCTCTGGGGAATCCACCACCCCCCTA ATAGCAAAGAGCAGCAAAACCTTTATCAGAATGAGAACGCTTATGTTTCTGTGGTGACCTCAAACTATAATAGACGTTTCACACCTG AAATCGCCGAAAGGCCCAAGGTTCGAGATCAAGCCGGTCGTATGAATTACTATTGGACGCTGCTAAAGCCAGGCGACACCATAAT TTTCGAAGCGAACGGTAACTTGATCGCCCCTATGTATGCATTCGCCCTGTCACGTGGCTTTGGCTCTGGTATTATCACCAGCAATG CCTCTATGCACGAGTGTAATACTAAATGTCAGACCCCACTCGGCGCTATCAACTCATCCTTGCCCTACCAGAACATACATCCAGTG ACAATTGGAGAATGCCCGAAATACGTGCGTAGTGCCAAGCTGAGAATGGTGACTGGTCTCCGGAATAACCCCTCTATACAGTCCA GGGGCCTGTTTGGGGCTATTGCCGGTTTTATTGAGGGAGGTTGGACTGGCATGATCGATGGATGGTACGGCTATCACCACCAGAA CGAGCAAGGCTCTGGCTATGCCGCAGACCAGAAATCAACACAAAACGCTATCAATGGAATAACTAACAAGGTCAACACAGTGATTG AAAAAATGAACATCCAGTTTACGGCTGTGGGCAAGGAGTTCAACAAGCTGGAAAAGAGGATGGAAAATCTCAACAAAAAAGTCGAC GACGGCTTCTTGGACATCTGGACTTACAATGCAGAGTTGCTTGTTCTGCTGGAAAACGAACGTACGCTGGACTTCCACGACAGTAA TGTCAAGAATTTGTATGAAAAAGTGAAATCACAGCTGAAAAATAATGCGAAGGAGATAGGCAATGGCTGTTTTGAGTTCTATCATAA GTGCGATAATGAGTGCATGGAGTCAGTGCGAAACGGGACCTACGACTACCCCAAATATAGTGAAGAGAGCAAGCTGAATAGAGAG AAAGTCGATGGCGTTAAACTTGAGAGCATGGGTATCTATCAAATCCTGGCTATTTACAGTACAGTGGCCTCCTCTCTAGTGCTGCT CGTATCTCTCGGAGCGATATCCTTCTGGATGTGTTCCAATGGCTCTCTACAGTGCCGGATATGCATA

2662	ATGAAAGCTAATCTGCTTGTACTGCTGTGCGCACTGGCAGCCGCAGACGCTGACACTATCTGTATCGGGTACCACGCCAACAATT CGACCGATACTGTGGACACAGTGTTGGAGAAGAACGTCACCGTAACGCACAGCGTGAACCTTTTAGAAGATAGTCATAATGGAAA GCTGTGTAGGCTCAAGGGAATCGCTCCACTTCAGTTGGGAAAATGCAATATCGCTGGCTGGCTACTGGGCAACCCTGAATGCGAC CCCCTGCTGCCTGTGCGATCCTGGTCCTATATTGTAGAGACCCCCAACTCAGAGAATGGCATTTGCTACCCAGGCGATTTTATTGA CTACGAGGAGCTGAGAGAGCAGTTGTCCTCCGTCTCCAGCTTTGAAAGGTTTGAGATTTTCCCAAAAGAGTCCAGCTGGCCAAAT CACAATACAAATGGTGTGACCGCCGCTTGTTCTCACGAAGGCAAAAGCTCCTTCTATAGAAATCTTCTGTGGCTAACAGAAAAAGA AGGATCATACCCAAAGTTGAAAAACAGCTACGTTAATAAGAAAGGCAAAGAGGTGCTGGTCCTGTGGGGTATCCACCACCCACCC AATTCTAAAGAACAACAAAATTTGTACCAAAATGAAAACGCCTACGTTTCTGTCGTCACTAGCAATTATAACCGGAGGTTCACACCC GAGATCGCAGAAAGACCTAAGGTGAGAGACCAAGCCGGCAGAATGAATTATTACTGGACTCTCCTCAAACCCGGGGACACCATCA TCTTCGAAGCTAACGGCAATCTGATTGCTCCTATGTACGCTTTTGCTCTGTCCCGGGGGTTTGGCAGCGGTATTATCACATCAAAC GCCAGTATGCACGAGTGTAATACGAAATGCCAGACTCCTCTGGGGGCTATCAATTCCAGTCTGCCCTATCAGAATATTCATCCTGT CACGATCGGTGAGTGCCCTAAGTATGTGCGAAGCGCCAAACTGCGGATGGTAACGGGCCTGAGGAATAACCCCTCTATTCAGTCT CGCGGGCTGTTTGGCGCAATCGCCGGCTTTATCGAGGGCGGTTGGACTGGCATGATCGACGGATGGTACGGGTACCATCACCAG AACGAACAGGGGTCCGGCTACGCCGCCGACCAGAAATCAACCCAGAACGCGATTAACGGGATCACCAACAAGGTGAATACTGTG ATTGAGAAGATGAATATTCAGTTTACAGCAGTGGGCAAGGAGTTTAACAAACTTGAGAAACGCATGGAGAATCTCAACAAAAAGGT TGACGATGGATTCCTTGATATTTGGACATATAATGCCGAGCTGCTGGTCCTTCTGGAAAATGAACGCACACTGGATTTTCACGACT CGAACGTGAAGAATCTGTACGAGAAGGTAAAGTCCCAATTAAAAAACAATGCTAAGGAGATAGGCAACGGCTGCTTCGAGTTTTAC CACAAGTGCGATAACGAGTGCATGGAGTCTGTGAGAAACGGTACGTACGATTACCCCAAATATTCCGAAGAGAGCAAGCTCAACC GCGAGAAAGTAGACGGCGTTAAACTGGAGAGCATGGGCATCTACCAGATCCTCGCCATTTACTCCACTGTGGCCTCGTCACTAGT TCTCTTGGTGTCCCTGGGAGCCATCTCATTTTGGATGTGCTCCAACGGCTCACTACAGTGCCGGATTTGTATC
2663	ATGAAGGCCAACCTCCTGGTGTTACTTTGTGCCCTAGCAGCCGCAGACGCCGACACTATCTGCATCGGATACCATGCTAATAACA GCACCGATACAGTAGATACCGTTTTGGAAAAGAATGTCACTGTCACCCACAGCGTTAATCTGTTGGAGGATAGCCACAACGGTAAA CTGTGCAGGCTGAAGGGAATTGCCCCCCTTCAACTGGGCAAGTGCAACATTGCAGGCTGGCTCCTCGGTAACCCTGAGTGCGAT CCTTTACTCCCTGTTAGAAGCTGGAGCTACATCGTGGAGACCCCCAATAGTGAGAACGGAATCTGCTACCCTGGAGATTTCATCGA CTATGAAGAACTTCGGGAACAGCTCAGTTCAGTGTCGAGCTTCGAGAGATTCGAGATCTTTCCAAAGGAGTCCAGCTGGCCTAATC ACAACACAAATGGCGTTACAGCAGCATGCTCACACGAAGGCAAGAGCAGCTTCTACCGCAACCTGCTGTGGCTTACCGAGAAGGA GGGATCCTATCCCAAGCTCAAGAATTCTTATGTGAATAAAAAGGGTAAGGAAGTGCTGGTCCTGTGGGGTATCCATCATCCGCCAA ATTCTAAGGAACAGCAGAATCTGTATCAGAATGAAAATGCCTACGTGAGTGTTGTGACTTCGAATTACAACAGGCGGTTTACCCCG GAGATTGCTGAGCGGCCCAAAGTCCGGGATCAGGCTGGCCGCATGAATTATTATTGGACTCTGCTCAAGCCTGGCGACACCATCA TTTTTGAGGCCAATGGCAACCTTATCGCTCCCATGTACGCCTTCGCCCTCAGTAGGGGATTCGGGTCTGGTATCATCACTTCAAAC GCCAGCATGCACGAGTGCAATACCAAGTGCCAGACACCACTCGGTGCCATTAATAGCTCTCTTCCATATCAGAATATACATCCAGT CACGATCGGGGAATGCCCCAAATATGTAAGAAGTGCCAAACTGCGCATGGTGACAGGCCTGCGGAACAACCCAAGCATCCAAAG CAGAGGGCTGTTTGGCGCCATCGCGGGCTTCATTGAAGGTGGTTGGACTGGCATGATTGATGGCTGGTACGGATATCATCATCAG AACGAGCAGGGATCAGGGTACGCGGCCGATCAGAAGAGTACCCAGAATGCCATTAACGGCATCACGAATAAAGTAAATACTGTCA TCGAAAAGATGAACATCCAATTTACTGCTGTCGGGAAGGAGTTCAACAAACTGGAAAAAAGGATGGAGAACCTGAATAAGAAAGTT GACGACGGCTTTTTAGATATTTGGACGTATAACGCAGAACTGCTCGTGCTCCTGGAGAACGAGAGGACCCTGGACTTTCACGACT CCAATGTGAAGAACCTTTATGAGAAAGTCAAATCTCAGCTCAAAAACAACGCTAAAGAAATTGGGAATGGTTGTTTTGAGTTTTATC ACAAATGCGACAACGAATGCATGGAGAGCGTCAGGAACGGGACATACGACTATCCAAAGTACTCAGAAGAGTCAAAACTCAACAG AGAGAAGGTTGACGGAGTCAAACTTGAGTCTATGGGCATATACCAGATCTTGGCGATCTATTCCACGGTGGCCTCAAGCCTTGTTC TGTTGGTCTCATTGGGTGCCATCAGCTTTTGGATGTGTAGTAACGGCAGTCTGCAGTGTAGAATTTGCATC

2664	ATGAAAGCAAACCTGCTCGTGCTCCTGTGCGCTCTGGCAGCCGCAGATGCCGACACTATTTGCATCGGTTACCATGCCAACAACA GCACTGATACGGTGGATACCGTACTAGAGAAGAATGTCACCGTAACCCACTCAGTGAACCTGCTCGAAGACTCTCACAACGGTAA GCTGTGTAGATTAAAGGGAATTGCCCCCTTGCAGTTAGGAAAGTGTAACATAGCAGGTTGGCTCCTTGGCAATCCTGAGTGTGAC CCCCTACTGCCAGTCCGCAGCTGGAGCTATATCGTTGAGACTCCCAACTCTGAAAATGGAATCTGTTATCCAGGTGACTTCATCGA TTATGAAGAATTACGCGAGCAGTTGTCCAGTGTTAGCTCTTTCGAGAGATTTGAAATATTCCCAAAGGAGTCGTCCTGGCCCAATC ACAACACCAATGGAGTGACCGCTGCTTGCTCCCACGAAGGGAAGAGCTCTTTCTATCGGAATCTGTTGTGGCTGACAGAAAAGGA GGGCAGCTATCCTAAACTAAAGAACAGCTACGTGAACAAGAAGGGAAAGGAGGTGCTCGTGCTGTGGGGCATTCATCATCCTCCA AATTCCAAGGAGCAGCAGAACTTATATCAGAATGAGAATGCTTATGTCTCAGTGGTGACGTCAAATTACAACAGGCGCTTCACACC CGAGATCGCTGAGCGGCCCAAGGTCCGCGACCAGGCTGGTAGAATGAATTACTATTGGACCCTCCTGAAGCCCGGAGATACTATT ATCTTCGAAGCAAATGGAAATTTGATCGCACCCATGTATGCTTTCGCTCTGTCCAGGGGATTCGGTAGTGGCATTATCACCAGCAA TGCAAGTATGCATGAGTGCAACACAAAATGTCAGACGCCGTTAGGGGCTATTAACAGCAGCCTCCCTTATCAGAACATTCATCCGG TTACCATCGGGGAGTGCCCTAAGTACGTGAGAAGTGCCAAACTGAGGATGGTGACAGGCCTGCGCAATAATCCCAGTATCCAGAG CCGAGGTCTCTTTGGTGCCATTGCGGGCTTTATTGAAGGGGGCTGGACCGGCATGATCGACGGTTGGTACGGATACCACCACCA GAACGAACAGGGCAGTGGCTACGCCGCCGACCAGAAGAGTACTCAGAACGCAATCAATGGAATTACGAACAAAGTTAATACCGTA ATCGAAAAAATGAATATCCAGTTTACCGCCGTAGGCAAGGAGTTTAACAAGCTCGAAAAACGCATGGAAAACCTGAATAAGAAAGT GGATGATGGATTCCTTGACATCTGGACCTACAACGCGGAACTTCTTGTCCTGCTCGAAAATGAGAGAACGCTCGACTTTCACGACA GCAATGTCAAGAATCTGTACGAAAAGGTAAAGTCACAGCTGAAGAACAACGCTAAGGAGATAGGCAACGGCTGCTTCGAGTTTTAC CATAAATGCGACAATGAATGCATGGAATCCGTGAGAAACGGCACTTACGATTACCCCAAATACAGCGAAGAATCAAAGCTAAACAG GGAGAAGGTGGACGGGGTGAAACTAGAAAGCATGGGCATCTATCAGATCCTCGCCATTTACTCCACAGTCGCATCTTCTTTAGTG CTCCTTGTGTCACTGGGCGCTATCTCGTTTTGGATGTGCTCAAACGGATCTCTGCAGTGTCGGATCTGTATT
2665	ATGAAAGCAAATCTGCTAGTGCTGCTGTGCGCCCTGGCCGCCGCAGATGCAGATACTATCTGCATTGGTTACCATGCCAACAACT CTACCGATACCGTGGATACTGTCCTCGAGAAGAATGTGACTGTGACTCATTCCGTGAACTTACTCGAAGACAGCCACAACGGAAA GCTGTGCCGATTGAAGGGCATCGCGCCACTCCAGCTGGGGAAATGCAACATCGCCGGCTGGCTTTTGGGAAACCCCGAGTGCGA TCCGCTATTGCCCGTAAGATCATGGTCCTACATCGTGGAAACACCAAATTCAGAAAATGGCATCTGCTACCCCGGAGATTTTATCG ATTACGAGGAACTGAGGGAGCAACTGAGCAGCGTTAGTTCTTTTGAGCGATTCGAAATTTTCCCAAAGGAGTCCTCATGGCCCAAT CACAATACCAATGGCGTTACAGCCGCTTGTAGCCATGAAGGTAAGAGTTCATTTTATCGAAATCTTCTCTGGCTGACGGAGAAGGA GGGTAGTTACCCTAAATTGAAAAATTCATACGTGAACAAGAAAGGAAAGGAGGTGCTGGTCCTCTGGGGGATTCACCATCCCCCC AATAGCAAAGAGCAACAGAATCTGTACCAGAACGAAAACGCTTATGTCTCCGTGGTCACCTCAAATTACAATAGACGGTTTACCCC GGAGATAGCCGAGAGGCCCAAAGTCCGCGATCAGGCCGGCCGCATGAACTACTACTGGACCCTCCTCAAGCCAGGGGATACCAT CATATTCGAGGCGAATGGGAACTTGATTGCCCCGATGTACGCATTTGCTTTGTCCAGAGGCTTTGGCTCCGGGATCATCACATCTA ATGCCTCCATGCATGAATGCAACACAAAATGCCAGACACCCCTGGGCGCTATTAACTCCTCTCTACCTTACCAGAACATCCACCCG GTTACTATTGGCGAGTGTCCTAAATATGTCCGCAGCGCCAAACTGAGGATGGTGACAGGGCTGAGGAATAACCCCAGCATTCAGT CCAGAGGCTTGTTCGGCGCTATAGCGGGGTTCATAGAAGGAGGCTGGACCGGAATGATCGATGGCTGGTACGGCTACCACCATC AGAACGAGCAGGGCTCCGGGTACGCCGCTGACCAGAAGAGTACGCAGAACGCCATTAATGGAATCACCAACAAGGTGAATACCG TAATCGAGAAAATGAATATTCAGTTCACAGCTGTGGGGAAGGAATTCAATAAGCTGGAAAAACGCATGGAGAACCTGAATAAAAAA GTGGATGATGGCTTCTTGGATATCTGGACCTATAATGCGGAACTCCTGGTCCTTTTGGAAAATGAGCGCACACTTGACTTTCATGA CTCAAACGTTAAAAACCTGTACGAAAAAGTCAAGTCACAGCTAAAGAATAATGCAAAGGAGATTGGCAACGGGTGCTTCGAATTTT ATCATAAATGCGACAACGAATGCATGGAAAGCGTAAGGAATGGTACATACGATTATCCTAAGTACAGTGAGGAGTCAAAGTTGAAT CGCGAAAAAGTTGACGGTGTCAAACTGGAATCCATGGGGATTTATCAGATTCTTGCAATTTACTCAACGGTGGCTAGTTCTCTGGT GCTGCTGGTCTCCCTGGGCGCAATTAGTTTCTGGATGTGTAGCAACGGCTCTCTCCAGTGCCGGATCTGTATT

2666	ATGAAAGCAAATCTTCTGGTCCTGCTGTGTGCACTAGCCGCCGCCGACGCAGACACAATTTGCATCGGATATCACGCCAACAACT CTACGGATACAGTGGACACAGTGCTGGAGAAGAACGTAACTGTTACACACAGCGTGAACCTGCTAGAAGATTCTCACAATGGCAA ACTGTGCAGACTGAAAGGAATTGCACCCTTGCAGTTGGGAAAATGTAATATCGCGGGTTGGCTCCTCGGTAATCCCGAATGCGAC CCCCTACTTCCCGTCCGTTCTTGGTCTTACATAGTGGAGACACCCAACTCGGAGAATGGGATTTGCTATCCTGGGGATTTTATTGA CTACGAGGAGCTCAGGGAACAGTTGAGTTCTGTGTCATCATTCGAGAGGTTTGAAATCTTTCCCAAAGAGTCTTCTTGGCCTAATC ACAACACTAACGGTGTGACCGCTGCTTGTTCTCACGAAGGCAAATCATCCTTCTACAGAAATCTGCTCTGGCTGACGGAGAAGGAA GGATCATACCCAAAGCTAAAGAATTCATATGTGAATAAAAAGGGGAAGGAGGTCCTCGTTCTGTGGGGTATTCACCATCCCCCTAA TTCTAAAGAACAGCAGAACCTTTATCAGAATGAAAATGCATACGTGTCAGTGGTGACAAGCAACTATAACAGGCGCTTTACCCCGG AAATCGCCGAACGCCCGAAGGTGAGGGACCAGGCCGGGAGGATGAACTACTACTGGACCTTGCTGAAGCCGGGCGACACAATCA TTTTTGAAGCTAACGGGAATCTAATTGCTCCCATGTACGCCTTTGCTCTGTCTCGCGGCTTTGGCAGCGGGATTATCACTAGCAAC GCGTCCATGCACGAGTGTAATACTAAATGCCAGACGCCACTGGGCGCAATCAATAGCAGCCTTCCATACCAGAACATACATCCCG TCACCATAGGAGAATGTCCAAAGTACGTCCGGTCTGCAAAACTAAGGATGGTGACCGGGCTTAGAAACAACCCAAGTATTCAGAG CCGGGGGCTCTTTGGTGCCATCGCAGGCTTTATCGAGGGAGGGTGGACAGGCATGATAGACGGATGGTATGGATACCATCACCA GAATGAGCAGGGCAGCGGATACGCTGCCGACCAAAAAAGCACCCAAAACGCTATCAATGGCATTACTAATAAGGTGAACACTGTG ATTGAGAAGATGAACATTCAGTTCACCGCAGTCGGGAAGGAATTTAACAAGCTGGAGAAAAGGATGGAGAACTTAAACAAGAAGG TTGATGACGGGTTTCTCGACATATGGACGTATAACGCGGAGTTGCTGGTATTACTGGAGAATGAACGGACTCTTGACTTCCACGAT TCAAATGTGAAGAATCTCTATGAAAAAGTCAAATCGCAGCTCAAAAATAATGCCAAGGAGATCGGCAACGGCTGTTTCGAGTTTTAT CACAAATGCGACAACGAATGTATGGAATCAGTGCGAAATGGCACATATGACTATCCGAAATACTCCGAGGAAAGTAAGCTGAATAG AGAAAAGGTGGACGGAGTAAAGCTCGAGTCCATGGGGATCTACCAGATTTTGGCTATCTATTCCACAGTTGCTTCATCTCTGGTGC TGCTGGTTTCCTTAGGCGCCATATCGTTCTGGATGTGCTCCAACGGGTCCCTGCAGTGTCGAATCTGCATC
2667	ATGAAGGCGAATCTACTTGTCCTGCTGTGCGCGCTTGCGGCTGCTGACGCCGACACGATCTGCATAGGCTACCATGCAAATAACT CCACTGACACCGTCGACACCGTTCTTGAGAAGAATGTGACGGTTACTCACTCCGTGAACCTCTTAGAGGATAGTCACAATGGCAA GTTGTGCAGATTAAAAGGCATAGCTCCTCTTCAGCTGGGGAAATGTAATATCGCCGGGTGGCTCCTCGGGAACCCTGAATGCGAT CCTCTGCTACCTGTGAGGAGTTGGAGCTACATAGTCGAAACTCCTAACTCGGAGAATGGAATTTGTTACCCTGGAGACTTCATTGA TTACGAAGAGCTGCGGGAACAATTGTCCTCTGTGTCATCATTTGAGCGGTTTGAGATTTTCCCCAAAGAAAGTAGTTGGCCAAATC ATAACACTAACGGGGTCACCGCCGCATGTTCCCATGAAGGAAAGAGCTCTTTTTATCGAAACCTCCTCTGGCTTACTGAGAAGGAA GGATCCTATCCCAAGCTCAAGAACTCCTACGTGAATAAAAAGGGCAAAGAAGTTCTGGTGCTGTGGGGTATTCACCATCCCCCCAA CTCAAAAGAGCAACAGAACCTCTACCAGAATGAGAATGCGTACGTGTCAGTGGTTACTAGCAACTACAATAGGCGGTTTACACCGG AGATCGCCGAGCGTCCTAAAGTGAGAGACCAGGCTGGAAGAATGAACTATTATTGGACGCTGCTCAAACCCGGTGACACTATCAT TTTCGAGGCTAATGGGAACTTGATTGCCCCTATGTATGCTTTCGCCTTATCACGCGGATTTGGCTCCGGGATCATTACCTCAAATG CCTCAATGCACGAGTGTAACACCAAGTGTCAGACACCCCTGGGCGCGATTAACAGCAGTCTGCCGTACCAAAATATCCACCCCGT GACGATTGGGGAGTGCCCTAAGTACGTGCGGAGTGCCAAACTGCGTATGGTGACAGGGCTGCGCAACAATCCTTCCATTCAGTC GAGAGGCCTGTTTGGCGCCATCGCCGGGTTCATCGAAGGGGGGTGGACCGGAATGATTGATGGATGGTATGGGTACCATCATCA GAATGAGCAGGGGTCTGGATATGCAGCAGACCAGAAATCGACTCAAAATGCAATAAATGGCATAACGAATAAAGTTAATACAGTGA TAGAGAAGATGAATATCCAGTTTACTGCCGTGGGCAAGGAGTTCAATAAACTCGAGAAGCGCATGGAGAACCTTAACAAAAAGGTG GACGATGGATTCCTGGATATCTGGACTTATAATGCCGAGCTGCTGGTACTTCTCGAAAATGAGCGTACACTAGACTTCCATGACTC AAATGTAAAAAATTTATATGAGAAGGTGAAATCCCAGCTCAAGAACAATGCAAAGGAGATCGGCAATGGGTGTTTCGAATTTTATCA CAAGTGTGACAACGAGTGCATGGAATCAGTGAGAAACGGAACATACGACTACCCTAAATATAGTGAAGAGTCTAAGCTGAATCGTG AGAAAGTGGACGGGGTGAAACTCGAGTCTATGGGCATTTACCAGATTCTGGCTATTTATAGTACAGTGGCTTCAAGCTTAGTCCTT CTAGTGAGCTTGGGTGCCATTTCATTTTGGATGTGTAGTAATGGGTCATTGCAGTGTAGGATTTGTATC

2668	ATGAAAGCAAACCTGCTGGTCTTATTGTGCGCCTTGGCAGCTGCCGATGCCGACACCATCTGCATCGGGTACCACGCCAATAATA GTACTGACACCGTCGATACCGTGCTCGAAAAGAACGTAACTGTGACTCACAGTGTGAACCTGCTGGAGGATAGCCATAACGGCAA ACTCTGCCGACTGAAAGGAATCGCTCCCTTGCAACTGGGCAAGTGCAACATAGCAGGGTGGTTACTGGGGAACCCCGAGTGCGA CCCGCTGCTCCCTGTAAGATCTTGGAGCTATATCGTGGAAACACCTAACTCAGAAAATGGGATATGTTACCCAGGTGACTTTATCG ATTACGAAGAATTGAGGGAACAGCTGAGCAGCGTTAGTAGCTTTGAACGATTTGAGATCTTCCCAAAGGAGTCCTCTTGGCCTAAT CACAACACAAATGGGGTGACCGCCGCTTGCTCTCATGAGGGTAAAAGTAGCTTTTACCGGAACTTACTATGGCTGACCGAGAAAG AAGGAAGCTACCCCAAACTGAAAAATTCGTACGTGAATAAAAAGGGGAAAGAGGTGTTAGTGCTCTGGGGGATACATCACCCTCC AAATTCCAAGGAGCAGCAGAATCTGTACCAAAACGAGAACGCCTACGTTTCCGTGGTGACTAGCAATTATAACAGGCGGTTTACAC CTGAAATCGCCGAACGGCCTAAAGTTCGAGATCAGGCCGGGCGGATGAACTACTATTGGACTTTGCTCAAGCCTGGCGACACAAT CATTTTTGAAGCCAACGGGAACCTGATCGCACCGATGTATGCTTTTGCACTGAGTCGGGGGTTTGGCTCTGGCATTATCACCTCTA ATGCGTCCATGCACGAATGTAATACCAAGTGTCAGACACCCCTGGGGGCTATAAACTCCTCCCTCCCCTATCAGAACATCCACCCT GTGACCATCGGAGAATGTCCTAAATATGTCCGCTCCGCCAAACTGCGGATGGTTACTGGCCTCAGAAATAATCCAAGCATTCAGAG TCGAGGCCTCTTCGGCGCGATCGCCGGCTTCATTGAGGGTGGCTGGACCGGCATGATCGATGGGTGGTACGGGTATCATCATCA GAATGAACAGGGATCCGGCTACGCAGCAGATCAAAAAAGCACACAGAACGCGATTAATGGCATTACTAACAAGGTGAACACGGTG ATCGAGAAGATGAATATACAATTCACTGCTGTCGGGAAAGAGTTCAATAAGCTCGAAAAGAGGATGGAGAATTTGAACAAGAAAGT GGATGATGGCTTCCTGGACATTTGGACCTACAATGCTGAACTCCTGGTCCTTCTGGAAAACGAAAGAACCCTAGATTTCCATGACT CTAACGTGAAGAACCTTTACGAAAAGGTGAAGAGCCAGTTGAAAAACAACGCAAAGGAAATTGGAAATGGGTGCTTCGAGTTTTAT CATAAGTGCGACAACGAGTGCATGGAAAGTGTCCGTAATGGCACTTACGATTACCCCAAATATTCTGAAGAATCCAAGCTGAACAG GGAGAAAGTCGATGGGGTTAAATTAGAATCCATGGGCATCTATCAGATACTGGCCATTTATTCCACTGTGGCGAGCTCCTTAGTGC TCTTGGTCTCTTTAGGTGCCATCTCCTTTTGGATGTGCAGTAACGGGTCCCTCCAATGCAGGATTTGTATC
2669	ATGAAGGCCAACCTCCTCGTTCTGCTGTGTGCGCTGGCAGCGGCAGATGCCGATACCATTTGTATCGGTTACCATGCCAACAATT CCACCGACACCGTCGACACTGTTCTTGAAAAGAATGTAACTGTCACCCATTCTGTCAACTTGCTTGAAGATAGTCACAACGGCAAA CTGTGTCGGCTTAAAGGCATAGCCCCGCTGCAACTGGGCAAATGTAACATCGCCGGATGGCTGCTGGGAAATCCAGAATGCGAC CCCCTGTTGCCTGTGCGGTCTTGGAGTTACATAGTAGAAACCCCCAATTCCGAAAATGGGATTTGTTATCCTGGCGACTTCATCGA TTATGAGGAGTTGAGAGAGCAATTGAGTAGCGTTTCTTCATTTGAGCGCTTCGAAATATTCCCGAAAGAGAGTTCATGGCCTAACC ACAACACAAATGGAGTGACAGCTGCCTGTTCCCACGAAGGAAAAAGCTCCTTTTACCGTAACCTGCTATGGCTGACTGAAAAGGA GGGGTCCTACCCCAAACTTAAGAACTCTTATGTCAACAAGAAGGGGAAAGAGGTACTGGTTCTGTGGGGAATTCATCATCCCCCG AATTCAAAAGAGCAGCAAAATCTGTACCAAAATGAAAACGCGTACGTCTCCGTTGTGACATCTAACTACAATAGGCGGTTCACGCC CGAGATCGCTGAACGGCCGAAAGTCCGGGACCAAGCTGGCAGAATGAACTATTACTGGACACTGCTCAAACCCGGCGACACCAT TATTTTCGAGGCAAATGGGAACCTGATTGCACCAATGTACGCCTTCGCTCTGTCCAGGGGCTTTGGTAGCGGAATTATCACCTCGA ATGCTTCAATGCATGAGTGCAACACCAAGTGCCAGACACCTTTGGGCGCCATCAATAGTAGCCTTCCTTACCAGAACATTCACCCT GTCACCATTGGTGAATGTCCTAAGTACGTGCGAAGCGCCAAACTGCGGATGGTTACTGGCCTGCGAAATAATCCCTCCATCCAGT CTCGCGGTCTGTTCGGAGCCATTGCAGGGTTCATCGAGGGAGGCTGGACCGGGATGATTGACGGCTGGTATGGGTACCACCATC AGAACGAGCAAGGATCCGGGTATGCCGCCGACCAGAAGAGCACCCAGAACGCTATCAATGGGATTACAAATAAAGTCAACACCGT CATCGAGAAGATGAACATCCAATTCACTGCGGTTGGCAAGGAGTTCAACAAACTCGAGAAAAGAATGGAAAACTTGAATAAGAAGG TAGACGACGGGTTTCTGGATATCTGGACTTATAACGCCGAATTATTGGTGCTGCTGGAGAATGAACGGACTTTAGATTTCCACGAC TCCAATGTGAAGAATCTATACGAGAAAGTGAAGTCTCAGTTGAAGAACAACGCCAAGGAGATAGGAAATGGCTGTTTCGAGTTTTA TCACAAGTGCGATAATGAGTGCATGGAAAGCGTGCGCAACGGGACATATGATTACCCGAAGTACTCCGAGGAAAGCAAACTAAAT CGCGAAAAGGTAGATGGCGTGAAGCTGGAATCGATGGGAATCTACCAGATCCTAGCTATTTACTCTACTGTTGCCAGCTCTCTGGT GCTTCTGGTATCTCTGGGGGCCATTAGTTTCTGGATGTGCAGTAACGGCTCATTACAGTGTAGGATATGCATT

2670	ATGAAAGCAAACTTGCTGGTTCTGTTGTGCGCTCTCGCGGCCGCTGACGCGGACACTATCTGCATCGGGTACCACGCGAATAATT CGACTGACACAGTTGACACTGTGCTCGAAAAGAATGTGACCGTCACGCACTCCGTCAATCTCCTGGAGGACTCACACAATGGCAA ACTTTGTCGTTTGAAGGGCATCGCACCCCTCCAACTCGGGAAATGCAACATTGCAGGTTGGCTTTTAGGTAATCCTGAGTGTGATC CCCTGCTGCCTGTGCGCAGCTGGAGCTATATTGTCGAGACACCCAACTCCGAAAATGGAATCTGTTACCCAGGAGACTTCATTGA CTATGAAGAATTAAGAGAGCAATTAAGCTCCGTGAGCTCCTTTGAGAGGTTTGAGATCTTCCCGAAGGAAAGCTCCTGGCCTAACC ATAACACCAATGGAGTGACTGCCGCCTGCAGCCACGAAGGAAAGTCTTCCTTCTATAGAAACCTACTGTGGTTGACAGAGAAGGA AGGCTCCTATCCTAAATTGAAGAACTCATATGTCAACAAGAAGGGGAAAGAGGTCCTCGTGCTGTGGGGAATCCATCATCCTCCTA ACTCTAAGGAGCAACAAAATTTATACCAGAACGAAAATGCATACGTTTCTGTGGTTACATCAAACTATAATAGGCGCTTCACACCTG AGATTGCTGAACGTCCCAAAGTGAGGGATCAAGCGGGGCGTATGAACTACTACTGGACACTGTTAAAACCGGGGGATACCATCAT CTTTGAGGCGAACGGCAATCTTATTGCACCTATGTATGCTTTTGCCCTCTCGCGCGGTTTCGGCAGTGGAATAATAACATCAAATG CCAGTATGCACGAGTGCAATACGAAGTGCCAGACTCCCTTAGGCGCTATAAATAGTTCCTTACCTTACCAGAATATTCACCCAGTG ACCATCGGCGAATGCCCAAAGTACGTCAGGTCTGCGAAACTGAGAATGGTCACAGGGTTGAGAAATAATCCATCTATCCAGTCAC GAGGCCTGTTTGGCGCGATCGCGGGCTTTATTGAAGGAGGGTGGACGGGTATGATTGACGGATGGTACGGTTACCATCATCAAAA TGAACAAGGTTCAGGGTATGCCGCGGACCAGAAGAGCACCCAGAATGCAATCAATGGCATTACCAACAAGGTGAACACGGTGATC GAGAAGATGAATATTCAGTTTACCGCTGTGGGCAAGGAGTTTAATAAGCTCGAGAAACGCATGGAAAATCTGAACAAAAAAGTAGA CGATGGATTCCTCGACATATGGACGTACAATGCCGAGCTTCTCGTCCTCTTAGAGAACGAGCGGACCCTGGATTTCCACGATTCAA ACGTCAAAAACCTGTATGAGAAAGTCAAGAGCCAGCTTAAGAATAACGCCAAGGAAATCGGGAACGGCTGCTTCGAGTTCTATCAT AAATGCGACAATGAGTGTATGGAATCTGTACGCAATGGAACATACGACTACCCTAAGTACTCCGAAGAAAGCAAACTGAATAGGGA GAAGGTAGACGGAGTGAAATTGGAGAGCATGGGCATATACCAGATCCTGGCCATTTACTCAACCGTCGCCTCATCCTTGGTGTTG CTGGTGTCCTTGGGTGCTATTTCTTTTTGGATGTGCAGCAATGGCAGTCTCCAGTGCAGGATTTGTATC
2671	ATGAAGGCTAACCTGCTGGTGCTCCTCTGCGCTCTGGCAGCTGCAGACGCTGATACCATTTGCATTGGCTACCATGCCAATAACA GCACCGATACAGTCGATACAGTTCTGGAGAAGAACGTGACAGTGACTCACAGCGTTAATCTGTTGGAGGACAGCCATAATGGGAA GCTGTGTCGCCTGAAGGGAATTGCTCCATTACAGCTGGGGAAATGTAACATCGCTGGATGGCTGTTGGGAAACCCAGAGTGTGAT CCCCTACTTCCTGTACGGAGTTGGAGTTATATCGTGGAGACACCCAATAGTGAGAACGGTATCTGTTATCCTGGAGATTTTATAGA CTATGAAGAGTTAAGGGAGCAGCTGAGCTCCGTCAGTTCATTCGAACGCTTCGAAATTTTCCCTAAGGAGTCTTCCTGGCCCAACC ACAATACCAACGGGGTCACCGCTGCTTGTAGTCATGAAGGAAAATCTAGCTTCTACCGAAACCTGCTCTGGCTAACTGAGAAAGAA GGATCATACCCGAAGCTGAAGAATAGCTACGTGAACAAGAAGGGGAAGGAAGTGCTGGTTCTGTGGGGCATTCACCACCCCCCTA ACAGTAAGGAACAGCAGAACCTCTATCAGAATGAGAACGCATACGTGAGTGTCGTTACAAGCAACTACAATCGCCGGTTTACTCCA GAGATCGCGGAAAGACCCAAAGTCAGGGACCAGGCTGGTAGGATGAATTACTATTGGACCCTTCTCAAACCTGGGGACACTATCA TATTCGAGGCCAATGGCAACCTGATTGCCCCTATGTACGCCTTTGCTCTGAGCCGCGGATTCGGCTCCGGCATAATCACGTCTAAT GCTTCCATGCACGAGTGTAACACAAAATGTCAGACGCCCTTGGGCGCTATCAATTCTAGTCTGCCTTATCAGAACATACATCCAGT CACTATCGGAGAGTGCCCAAAGTACGTTCGCTCTGCAAAGCTCCGGATGGTTACAGGTCTAAGAAATAATCCTAGTATTCAGAGCA GGGGCCTGTTTGGAGCCATAGCTGGATTTATAGAGGGCGGATGGACTGGGATGATAGATGGGTGGTATGGCTACCATCACCAGA ATGAACAGGGGAGCGGATACGCTGCCGACCAAAAATCTACTCAGAACGCAATTAATGGTATCACAAACAAAGTGAACACAGTGATC GAAAAGATGAACATTCAGTTCACTGCTGTGGGGAAAGAGTTCAACAAGCTGGAGAAACGAATGGAAAATCTGAATAAAAAGGTGGA CGACGGATTCCTTGACATTTGGACTTACAACGCCGAATTACTTGTCCTCTTGGAAAACGAACGGACCCTGGATTTTCATGACTCCA ATGTTAAGAACCTGTATGAGAAGGTGAAGAGCCAGCTGAAAAATAACGCAAAGGAAATCGGGAACGGATGTTTTGAGTTTTACCAC AAGTGCGATAATGAATGTATGGAGAGTGTTCGGAATGGGACATACGACTACCCTAAGTATTCTGAGGAGTCTAAGCTCAACCGGG AGAAGGTGGATGGTGTTAAGTTAGAGAGTATGGGAATTTACCAGATTCTGGCCATATACTCTACAGTAGCATCTTCTCTGGTGTTAT TGGTCAGTCTCGGTGCTATTTCTTTTTGGATGTGCTCAAACGGCTCACTGCAGTGCCGCATCTGTATC

2672	ATGAAAGCAAACCTGCTGGTCCTCTTGTGCGCGCTGGCAGCCGCCGATGCTGACACCATATGTATTGGCTATCACGCTAACAATT CTACGGACACTGTGGATACGGTGCTGGAGAAAAACGTAACCGTCACACACAGCGTGAATCTGCTGGAGGATTCGCATAATGGTAA ACTTTGCAGACTGAAGGGGATAGCCCCGCTCCAGTTGGGTAAGTGTAATATCGCCGGATGGTTGCTTGGGAACCCCGAGTGTGAT CCTCTGTTGCCGGTGCGCAGTTGGTCGTACATCGTGGAAACCCCGAATAGTGAGAATGGCATTTGTTATCCAGGGGACTTCATCG ATTACGAAGAGCTCAGGGAGCAGCTCAGCTCTGTAAGTAGCTTCGAACGGTTCGAGATCTTCCCCAAAGAAAGCTCCTGGCCCAA TCACAACACTAACGGGGTGACAGCGGCCTGCAGCCACGAGGGCAAAAGTTCTTTCTATCGCAACCTACTGTGGCTCACCGAGAAG GAGGGAAGTTACCCTAAGCTGAAGAACAGCTATGTCAACAAGAAAGGCAAGGAGGTCCTTGTGTTATGGGGCATTCACCACCCTC CCAATAGCAAAGAGCAGCAAAACCTTTATCAAAATGAGAACGCCTACGTGAGCGTGGTGACCTCTAACTACAACAGAAGGTTTACA CCAGAGATCGCCGAAAGGCCTAAAGTTAGGGACCAGGCCGGACGGATGAATTACTATTGGACGTTATTAAAGCCTGGGGATACCA TTATCTTTGAGGCCAATGGCAACTTGATCGCCCCTATGTACGCCTTTGCTCTCTCGCGCGGATTTGGTTCCGGAATCATTACCAGC AATGCTTCAATGCATGAGTGCAATACAAAGTGTCAGACCCCACTGGGCGCTATTAACTCGTCTTTACCGTATCAAAATATACATCCC GTGACTATTGGTGAGTGCCCTAAATATGTTCGCTCTGCTAAGCTCAGGATGGTTACTGGACTGCGGAACAATCCGTCGATTCAGTC CAGGGGACTCTTCGGCGCAATTGCGGGTTTCATCGAGGGTGGCTGGACGGGCATGATCGACGGCTGGTACGGTTACCACCATCA GAACGAACAGGGATCCGGCTACGCAGCCGATCAAAAGTCTACGCAGAACGCTATTAATGGAATCACCAACAAGGTGAATACAGTT ATCGAAAAGATGAATATACAGTTTACTGCCGTGGGGAAAGAATTCAACAAACTGGAGAAGAGAATGGAGAATCTTAACAAGAAGGT CGACGATGGCTTCCTTGATATCTGGACGTACAATGCCGAGCTGCTGGTCCTCCTCGAGAATGAACGAACTCTGGACTTCCACGAC TCCAATGTAAAGAATCTCTATGAAAAGGTGAAATCCCAGCTGAAAAATAACGCCAAGGAGATAGGAAATGGCTGCTTCGAATTTTAC CACAAATGTGACAATGAATGCATGGAGTCCGTGAGAAATGGAACCTACGACTATCCCAAATATTCAGAAGAGTCAAAGCTAAATCG CGAAAAAGTTGACGGAGTCAAGCTCGAGTCTATGGGCATATATCAGATTCTGGCCATCTACTCCACCGTGGCCTCTTCCCTTGTCC TTCTAGTGTCCTTGGGTGCAATTTCTTTTTGGATGTGCAGCAACGGGTCCCTGCAATGCAGGATTTGTATA
2673	ATGAAGGCCAACCTTCTCGTCTTACTATGTGCACTTGCCGCAGCAGATGCCGATACTATCTGCATCGGCTATCATGCCAATAACAG CACAGACACCGTGGATACGGTGCTGGAGAAGAACGTTACAGTGACCCACTCCGTTAATCTGCTTGAAGATAGTCATAACGGCAAG CTGTGTCGCTTGAAAGGGATCGCCCCTCTCCAGCTGGGTAAGTGCAACATTGCTGGGTGGCTCCTTGGCAACCCTGAGTGTGACC CACTACTTCCTGTGAGAAGTTGGAGCTATATCGTAGAGACACCAAATTCTGAAAACGGAATCTGTTATCCCGGAGACTTCATCGAC TACGAGGAACTGCGCGAACAACTGTCTTCCGTCTCTTCCTTTGAGAGATTTGAGATCTTTCCCAAAGAAAGTAGTTGGCCAAACCA TAATACTAACGGCGTGACAGCAGCCTGTAGTCACGAAGGCAAGTCCAGTTTTTACCGCAACCTGCTCTGGCTGACAGAAAAGGAG GGGAGTTATCCAAAGCTGAAAAACTCTTACGTCAACAAGAAAGGGAAAGAGGTCCTGGTTCTTTGGGGCATTCATCATCCCCCAAA TTCAAAGGAACAGCAGAATCTTTACCAGAACGAAAATGCTTACGTGAGTGTGGTCACCAGTAACTACAACAGGAGGTTCACACCAG AGATAGCAGAACGGCCAAAAGTAAGAGATCAGGCTGGTAGGATGAACTATTACTGGACACTGCTCAAGCCGGGTGACACCATAAT CTTTGAGGCTAATGGCAACCTTATTGCTCCCATGTACGCTTTCGCTCTTAGCAGAGGGTTTGGCAGCGGCATTATCACATCAAATG CAAGCATGCACGAGTGCAACACCAAGTGTCAAACACCCCTCGGCGCAATAAATAGCAGCTTGCCATATCAGAATATACACCCCGT GACAATCGGCGAGTGTCCTAAATACGTACGCTCTGCAAAACTCCGCATGGTGACCGGCCTCCGTAACAACCCCAGCATCCAGAGT CGGGGGCTCTTTGGCGCCATCGCCGGGTTTATTGAAGGGGGGTGGCACAGGGATGATTGATGGTTGGTATGGATATCACCACCAG AACGAACAAGGATCAGGTTACGCGGCGGACCAGAAGAGCACCCAAAACGCTATAAACGGCATAACCAACAAGGTAAACACCGTTA TTGAAAAGATGAATATTCAGTTTACCGCAGTCGGTAAAGAGTTTAATAAGCTCGAGAAGAGGATGGAAAATCTGAACAAGAAGGTT GATGATGGGTTTTTAGATATTTGGACCTACAATGCGGAGCTGCTCGTCCTGTTAGAAAATGAGAGGACACTGGACTTCCACGATAG TAATGTGAAGAACCTCTATGAGAAGGTGAAAAGTCAGCTCAAGAATAACGCTAAAGAGATTGGGAATGGTTGTTTTGAATTTTATCA CAAATGCGACAATGAATGTATGGAGAGTGTAAGGAACGGTACGTACGACTACCCAAAATATTCAGAAGAGTCTAAACTCAATAGAG AGAAGGTTGACGGAGTGAAACTGGAGTCCATGGGCATTTACCAGATCCTGGCAATCTACTCTACTGTTGCTAGCTCGTTGGTGTTG CTCGTCAGTCTGGGAGCTATCTCCTTCTGGATGTGTAGTAATGGGAGCCTGCAGTGCCGGATCTGTATT

2674	ATGAAGGCTAATCTACTGGTTCTTCTGTGCGCGCTCGCCGCAGCAGATGCTGATACTATCTGCATCGGGTATCATGCTAATAACAG CACCGACACCGTCGATACGGTACTAGAGAAAAACGTTACTGTGACTCACTCCGTGAACCTGCTGGAAGACTCACATAACGGCAAA CTGTGTCGCCTGAAGGGGATTGCCCCCCTGCAGCTGGGGAAGTGTAATATAGCGGGATGGCTGTTGGGGAATCCTGAGTGTGAT CCACTTCTCCCCGTGAGATCCTGGAGTTACATTGTGGAGACTCCAAATTCGGAGAACGGGATCTGCTATCCAGGAGACTTCATCG ACTACGAAGAATTGCGAGAGCAACTGTCTAGCGTGTCCTCCTTTGAAAGATTCGAGATTTTTCCAAAGGAGTCCAGTTGGCCTAAC CATAACACTAACGGCGTGACTGCCGCTTGTTCTCACGAGGGCAAGTCTTCTTTTTATAGAAACCTTCTCTGGCTCACCGAAAAAGA AGGTTCCTATCCCAAACTCAAAAACTCCTACGTGAATAAGAAAGGCAAAGAAGTCCTGGTGTTGTGGGGTATACACCATCCGCCTA ACTCCAAGGAACAGCAGAATCTCTACCAGAACGAGAATGCATACGTGTCAGTTGTCACCTCCAACTACAACCGGCGGTTCACCCC TGAGATCGCCGAAAGGCCAAAAGTGCGTGACCAGGCCGGACGCATGAACTACTACTGGACTCTCTTGAAGCCAGGAGACACCATT ATCTTCGAAGCCAATGGCAACCTGATTGCCCCCATGTACGCTTTCGCACTCAGCAGAGGTTTCGGGTCCGGCATAATTACTAGCAA CGCCAGCATGCACGAATGCAACACTAAATGTCAGACCCCACTCGGAGCCATCAACTCCAGTCTGCCCTACCAGAATATTCACCCT GTCACCATCGGCGAATGCCCCAAGTACGTGAGGAGTGCTAAACTTCGAATGGTGACCGGCCTGCGGAACAATCCGAGCATTCAGT CCCGGGGATTGTTTGGAGCCATCGCTGGATTTATAGAGGGGGGCTGGACCGGCATGATCGACGGCTGGTATGGCTATCATCATC AAAATGAACAGGGCAGCGGCTATGCAGCTGACCAAAAGTCAACCCAGAACGCTATTAACGGAATCACAAACAAGGTGAATACCGT GATAGAAAAGATGAATATTCAGTTCACGGCTGTGGGAAAGGAATTCAACAAGTTAGAGAAGAGGATGGAAAATCTGAATAAGAAGG TGGACGACGGGTTCTTGGATATCTGGACATACAACGCTGAGCTCTTGGTGCTGCTCGAAAATGAACGGACTCTTGACTTTCACGAC TCAAACGTAAAAAATCTCTATGAAAAGGTGAAATCTCAGTTGAAGAATAACGCAAAGGAGATTGGCAATGGGTGCTTCGAATTCTAT CACAAGTGTGATAACGAGTGTATGGAGTCAGTCCGGAATGGGACTTACGACTATCCCAAGTATAGTGAAGAATCTAAGCTCAATCG CGAAAAGGTGGACGGGGTGAAGTTGGAATCAATGGGTATTTATCAGATCCTCGCAATTTACTCTACCGTGGCCTCCTCATTGGTTT TGCTGGTGTCCCTTGGAGCTATAAGCTTCTGGATGTGCAGTAATGGGTCACTACAGTGCCGTATTTGCATT
2675	ATGAAGGCCAACTTGCTGGTGCTTCTTTGCGCCCTCGCAGCAGCTGACGCCGATACCATCTGTATAGGCTATCACGCAAACAATA GCACCGATACTGTGGATACCGTGCTTGAGAAGAACGTTACAGTGACACATTCAGTGAACCTACTCGAGGACTCCCACAATGGAAA GCTGTGTAGGCTGAAAGGCATTGCACCACTGCAGCTCGGGAAATGTAATATAGCAGGGTGGCTCCTCGGCAATCCCGAGTGCGA TCCCCTGCTGCCCGTCAGGTCATGGAGCTACATTGTAGAAACCCCGAACTCTGAGAATGGCATTTGTTATCCCGGAGATTTCATCG ACTACGAAGAACTGCGAGAGCAACTCTCATCCGTGTCTTCATTTGAAAGGTTCGAGATCTTCCCCAAGGAGTCTAGCTGGCCTAAT CACAACACAAACGGCGTGACTGCCGCCTGTTCACACGAGGGTAAGAGTTCATTTTACAGGAACCTCCTGTGGCTGACTGAGAAGG AAGGCAGCTATCCAAAGTTGAAGAATTCGTACGTGAACAAGAAGGGGAAAGAAGTTCTGGTACTCTGGGGGATCCACCACCCGCC TAATTCCAAAGAACAGCAGAATCTATACCAGAACGAAAATGCTTACGTCAGCGTTGTGACATCCAATTACAACAGAAGATTTACCCC AGAGATAGCCGAGCGACCCAAGGTGAGGGACCAGGCGGGACGAATGAATTATTACTGGACCCTGTTGAAGCCAGGAGACACCAT CATTTTTGAAGCCAACGGAAATCTTATCGCGCCGATGTACGCGTTCGCACTGTCACGAGGATTTGGCTCGGGGATTATTACCTCAA ATGCTAGCATGCACGAATGCAACACTAAGTGTCAGACTCCGCTGGGAGCTATCAATTCATCTCTCCCATATCAGAATATTCATCCC GTGACCATAGGCGAGTGCCCCAAATACGTCCGGTCGGCAAAATTGCGGATGGTTACTGGTCTGAGAAATAATCCATCCATACAGT CTCGCGGTTTATTTGGGGCGATTGCAGGCTTTATTGAGGGGGGATGGACGGGCATGATCGATGGTTGGTACGGATATCACCACCA AAACGAGCAAGGGAGTGGTTATGCCGCCGATCAGAAGTCGACACAAAACGCAATAAATGGTATAACAAATAAAGTTAACACCGTAA TCGAGAAGATGAACATCCAGTTTACAGCCGTGGGGAAGGAATTCAACAAGCTGGAAAAGCGCATGGAGAACCTGAACAAAAAGGT GGATGACGGCTTCCTGGACATTTGGACCTACAACGCAGAGTTGTTAGTGCTGCTGGAGAATGAGCGAACACTGGATTTTCATGAC TCCAATGTGAAGAATCTCTACGAGAAGGTGAAGTCACAGCTCAAGAACAATGCTAAAGAGATTGGGAACGGGTGCTTCGAATTTTA CCATAAATGCGACAACGAGTGTATGGAGTCCGTCCGTAATGGCACTTATGACTACCCTAAATACTCTGAAGAGTCCAAACTCAATC GGGAGAAAGTGGACGGTGTGAAGCTGGAGTCAATGGGGATATACCAGATCTTAGCAATTTACAGCACCGTCGCCTCCTCATTGGT GCTTCTGGTCTCTCTCGGAGCAATTAGTTTTTGGATGTGTTCCAATGGGTCTCTCCAGTGCCGGATATGCATT

2676	ATGAAAGCCAACCTTCTAGTGTTACTTTGCGCTCTCGCCGCTGCCGATGCCGATACAATCTGTATCGGTTATCACGCGAATAATTC CACCGACACTGTCGATACAGTGTTGGAAAAGAATGTCACAGTCACACATTCCGTCAATTTGTTGGAGGACAGCCATAACGGAAAGT TGTGTAGATTGAAAGGGATTGCCCCACTGCAGCTGGGTAAGTGTAATATTGCTGGTTGGCTACTGGGCAACCCAGAATGTGACCC CCTTCTACCCGTCAGGAGTTGGAGCTACATTGTGGAGACCCCTAACAGCGAGAATGGCATTTGCTACCCAGGCGACTTTATCGATT ACGAAGAACTGCGGGAACAGTTGAGCTCTGTTAGTAGCTTCGAGAGATTCGAGATATTCCCAAAGGAGTCAAGCTGGCCCAATCA CAACACCAATGGGGTTACAGCAGCTTGTAGCCATGAAGGCAAAAGCAGCTTCTATCGAAATCTACTCTGGCTCACCGAAAAGGAA GGATCCTACCCGAAACTCAAAAACAGCTACGTCAACAAGAAGGGAAAAGAGGTGCTGGTGCTGTGGGGAATCCATCATCCTCCCA ACAGCAAAGAGCAACAGAATTTGTATCAAAACGAAAACGCTTACGTGAGCGTGGTAACCTCAAACTATAACAGACGGTTTACACCA GAAATCGCAGAGAGGCCAAAAGTGCGGGACCAAGCCGGTAGAATGAACTATTACTGGACCCTCTTGAAACCTGGAGACACCATAA TATTCGAGGCCAACGGAAATCTGATCGCCCCAATGTATGCCTTCGCTCTCTCTAGGGGCTTCGGCAGCGGGATAATTACCTCCAA CGCCAGCATGCATGAGTGCAATACCAAGTGTCAAACTCCCCTGGGCGCAATTAACAGCTCCCTGCCTTACCAAAACATTCACCCA GTCACCATCGGAGAGTGCCCCAAATACGTTAGATCTGCAAAGCTGCGGATGGTGACAGGTCTACGCAACAACCCCTCAATCCAGT CACGGGGGCTTTTCGGTGCCATTGCCGGGTTTATTGAAGGTGGCTGGACCGGCATGATAGATGGGTGGTATGGCTACCACCATC AGAATGAACAGGGAAGTGGGTATGCAGCGGACCAGAAAAGCACGCAGAACGCAATAAACGGGATCACGAATAAAGTGAACACCG TCATTGAAAAGATGAACATCCAGTTTACCGCGGTTGGAAAAGAATTTAACAAGCTCGAAAAAAGAATGGAAAATCTGAACAAGAAG GTCGACGACGGATTCCTTGATATTTGGACTTACAATGCTGAGCTTCTTGTACTGTTAGAAAACGAGCGAACCTTGGATTTCCACGAT TCAAACGTCAAAAACTTGTACGAGAAAGTGAAGAGCCAACTCAAGAACAACGCCAAGGAAATTGGGAATGGTTGCTTTGAATTTTA CCATAAGTGCGACAATGAATGCATGGAGTCGGTACGAAACGGGACTTATGATTATCCAAAGTATAGCGAAGAGAGTAAACTCAACC GGGAGAAAGTCGATGGGGTGAAACTGGAGTCAATGGGAATTTATCAAATTCTGGCTATCTATTCTACAGTGGCCTCTTCTCTCGTC TTGCTAGTGAGCCTCGGAGCAATCTCTTTTTGGATGTGCAGCAATGGCTCCCTGCAATGCCGCATCTGCATA
2677	ATGAAGGCTAACCTCCTGGTGTTGTTGTGTGCCCTGGCTGCAGCTGACGCCGATACTATTTGCATAGGCTATCATGCTAATAATAG TACAGATACGGTTGACACCGTTCTCGAAAAGAACGTGACGGTGACACATAGCGTTAATCTGCTAGAAGACTCCCACAATGGGAAG CTGTGCCGGCTGAAGGGGATCGCACCACTCCAGCTAGGTAAGTGCAATATCGCCGGGTGGCTGTTAGGAAATCCGGAGTGCGAC CCTCTTCTCCCCGTGAGGTCTTGGTCCTACATCGTGGAAACTCCAAACAGTGAAAACGGAATCTGCTATCCTGGCGATTTTATAGA CTATGAAGAGCTGCGTGAGCAACTGAGCAGCGTATCCAGCTTCGAAAGATTCGAGATTTTCCCTAAGGAATCGTCATGGCCGAAT CACAACACTAATGGCGTCACTGCTGCATGCAGCCATGAGGGCAAAAGCAGCTTCTATAGAAATCTCCTGTGGCTTACCGAGAAAG AGGGTAGCTATCCTAAGCTCAAGAATAGCTATGTCAATAAGAAAGGCAAGGAGGTGCTGGTGTTGTGGGGAATCCATCACCCACC TAACTCCAAAGAGCAGCAGAACCTGTATCAAAACGAGAACGCCTACGTGTCTGTAGTGACGAGCAACTACAACAGGCGTTTCACA CCTGAGATCGCGGAGCGCCCCAAGGTTCGCGACCAAGCGGGGCGCATGAATTACTATTGGACTCTTTTAAAGCCGGGAGATACG ATCATCTTTGAGGCTAACGGCAACTTAATCGCCCCTATGTACGCCTTTGCGTTGTCGCGGGGATTCGGATCCGGTATAATAACAAG CAACGCAAGTATGCACGAATGTAATACTAAATGTCAAACCCCTTTAGGGGCCATTAACAGCAGCCTTCCGTACCAGAACATTCATC CCGTCACCATCGGCGAATGCCCAAAATATGTGCGGTCCGCGAAACTCAGAATGGTAACAGGCCTGCGGAATAATCCTAGTATACA GTCCAGAGGACTTTTTGGGGCGATAGCTGGATTCATTGAAGGCGGATGGACTGGGATGATTGACGGATGGTACGGCTACCACCAT CAGAACGAACAGGGCAGTGGGTACGCTGCTGATCAAAAATCCACACAAAATGCCATTAATGGAATTACAAACAAAGTTAATACAGT CATCGAGAAAATGAATATTCAGTTCACAGCTGTGGGCAAAGAGTTTAACAAACTGGAGAAACGAATGGAGAACCTGAACAAAAAGG TTGATGATGGCTTTCTGGACATCTGGACATATAATGCTGAGCTGTTGGTCCTACTGGAGAATGAGAGGACCTTGGACTTCCACGAC TCCAATGTTAAGAACCTATATGAGAAGGTTAAGTCGCAATTGAAAAACAATGCAAAGGAAATCGGGAACGGGTGCTTTGAATTCTAT CATAAATGTGATAATGAGTGCATGGAAAGCGTGCGTAATGGGACTTACGATTATCCTAAGTATAGTGAGGAATCTAAGTTGAACAG AGAGAAAGTGGATGGAGTCAAGTTAGAAAGTATGGGTATCTACCAAATTCTTGCTATATATAGCACAGTGGCTTCCAGCCTCGTCT TGCTTGTCTCTCTCGGGGCTATCAGCTTTTGGATGTGCTCAAACGGCAGCCTCCAGTGTCGGATATGTATC

2678	ATGAAGGCTAACCTGCTGGTGCTGTTGTGTGCACTCGCAGCCGCGGACGCCGACACTATCTGCATAGGCTATCACGCTAATAACT CAACGGATACGGTGGATACCGTTCTGGAGAAGAATGTAACCGTCACACACAGCGTCAATCTGCTCGAGGATAGCCACAACGGCAA ACTATGCCGCCTGAAGGGGATAGCTCCCCTCCAGCTGGGAAAGTGTAACATTGCCGGATGGCTCCTGGGAAATCCAGAGTGCGA TCCCCTGTTACCTGTCAGATCATGGTCTTACATCGTGGAGACCCCTAACTCCGAGAATGGGATCTGCTACCCCGGAGACTTTATTG ACTACGAGGAGCTGAGAGAGCAATTGTCTTCAGTGAGTTCTTTTGAGAGGTTTGAAATCTTCCCGAAGGAGAGTTCATGGCCAAAT CACAATACGAATGGAGTTACCGCAGCTTGCTCTCATGAGGGCAAGAGCAGTTTCTACAGGAACCTACTGTGGCTGACCGAGAAGG AAGGTAGCTACCCGAAGCTGAAGAATTCATACGTCAATAAAAAGGGCAAGGAAGTGTTAGTGCTCTGGGGCATCCATCACCCCCC CAACTCCAAAGAGCAGCAGAACCTTTATCAGAATGAAAACGCTTATGTTTCAGTGGTGACCTCTAATTATAACCGACGTTTTACTCC AGAGATCGCTGAAAGACCCAAGGTCCGTGACCAAGCAGGGCGGATGAACTATTACTGGACCCTGCTAAAGCCTGGCGATACTATA ATTTTCGAAGCTAACGGCAACCTTATTGCTCCCATGTATGCATTCGCCCTGAGTCGTGGATTTGGCAGCGGTATTATCACCTCTAAT GCATCGATGCATGAATGTAATACAAAATGCCAGACTCCATTGGGCGCTATTAACTCAAGCCTTCCTTATCAAAATATCCACCCTGTA ACTATCGGGGAGTGTCCCAAATATGTGCGCTCAGCCAAGCTCCGCATGGTAACTGGACTCCGCAACAATCCATCAATACAGAGCA GAGGACTGTTCGGGGCGATAGCCGGGTTCATCGAAGGAGGCTGGACCGGCATGATCGACGGGTGGTACGGTTACCATCACCAAA ACGAGCAGGGCAGCGGCTACGCTGCCGATCAAAAGTCCACACAAAATGCCATAAACGGAATAACCAATAAGGTCAACACTGTGAT CGAAAAGATGAATATTCAGTTCACTGCTGTCGGTAAAGAATTCAATAAGCTGGAAAAGCGCATGGAGAACCTTAACAAAAAGGTGG ACGACGGCTTTTTAGACATCTGGACCTACAATGCCGAACTGCTAGTGCTACTTGAAAACGAAAGGACCTTGGATTTTCACGACAGT AATGTTAAAAATCTCTACGAGAAAGTAAAGTCTCAGTTGAAGAACAATGCTAAGGAGATCGGCAATGGATGCTTTGAGTTCTATCAC AAGTGCGATAACGAATGCATGGAAAGTGTGCGCAACGGAACCTATGACTACCCCAAGTATTCCGAAGAGAGCAAGCTTAACAGAG AGAAGGTCGACGGAGTGAAGCTGGAAAGTATGGGGATCTACCAAATACTGGCAATATACAGCACAGTTGCAAGCAGTCTCGTGTT ACTCGTCTCCCTTGGAGCAATCTCTTTCTGGATGTGTTCTAACGGCAGCCTGCAGTGCAGAATTTGTATC
2679	ATGAAGGCAAACTTGCTCGTGCTGTTGTGCGCATTAGCCGCCGCCGATGCCGACACAATCTGCATCGGCTATCACGCTAACAACT CCACTGACACTGTGGATACCGTACTTGAAAAAAATGTGACCGTCACACACTCGGTTAATCTCTTAGAGGACTCGCACAATGGCAAA CTGTGTCGTCTGAAAGGGATCGCTCCACTTCAACTCGGGAAGTGTAATATTGCCGGGTGGCTGCTGGGCAACCCCGAGTGCGAC CCTCTTCTGCCTGTACGGTCGTGGTCTTATATAGTGGAAACACCCAATAGTGAGAACGGCATATGCTATCCCGGAGACTTCATTGA TTATGAAGAACTTAGGGAACAGCTAAGCTCCGTGTCAAGCTTTGAAAGGTTTGAAATATTCCCCAAGGAGAGCTCCTGGCCTAATC ACAATACCAACGGTGTGACCGCCGCCTGTTCGCACGAGGGCAAGAGCTCATTTTACAGGAATCTACTGTGGCTCACCGAGAAGGA AGGTTCGTACCCTAAGCTCAAGAATAGCTACGTGAATAAAAAGGGCAAAGAGGTACTGGTGCTGTGGGGGATTCACCACCCCCCT AATTCGAAGGAGCAGCAAAACTTGTACCAGAATGAGAATGCCTACGTCTCTGTTGTCACTAGTAATTACAACCGTAGATTCACGCC TGAAATTGCTGAGCGACCAAAAGTTCGAGATCAGGCTGGCAGGATGAATTATTATTGGACCCTCCTGAAGCCCGGAGATACCATTA TCTTCGAGGCTAATGGTAACCTGATCGCTCCTATGTACGCCTTCGCTTTGAGTCGAGGCTTCGGCAGCGGGATAATTACGTCTAAC GCTTCTATGCACGAGTGCAATACAAAGTGTCAGACTCCCCTTGGGGCAATCAATTCAAGCCTTCCCTATCAGAATATCCACCCTGT TACTATAGGGGAGTGTCCGAAATACGTTAGGTCGGCAAAGCTGAGGATGGTCACTGGCCTGCGAAATAACCCGTCCATTCAGTCC AGGGGGCTTTTCGGTGCTATCGCTGGATTTATTGAAGGTGGTTGGACTGGCATGATCGACGGATGGTACGGGTATCACCACCAGA ACGAACAGGGATCTGGGTACGCTGCCGATCAAAAATCGACCCAGAACGCCATAAACGGCATTACAAATAAGGTGAATACCGTCAT CGAAAAAATGAATATTCAATTCACCGCCGTAGGCAAAGAGTTTAATAAACTTGAGAAAAGGATGGAGAACCTCAATAAAAAAGTCGA CGATGGGTTTCTGGATATCTGGACCTATAACGCTGAGCTTCTTGTGCTGCTCGAAAATGAACGCACACTGGATTTCCACGATTCCA ACGTGAAAAATTTGTACGAAAAGGTGAAGTCACAACTGAAGAATAACGCAAAGGAAATCGGGAATGGCTGTTTTGAATTCTATCAC AAGTGTGACAACGAATGCATGGAGAGTGTCCGAAACGGAACCTACGATTATCCAAAGTACTCGGAAGAATCTAAGCTGAATCGGG AGAAGGTGGACGGGGTCAAGCTGGAGTCAATGGGTATTTATCAGATCCTAGCGATTTACTCAACCGTGGCATCGAGCCTGGTCCT ACTTGTTAGTCTTGGGGCAATAAGCTTCTGGATGTGCAGCAACGGCAGCTTGCAGTGCAGAATATGTATA

2680	ATGAAGGCTAACCTGCTCGTTCTCTTATGTGCACTGGCCGCCGCAGACGCGGATACTATTTGTATAGGCTACCACGCGAATAACTC TACTGATACGGTGGACACAGTTCTCGAGAAAAACGTGACAGTGACTCACTCCGTGAATCTGCTGGAGGATTCACACAACGGCAAA CTGTGTCGACTGAAAGGAATCGCACCGTTGCAACTTGGTAAGTGCAATATTGCCGGATGGCTCCTGGGAAACCCCGAATGCGATC CTTTGCTTCCCGTGCGGAGTTGGTCCTACATCGTGGAGACCCCTAACTCTGAGAACGGAATCTGTTATCCCGGGGATTTCATTGAT TATGAAGAGCTCAGAGAACAATTGAGTTCTGTTTCTAGCTTTGAGAGATTCGAGATTTTCCCTAAGGAGTCGTCCTGGCCAAATCAC AACACTAACGGAGTAACCGCCGCCTGTTCTCACGAGGGCAAGTCAAGCTTTTACAGGAATCTCCTGTGGTTAACAGAAAAGGAAG GGAGTTACCCAAAACTAAAAAACTCGTACGTGAACAAGAAAGGCAAGGAGGTCCTGGTGCTGTGGGGAATTCATCACCCCCCGAA TTCCAAGGAACAACAGAATCTGTATCAAAACGAAAACGCTTATGTGTCCGTCGTGACATCGAACTACAATAGAAGATTTACCCCCG AGATTGCAGAGCGCCCTAAAGTCAGGGATCAGGCTGGGAGAATGAACTATTATTGGACACTGCTAAAGCCCGGCGATACAATTAT CTTTGAGGCCAATGGCAATCTGATTGCTCCCATGTATGCATTTGCCCTGAGCAGAGGGTTTGGCAGTGGAATCATTACCAGCAATG CTAGCATGCATGAATGTAATACAAAGTGTCAGACACCATTAGGAGCCATAAACAGTTCACTCCCTTATCAGAACATCCATCCCGTGA CAATCGGAGAGTGTCCTAAATACGTCAGAAGTGCAAAATTGAGAATGGTTACCGGCTTGAGGAATAATCCATCTATTCAATCCCGC GGGCTCTTTGGAGCCATCGCTGGCTTTATTGAAGGAGGATGGACCGGAATGATTGATGGATGGTATGGTTACCACCATCAGAATG AGCAGGGCTCCGGCTACGCCGCCGACCAGAAGTCTACCCAAAACGCCATAAACGGAATCACCAATAAGGTGAACACTGTCATTGA AAAGATGAATATACAATTCACTGCCGTCGGGAAGGAGTTTAACAAACTCGAAAAGCGGATGGAGAATTTGAATAAAAAGGTCGACG ACGGCTTCCTGGATATCTGGACATACAATGCCGAGCTCCTGGTCCTGTTGGAGAATGAACGCACACTCGATTTTCACGACAGCAAT GTGAAGAATCTTTACGAAAAGGTTAAAAGTCAGTTGAAAAACAACGCTAAAGAAATAGGTAATGGGTGTTTCGAGTTTTACCACAAG TGCGACAATGAGTGCATGGAATCTGTCAGAAATGGGACCTACGACTACCCTAAGTACTCCGAAGAGTCTAAGCTGAATAGGGAGA AGGTGGATGGGGTGAAGCTGGAATCCATGGGCATCTACCAGATCCTCGCTATCTACAGCACAGTGGCCTCTAGCTTGGTGCTTTT GGTGTCACTTGGGGCGATTTCGTTCTGGATGTGCTCCAACGGAAGCCTGCAATGCCGAATCTGTATC
2681	ATGAAGGCTAACCTTCTTGTGCTTCTCTGTGCACTTGCTGCCGCAGATGCTGATACAATATGCATCGGTTACCACGCTAATAATTCT ACCGATACGGTTGACACGGTTCTTGAAAAGAACGTCACAGTGACTCACAGCGTCAACCTCCTGGAGGACTCTCATAACGGAAAATT GTGCCGCCTCAAAGGCATTGCCCCATTGCAATTAGGCAAGTGTAATATTGCAGGTTGGCTCCTGGGTAACCCGGAATGCGACCCT CTGTTGCCTGTCAGGTCTTGGAGTTATATAGTCGAAACCCCAAATTCCGAAAATGGAATATGTTACCCTGGCGACTTTATCGACTAC GAAGAACTGAGGGAGCAGCTTAGTTCCGTCAGTTCTTTCGAACGTTTCGAAATCTTCCCCAAGGAATCTAGTTGGCCCAACCATAA CACCAACGGTGTGACGGCGGCATGCAGTCACGAGGGCAAGTCCAGCTTCTATAGAAACTTGCTGTGGTTAACCGAGAAAGAGGG AAGTTACCCTAAGCTGAAGAACAGCTACGTAAACAAAAAAGGCAAAGAAGTCCTGGTTCTCTGGGGCATTCACCACCCACCCAATT CCAAGGAACAGCAAAATCTCTATCAAAATGAAAATGCTTATGTCTCTGTTGTGACCTCGAACTATAATCGGCGTTTTACCCCTGAGA TCGCCGAGAGGCCTAAGGTCAGGGACCAAGCTGGCCGAATGAATTATTACTGGACCCTACTGAAGCCTGGCGATACCATAATCTT CGAGGCTAATGGAAATCTGATTGCGCCAATGTATGCTTTTGCACTGAGCCGCGGGTTTGGGAGCGGAATTATCACCTCTAACGCA TCTATGCACGAGTGTAACACAAAGTGTCAGACCCCGCTGGGGGCTATTAACTCCAGTTTGCCATACCAAAATATTCATCCGGTCAC AATCGGTGAATGCCCCAAGTATGTTAGAAGCGCTAAGTTAAGAATGGTGACCGGGCTGCGTAACAACCCTAGTATCCAGTCACGC GGCCTCTTTGGGGCTATTGCCGGGTTCATCGAGGGAGGTTGGACTGGGATGATCGACGGATGGTATGGGTACCACCATCAAAAC GAGCAGGGATCAGGCTACGCCGCAGACCAGAAGAGCACTCAGAACGCCATCAACGGGATCACTAATAAAGTTAATACCGTGATTG AAAAAATGAACATACAATTTACTGCAGTGGGCAAGGAGTTCAACAAACTAGAGAAACGCATGGAGAACTTGAACAAAAAAGTTGAC GACGGATTTCTGGACATCTGGACATATAATGCAGAGCTGCTGGTCCTGCTAGAGAACGAACGGACCCTAGACTTCCATGACTCAA ACGTCAAGAATCTGTACGAGAAGGTGAAATCTCAGCTCAAAAATAATGCCAAAGAGATCGGCAATGGATGCTTTGAATTCTATCAC AAGTGTGACAATGAGTGCATGGAGAGCGTGCGAAATGGAACTTACGACTACCCAAAGTACAGCGAGGAGAGTAAGCTTAACAGGG AGAAGGTGGACGGCGTGAAGCTCGAAAGCATGGGGATATACCAGATCCTCGCCATATACTCTACTGTGGCGTCGTCTCTCGTGCT TTTGGTATCACTCGGCGCTATCAGCTTCTGGATGTGTTCCAATGGCAGCTTGCAGTGCCGGATATGCATT

2682	ATGAAAGCCAATCTCTTAGTGCTACTGTGCGCCCTTGCCGCCGCAGACGCAGATACGATTTGCATTGGGTATCACGCAAACAATTC CACCGACACAGTGGATACCGTGCTCGAAAAGAACGTAACTGTGACTCACTCTGTGAATTTACTCGAGGACAGTCACAACGGGAAG CTCTGCAGACTTAAGGGGATCGCACCCTTGCAACTCGGAAAGTGTAACATTGCCGGGTGGCTTCTCGGCAACCCCGAGTGTGACC CTCTTCTTCCAGTTCGGAGCTGGAGTTATATAGTGGAAACGCCAAATTCCGAGAACGGCATATGTTACCCAGGAGACTTTATCGAT TATGAAGAACTCCGCGAACAGCTCAGCAGCGTGAGTTCTTTCGAACGTTTCGAGATCTTCCCAAAAGAGTCCAGTTGGCCAAATCA TAACACCAATGGCGTAACAGCAGCGTGCAGCCACGAGGGAAAATCTAGCTTTTACAGGAACCTACTTTGGCTCACAGAAAAGGAG GGATCCTATCCTAAACTGAAGAATTCATACGTCAATAAAAAAGGTAAAGAGGTGTTGGTGCTGTGGGGGATTCATCACCCACCAAA CTCCAAAGAACAGCAGAACCTGTACCAAAACGAGAACGCCTACGTCTCCGTGGTGACGAGCAACTATAACCGACGCTTTACCCCC GAAATCGCGGAGCGTCCAAAGGTGCGGGACCAGGCAGGGCGTATGAACTATTATTGGACACTTCTAAAACCCGGTGACACCATTA TCTTTGAAGCGAACGGCAACTTGATTGCGCCAATGTATGCTTTCGCTCTGAGTCGGGGATTTGGATCGGGGATTATTACCTCCAAC GCGTCTATGCATGAATGTAACACCAAATGTCAAACCCCTCTTGGGGCCATCAACAGCTCCCTGCCATATCAGAACATTCACCCAGT CACCATCGGAGAATGCCCCAAGTATGTCCGCAGTGCTAAGCTCCGCATGGTAACCGGCCTGAGAAACAATCCCAGTATTCAGTCT AGGGGCCTCTTCGGCGCTATTGCAGGATTCATTGAGGGCGGATGGACAGGCATGATCGATGGCTGGTACGGTTATCATCACCAG AACGAGCAAGGATCCGGCTACGCTGCGGACCAGAAGAGCACCCAGAACGCTATTAACGGCATTACGAATAAGGTGAACACTGTCA TCGAGAAGATGAACATTCAGTTTACCGCGGTTGGGAAGGAGTTCAACAAACTTGAAAAACGAATGGAGAATCTAAACAAAAAAGTC GACGATGGATTCCTGGATATCTGGACCTACAACGCTGAGCTGCTGGTGCTGCTGGAAAATGAGCGGACATTGGACTTCCACGATT CCAACGTGAAGAATCTTTACGAGAAAGTTAAATCACAGTTGAAGAATAATGCCAAAGAGATTGGGAACGGCTGTTTTGAATTCTACC ACAAATGCGATAACGAATGTATGGAATCTGTACGGAATGGTACGTATGACTATCCAAAATACTCCGAAGAGTCCAAGCTGAACCGA GAAAAAGTGGACGGGGTCAAACTGGAAAGCATGGGGATATACCAGATCCTGGCCATATATAGCACTGTGGCTTCTTCTTTGGTTCT CCTTGTGTCACTCGGGGCCATAAGCTTCTGGATGTGTTCAAACGGAAGCTTGCAGTGCAGAATTTGCATC
2683	ATGAAGGCCAACTTACTGGTGTTGCTATGCGCTCTGGCCGCCGCTGACGCCGATACCATCTGTATTGGATACCACGCCAACAACA GCACCGACACTGTGGACACAGTGCTTGAGAAGAATGTCACCGTCACACATAGTGTAAACTTGCTCGAAGATAGTCACAACGGGAA GCTGTGCCGCCTTAAAGGTATTGCACCACTGCAGTTAGGCAAGTGCAATATTGCCGGGTGGCTGCTTGGGAACCCCGAGTGTGAT CCTCTATTGCCAGTTCGGTCGTGGAGCTATATCGTGGAGACACCGAACAGCGAAAACGGCATCTGTTATCCTGGAGATTTTATCGA CTATGAGGAACTCAGGGAACAGCTCTCGTCTGTGTCTAGCTTCGAACGCTTTGAAATTTTCCCAAAAGAGAGTTCCTGGCCAAATC ATAACACTAACGGTGTGACTGCCGCTTGCTCCCATGAAGGCAAATCAAGCTTTTATAGGAATCTGTTGTGGTTGACTGAAAAAGAA GGCTCCTACCCGAAACTCAAGAACAGCTATGTGAACAAGAAAGGTAAGGAAGTGCTGGTGCTGTGGGGCATCCATCATCCTCCTA ACTCAAAGGAGCAGCAAAATCTTTACCAGAATGAAAACGCATACGTGTCCGTTGTAACCTCTAATTATAACCGCCGGTTCACCCCA GAAATCGCCGAGCGTCCAAAGGTTAGAGATCAAGCAGGGCGCATGAATTATTACTGGACACTGCTCAAGCCCGGCGATACCATCA TCTTTGAGGCTAACGGCAACCTCATTGCTCCGATGTATGCGTTCGCTCTGAGTCGCGGCTTTGGCTCAGGCATTATTACTTCCAAT GCCTCTATGCACGAGTGCAACACTAAATGCCAGACGCCCCTGGGTGCAATCAACTCTAGCCTCCCATATCAGAACATACATCCCGT GACCATCGGCGAATGCCCTAAATACGTCCGGTCCGCAAAGCTGCGAATGGTGACGGGCCTGAGAAATAACCCCTCAATACAGTCG CGAGGGCTGTTCGGCGCCATTGCCGGGTTTATTGAGGGGGGATGGACTGGGATGATAGATGGTTGGTATGGGTATCATCATCAG AACGAGCAAGGGTCTGGGTATGCCGCTGACCAAAAAAGCACCCAGAACGCCATAAACGGGATTACCAATAAGGTTAACACAGTTA TTGAGAAGATGAATATTCAGTTTACCGCTGTCGGGAAAGAATTCAATAAATTGGAAAAAAGGATGGAGAACCTGAACAAAAAGGTG GACGATGGGTTTCTCGATATCTGGACATATAACGCTGAGCTTCTCGTACTGCTGGAAAATGAAAGGACGCTCGATTTCCATGATTC CAATGTCAAGAATCTCTACGAGAAGGTCAAGTCTCAGCTGAAGAACAATGCCAAAGAGATCGGGAATGGATGTTTTGAGTTCTATC ACAAGTGCGATAACGAATGCATGGAGTCCGTCAGGAACGGCACCTATGATTACCCTAAGTACAGTGAAGAAAGCAAACTTAATAGG GAAAAGGTGGATGGTGTGAAGCTGGAGTCAATGGGCATTTACCAGATTCTGGCCATTTACAGTACAGTAGCCTCTAGTCTGGTGCT GCTTGTCAGCCTTGGCGCCATCTCCTTTTGGATGTGTAGCAACGGATCACTGCAGTGCAGGATCTGCATT

2684	ATGAAAGCAAACTTGCTTGTGCTGTTATGCGCTCTGGCAGCTGCGGACGCCGATACTATCTGCATCGGTTACCACGCTAACAATTC AACAGATACCGTTGATACTGTGCTGGAGAAGAACGTGACCGTCACCCACAGTGTGAACCTCTTGGAAGACAGCCACAACGGAAAA CTGTGCCGCCTGAAGGGCATCGCGCCGTTACAGCTGGGAAAGTGCAATATCGCCGGATGGCTTTTGGGTAATCCAGAATGTGATC CATTGCTCCCCGTGAGAAGCTGGAGCTATATTGTTGAAACCCCTAACTCAGAGAATGGGATCTGCTACCCCGGGGACTTCATTGA CTACGAGGAACTGCGCGAGCAGTTGTCATCCGTCAGTTCATTCGAGCGGTTTGAAATATTTCCCAAGGAGTCAAGTTGGCCAAAC CACAACACGAACGGAGTAACAGCGGCTTGTAGCCATGAAGGTAAAAGCTCTTTTTACAGGAATCTGCTGTGGTTGACTGAGAAGG AGGGCTCCTACCCCAAACTGAAAAACAGCTACGTGAACAAAAAGGGCAAGGAGGTCCTGGTGCTCTGGGGAATTCACCATCCGCC CAATTCCAAAGAGCAGCAGAATTTATACCAAAATGAGAACGCCTACGTTTCTGTCGTCACGTCAAATTATAACCGACGCTTTACACC TGAAATAGCTGAGAGACCCAAGGTGAGAGATCAGGCTGGGAGGATGAATTACTATTGGACTCTGCTAAAACCTGGAGACACTATA ATCTTCGAAGCCAACGGGAACCTGATTGCCCCTATGTACGCATTTGCTTTATCCCGAGGGTTTGGAAGCGGGATCATTACTTCAAA TGCTTCTATGCACGAGTGCAATACCAAATGCCAGACACCTTTAGGCGCTATCAATTCATCTCTGCCTTATCAGAACATTCACCCAGT GACAATAGGAGAGTGTCCTAAGTATGTCCGCTCCGCGAAGCTTCGTATGGTCACAGGCCTCAGGAATAACCCAAGTATCCAGTCG AGAGGCCTTTTTGGTGCCATCGCTGGCTTTATCGAAGGAGGGTGGACCGGCATGATTGATGGGTGGTATGGATACCATCATCAGA ACGAGCAAGGCTCAGGTTACGCCGCGGACCAGAAGTCCACGCAAAATGCCATTAACGGCATCACGAACAAAGTCAACACCGTTAT CGAGAAGATGAATATACAGTTCACAGCCGTAGGCAAAGAATTTAATAAACTCGAAAAGCGGATGGAGAATTTAAACAAAAAAGTCG ATGATGGATTTCTCGATATCTGGACTTATAACGCCGAACTTCTAGTGCTGCTTGAGAACGAGAGAACGCTTGATTTTCACGACAGTA ACGTTAAAAACCTGTACGAAAAGGTAAAGTCCCAACTGAAAAACAATGCCAAAGAAATTGGAAACGGTTGCTTTGAATTTTACCACA AGTGCGATAATGAGTGCATGGAAAGTGTAAGAAATGGCACGTATGACTATCCCAAATACTCCGAAGAAAGCAAGCTCAATCGTGAG AAGGTGGATGGTGTGAAGCTGGAATCGATGGGGATTTATCAGATCCTAGCTATCTATTCAACAGTCGCCTCTAGTCTTGTTCTTCTT GTATCTCTTGGAGCCATCTCATTTTGGATGTGTTCAAACGGATCCCTGCAGTGTCGGATCTGTATC
2685	ATGAAGGCTAACCTGCTGGTTCTACTCTGCGCACTCGCAGCTGCGGACGCCGACACAATTTGTATTGGATATCACGCTAACAATAG CACCGATACCGTAGACACCGTACTGGAAAAGAATGTGACGGTCACACACTCAGTAAATCTCCTGGAAGACAGCCATAATGGCAAG CTGTGTAGATTAAAAGGGATTGCTCCTCTCCAGTTGGGCAAATGTAACATCGCCGGTTGGCTGCTGGGGAACCCCGAGTGCGACC CATTGCTGCCCGTAAGGAGCTGGTCCTATATAGTTGAGACACCCAATTCTGAGAATGGCATCTGCTACCCCGGAGATTTTATCGAT TATGAGGAGTTGCGTGAGCAGCTCTCTAGCGTGAGCTCGTTTGAAAGATTTGAAATTTTTCCGAAAGAATCTAGTTGGCCCAACCA CAATACGAACGGAGTGACTGCCGCCTGCTCCCACGAGGGAAAATCCTCATTTTACAGGAACTTACTCTGGCTTACTGAGAAGGAA GGTAGCTACCCAAAGCTCAAGAACAGCTACGTCAACAAGAAGGGCAAGGAAGTCCTGGTCCTGTGGGGAATCCATCATCCGCCAA ACTCCAAGGAGCAACAGAATCTATATCAGAATGAGAATGCCTACGTTTCAGTGGTTACGTCTAATTATAACCGCAGGTTTACCCCC GAGATCGCTGAGCGCCCCAAAGTTAGAGACCAGGCAGGGCGCATGAACTACTATTGGACCCTCCTGAAACCTGGCGACACAATTA TCTTCGAGGCAAACGGTAATCTCATCGCGCCTATGTACGCTTTTGCCCTTAGCCGGGGTTTTGGATCAGGCATCATAACTTCTAAC GCCTCTATGCACGAATGTAACACCAAGTGTCAGACCCCCCTAGGAGCTATCAACAGCTCGCTTCCTTACCAGAACATTCATCCCGT GACCATAGGTGAGTGTCCAAAATACGTCAGGAGTGCAAAACTAAGGATGGTTACCGGACTCCGGAATAACCCTTCGATTCAGTCG CGCGGGCTCTTTGGAGCAATCGCCGGGTTCATCGAAGGCGGGTGGACCGGGATGATTGACGGTTGGTACGGCTACCATCACCAA AACGAACAAGGCTCAGGTTATGCAGCTGACCAGAAGTCCACTCAGAATGCCATAAATGGCATTACCAACAAAGTTAATACAGTCAT TGAGAAGATGAATATACAGTTCACCGCGGTGGGCAAAGAGTTCAACAAACTGGAGAAACGTATGGAAAACTTGAACAAAAAAGTG GACGATGGGTTCTTGGACATCTGGACGTATAATGCCGAGTTATTAGTGCTGCTGGAAAATGAAAGAACGTTAGACTTCCACGACTC TAATGTTAAAAATTTGTACGAAAAGGTCAAATCCCAACTGAAAAATAACGCCAAGGAAATAGGCAATGGCTGTTTTGAATTCTATCAT AAGTGCGATAACGAGTGTATGGAGAGTGTTCGCAATGGTACATACGACTACCCTAAATACTCCGAGGAAAGTAAGCTGAACAGGG AAAAGGTCGATGGAGTCAAGCTGGAATCCATGGGGATTTATCAGATCCTGGCCATTTATAGCACTGTAGCCTCCTCCCTGGTGCTG CTGGTATCACTCGGCGCAATAAGCTTCTGGATGTGTTCTAACGGGAGCCTGCAATGCAGAATATGCATT

2686	ATGAAAGCAAACCTCCTCGTGTTGCTCTGTGCCCTGGCTGCAGCCGACGCCGACACCATATGTATCGGGTACCACGCCAATAACT CCACTGATACCGTCGATACCGTCCTTGAGAAAAATGTGACAGTCACACACAGCGTTAATTTATTGGAGGACAGCCACAACGGCAAG CTCTGTAGGCTGAAAGGGATCGCCCCCTTACAGCTGGGGAAATGTAATATCGCTGGATGGCTGCTGGGAAACCCCGAGTGCGAC CCCTTGCTGCCTGTGAGGTCTTGGAGCTACATTGTTGAGACCCCTAACAGCGAGAACGGCATTTGTTACCCTGGCGACTTCATTGA CTACGAGGAGCTGAGGGAACAATTGTCATCTGTCAGCTCATTTGAGCGATTCGAGATCTTTCCCAAGGAATCGTCATGGCCAAATC ACAACACAAACGGGGTGACCGCGGCTTGCTCACATGAAGGAAAGAGCTCCTTTTACCGAAATCTTCTGTGGCTGACGGAGAAGGA AGGATCATACCCAAAATTGAAAAACTCCTATGTGAACAAGAAAGGTAAAGAGGTCCTTGTTTTATGGGGCATTCACCACCCACCTAA CTCAAAAGAACAGCAAAACCTGTACCAGAACGAGAATGCATACGTTTCAGTAGTCACCTCTAACTATAATAGGAGATTCACCCCCG AAATCGCCGAGAGGCCAAAGGTACGGGATCAGGCTGGCAGGATGAATTACTATTGGACCCTACTTAAACCTGGAGACACGATCAT CTTCGAGGCAAATGGAAACCTCATCGCCCCTATGTATGCCTTTGCCCTTAGCCGAGGGTTTGGATCAGGGATAATAACCTCTAACG CCAGTATGCACGAATGCAATACGAAGTGTCAAACCCCTCTGGGAGCAATAAATTCTTCCCTTCCTTATCAGAACATTCACCCCGTC ACAATTGGAGAGTGCCCCAAATACGTCAGGTCCGCCAAACTGAGAATGGTTACCGGCTTGCGGAATAATCCGAGCATTCAGAGCC GCGGATTATTTGGTGCTATTGCTGGCTTTATCGAAGGAGGGTGGACCGGAATGATCGACGGGTGGTACGGCTACCATCATCAGAA CGAGCAGGGTAGCGGTTACGCTGCCGATCAGAAATCCACTCAGAATGCCATCAATGGCATTACTAACAAAGTAAATACTGTGATTG AAAAGATGAACATCCAGTTTACCGCAGTAGGAAAAGAGTTCAACAAATTGGAGAAAAGGATGGAAAACCTCAACAAAAAAGTCGAC GATGGATTCCTTGATATTTGGACATACAACGCTGAGCTGCTAGTGCTGCTGGAGAACGAGCGAACACTCGATTTCCATGATAGCAA CGTCAAAAACCTGTACGAGAAGGTCAAAAGCCAGCTGAAAAATAACGCTAAGGAGATTGGGAATGGCTGTTTCGAATTCTACCACA AATGTGATAATGAATGCATGGAGAGTGTGCGGAATGGGACATACGACTACCCAAAATATTCTGAAGAGAGTAAACTTAATCGAGAG AAAGTGGATGGTGTTAAACTTGAGTCTATGGGCATTTACCAGATCCTGGCAATTTACAGTACCGTCGCTTCTTCTTTGGTCCTACTA GTGAGCCTCGGAGCAATCAGCTTCTGGATGTGTAGTAACGGGAGCCTTCAGTGTCGAATTTGCATC
2687	ATGAAAGCTAATCTGCTCGTACTGCTCTGCGCTCTGGCTGCCGCTGATGCCGACACCATCTGTATTGGATATCACGCAAACAATTC TACTGATACCGTGGACACCGTGCTCGAGAAAAACGTGACCGTGACGCATTCCGTTAATCTGCTGGAAGATTCACACAACGGAAAG CTGTGTAGACTGAAGGGAATCGCCCCACTGCAATTGGGTAAATGTAACATTGCAGGATGGCTTCTGGGTAATCCTGAATGCGATC CACTGCTCCCTGTTAGATCATGGTCATATATCGTCGAAACACCGAATTCTGAAAATGGTATTTGCTACCCTGGCGATTTCATCGACT ACGAAGAGCTGAGAGAGCAGCTATCTAGCGTGAGTAGCTTCGAACGCTTCGAAATTTTTCCCAAAGAGAGCTCTTGGCCGAACCA CAATACCAATGGCGTGACCGCCGCATGCAGCCATGAGGGCAAATCTTCTTTCTACAGAAATCTACTTTGGCTGACCGAAAAAGAG GGAAGTTACCCAAAGCTGAAAAATTCCTATGTCAACAAAAAAGGGAAGGAGGTACTTGTGTTATGGGGAATTCACCACCCCCCTAA CTCCAAGGAGCAGCAGAACCTGTATCAGAATGAGAACGCGTATGTTTCAGTGGTCACCAGCAACTATAATCGTCGGTTCACCCCT GAGATCGCCGAACGGCCCAAGGTGAGAGATCAGGCTGGAAGAATGAATTACTATTGGACCTTGCTGAAGCCTGGGGATACCATCA TCTTTGAAGCTAATGGAAACCTGATCGCGCCTATGTACGCATTCGCCCTATCCCGTGGCTTCGGTTCCGGCATTATCACCTCAAAT GCCTCAATGCACGAGTGCAACACTAAATGTCAGACCCCGCTGGGTGCCATCAACTCCAGCCTTCCCTACCAAAACATACATCCAGT GACGATTGGCGAGTGCCCAAAGTACGTCCGCTCTGCCAAGCTACGCATGGTCACGGGCCTGCGAAATAACCCCTCTATACAATCC CGTGGGCTGTTTGGCGCGATTGCCGGGTTCATCGAGGGAGGATGGACAGGTATGATCGATGGCTGGTACGGATACCATCATCAG AACGAACAGGGCTCCGGCTACGCTGCGGACCAGAAGTCTACGCAAAACGCAATTAATGGAATCACAAACAAGGTGAACACTGTGA TCGAGAAAATGAATATTCAGTTTACCGCCGTCGGTAAAGAATTCAATAAGCTCGAGAAGAGAATGGAGAACTTGAACAAAAAAGTG GATGATGGGTTTCTCGATATTTGGACATATAACGCGGAACTCCTGGTGCTGCTCGAGAATGAGCGCACCCTGGATTTTCACGACTC TAACGTCAAGAATTTATACGAGAAAGTAAAATCCCAGCTGAAGAACAATGCTAAAGAAATTGGCAACGGATGTTTCGAGTTTTATCA TAAATGCGATAATGAGTGCATGGAGAGCGTGCGCAATGGAACATACGACTACCCTAAATACAGCGAGGAATCTAAGCTGAATAGA GAGAAGGTAGACGGAGTCAAGCTGGAATCTATGGGAATATATCAAATTTTAGCTATTTACTCCACAGTTGCTTCATCCCTGGTCTTA CTGGTTTCCCTCGGCGCGATCTCGTTTTGGATGTGCAGTAATGGCTCCCTTCAGTGTCGAATCTGCATC

2688	ATGAAAGCTAATCTGTTGGTGCTGCTCTGCGCTTTGGCCGCTGCCGATGCAGATACCATCTGTATCGGCTATCACGCGAACAATTC CACAGATACAGTGGACACCGTCCTGGAGAAAAATGTGACCGTGACCCACTCTGTCAACCTGCTCGAAGACAGCCATAACGGCAAA CTGTGTAGACTGAAGGGCATTGCACCTTTACAGCTGGGCAAGTGCAATATAGCTGGATGGCTCCTCGGCAATCCCGAATGCGACC CTCTACTGCCCGTGAGAAGCTGGTCATATATCGTTGAAACCCCGAATTCCGAGAATGGAATCTGCTATCCCGGAGATTTCATCGAC TATGAAGAGCTCAGGGAACAGCTGTCCTCTGTATCCTCCTTTGAACGTTTTGAGATTTTCCCAAAGGAGTCATCTTGGCCTAACCA CAACACGAACGGGGTAACCGCAGCCTGCAGTCACGAAGGTAAGAGCAGCTTTTACAGGAATCTGCTGTGGCTAACCGAGAAAGA GGGCAGTTATCCTAAACTGAAAAATTCATATGTGAACAAGAAAGGAAAGGAAGTGCTGGTGCTTTGGGGAATCCACCATCCGCCCA ACAGCAAGGAGCAGCAGAACCTATACCAGAACGAAAACGCTTATGTGTCAGTGGTGACTAGCAACTACAACCGACGTTTTACCCC CGAGATTGCCGAGCGACCCAAGGTGAGAGACCAGGCCGGTCGCATGAACTACTACTGGACACTCCTTAAGCCAGGCGATACAAT CATATTTGAAGCTAACGGGAACCTCATTGCTCCTATGTACGCTTTCGCGCTGTCTAGGGGTTTCGGCTCCGGAATAATTACAAGCA ATGCTTCCATGCACGAGTGCAATACCAAGTGTCAGACTCCCCTGGGGGCCATTAACAGCAGTCTGCCATACCAAAATATTCATCCA GTAACAATCGGCGAGTGTCCAAAGTACGTGCGATCTGCCAAGTTACGAATGGTGACCGGCCTGCGCAATAACCCCTCAATTCAGT CTAGGGGCTTATTCGGGGCTATCGCCGGCTTCATTGAGGGAGGATGGACAGGAATGATCGATGGATGGTACGGCTACCATCACC AGAATGAGCAAGGCAGCGGTTACGCTGCCGACCAGAAATCGACACAGAATGCTATCAATGGAATCACTAACAAAGTCAACACGGT GATTGAAAAGATGAATATCCAATTCACTGCAGTGGGCAAGGAATTCAACAAATTAGAGAAGCGGATGGAGAACCTGAATAAAAAAG TTGATGACGGTTTCCTGGATATCTGGACATACAATGCAGAACTTCTCGTCCTGCTCGAGAACGAGCGAACTCTCGATTTTCACGAT TCTAATGTCAAAAATCTGTATGAAAAGGTGAAGTCCCAGCTGAAGAACAACGCAAAGGAAATCGGGAATGGATGCTTCGAGTTTTA CCATAAATGTGATAACGAGTGCATGGAAAGCGTCCGAAATGGCACTTATGACTATCCAAAATACTCAGAGGAGAGTAAACTTAACC GCGAGAAGGTTGATGGCGTTAAACTGGAATCCATGGGCATTTACCAGATCTTGGCTATCTATAGCACGGTGGCTTCTTCGCTGGT GCTCTTGGTCAGTCTCGGTGCCATCTCATTCTGGATGTGTTCGAACGGATCTCTTCAATGCCGAATCTGTATA
2689	ATGAAAGCCAACCTTCTAGTTTTACTATGTGCTCTCGCTGCTGCTGACGCCGACACTATTTGTATTGGATATCACGCAAATAATTCC ACAGATACTGTCGACACCGTTTTGGAAAAGAACGTGACTGTCACTCACTCAGTCAACCTGCTCGAAGATTCTCACAATGGCAAATT ATGCCGCCTCAAGGGAATCGCTCCGCTCCAATTGGGAAAATGTAACATCGCAGGATGGCTCCTCGGCAATCCTGAATGCGACCCT CTCCTGCCCGTCCGCTCTTGGTCATACATAGTGGAAACTCCCAACAGTGAAAATGGAATATGTTACCCCGGTGACTTCATCGACTA TGAGGAATTGCGCGAGCAACTGTCCTCAGTTTCCTCGTTCGAAAGATTTGAGATCTTCCCCAAGGAGTCCTCCTGGCCCAATCACA ATACGAATGGCGTTACGGCAGCCTGCTCTCATGAGGGGAAATCCTCATTCTACCGCAACCTGCTCTGGCTGACAGAGAAGGAGGG GTCCTACCCTAAGCTCAAAAATTCCTATGTGAATAAAAAGGGCAAAGAAGTACTGGTCCTGTGGGGCATTCACCATCCCCCAAATA GTAAAGAACAACAGAACCTCTACCAGAATGAGAACGCTTACGTGAGTGTGGTCACTAGCAACTACAACAGGAGATTCACACCCGA GATAGCGGAACGACCCAAGGTCCGCGATCAGGCTGGTCGCATGAATTACTACTGGACCTTGCTGAAGCCCGGAGATACGATTATT TTTGAGGCCAATGGCAATCTTATCGCCCCAATGTACGCCTTTGCACTCAGCCGTGGGTTCGGAAGTGGAATCATAACTTCAAATGC CTCCATGCATGAGTGCAATACAAAATGTCAGACCCCTCTCGGAGCCATCAACTCATCACTGCCATACCAGAATATTCACCCTGTTA CAATTGGAGAGTGTCCAAAATACGTCAGAAGTGCAAAACTGCGCATGGTTACCGGACTACGTAACAACCCGTCCATCCAGTCAAG AGGGCTGTTTGGAGCCATCGCAGGATTTATAGAGGGAGGCTGGACCGGGATGATTGATGGCTGGTACGGCTACCATCACCAAAAT GAGCAGGGAAGTGGGTATGCTGCCGACCAGAAATCCACCCAAAACGCGATCAATGGTATCACCAACAAAGTCAATACCGTGATCG AAAAAATGAACATACAGTTTACAGCAGTGGGAAAGGAATTCAATAAACTTGAGAAACGCATGGAGAATCTGAATAAAAAGGTCGAC GATGGCTTCCTAGATATTTGGACCTACAATGCAGAATTACTGGTATTGTTGGAGAATGAGCGGACCCTCGACTTCCATGATTCGAA CGTTAAAAACCTCTACGAGAAGGTAAAGTCGCAACTGAAAAACAACGCTAAGGAGATCGGGAACGGCTGCTTCGAATTTTATCACA AATGCGACAACGAATGCATGGAATCAGTGCGAAATGGCACCTATGATTATCCAAAGTATTCGGAGGAGTCAAAGCTGAATAGAGAG AAGGTGGATGGTGTGAAGTTGGAGAGTATGGGGATTTACCAGATCTTAGCAATCTATTCCACGGTTGCTAGTTCCCTCGTGTTACT TGTGAGCCTGGGTGCGATCAGTTTCTGGATGTGCAGTAACGGTTCACTGCAATGCAGAATCTGCATC

2690	ATGAAAGCTAATCTCCTGGTTTTGTTGTGTGCACTGGCCGCTGCCGACGCCGACACAATCTGCATAGGATACCATGCTAACAACAG TACAGATACAGTTGACACCGTGCTGGAAAAGAATGTTACTGTAACCCATAGCGTTAACTTGCTAGAGGACTCTCATAATGGAAAGC TCTGTAGGCTTAAAGGGATCGCCCCCTTACAATTGGGCAAGTGCAATATCGCCGGATGGCTGCTCGGGAACCCTGAGTGTGATCC ATTGCTGCCAGTTCGTTCCTGGTCTTACATAGTTGAAACGCCAAACTCCGAGAACGGGATCTGTTACCCAGGTGACTTCATTGATT ACGAAGAACTGAGAGAACAGCTAAGTTCTGTGTCTAGCTTCGAGAGATTCGAGATCTTTCCCAAAGAGTCAAGTTGGCCTAACCAC AATACCAACGGGGTTACCGCGGCTTGCTCCCACGAAGGTAAAAGCTCATTCTACCGGAACCTTTTGTGGCTCACTGAGAAAGAGG GTAGCTACCCAAAGCTCAAGAACTCCTACGTGAATAAGAAGGGGAAGGAGGTGCTCGTTTTATGGGGTATCCACCACCCCCCTAA TAGTAAGGAACAGCAAAATCTCTATCAGAACGAAAACGCATATGTCTCGGTGGTGACCTCAAACTATAATCGGAGATTCACTCCAG AGATCGCCGAGAGACCCAAAGTCCGGGACCAGGCAGGAAGGATGAACTATTATTGGACTTTACTGAAGCCCGGGGATACTATTAT TTTTGAAGCCAATGGCAATCTTATTGCCCCCATGTACGCTTTCGCCCTGAGTCGCGGCTTCGGGAGTGGCATTATTACTTCCAATG CCTCGATGCATGAGTGCAATACGAAGTGTCAGACCCCATTAGGAGCTATCAATAGCAGCCTGCCATATCAGAACATCCACCCGGT GACTATTGGAGAGTGTCCAAAGTACGTTAGGTCCGCCAAGTTACGGATGGTAACAGGCCTCCGCAATAATCCTTCTATCCAAAGTC GGGGGCTGTTTGGAGCAATCGCCGGCTTCATCGAGGGAGGTTGGACTGGGATGATTGACGGATGGTATGGGTACCATCATCAGA ATGAGCAAGGCAGTGGCTACGCAGCCGACCAGAAGTCAACCCAAAACGCCATCAATGGAATCACGAATAAAGTGAACACAGTGAT CGAGAAGATGAACATTCAGTTCACTGCAGTGGGGAAGGAATTTAACAAGTTGGAGAAGAGAATGGAGAACCTGAACAAGAAGGTA GACGATGGATTCCTAGACATATGGACATATAATGCCGAGCTGTTGGTCCTCCTCGAAAACGAAAGAACACTGGACTTTCATGACAG TAACGTGAAGAATTTGTATGAGAAAGTTAAGTCTCAACTGAAAAACAACGCCAAAGAAATAGGGAACGGTTGTTTCGAGTTCTATCA TAAGTGCGATAACGAGTGCATGGAGTCTGTGCGGAATGGCACGTACGATTATCCAAAATACTCCGAGGAGTCCAAACTTAACAGA GAGAAAGTGGATGGCGTGAAACTTGAAAGCATGGGCATTTACCAAATCCTCGCCATTTACAGCACAGTCGCTTCTAGTCTAGTACT GCTTGTATCACTCGGCGCCATATCATTCTGGATGTGTAGTAATGGCAGCCTGCAGTGCCGAATTTGCATT
2691	ATGAAAGCCAACCTGCTCGTACTCCTGTGTGCCCTGGCTGCCGCTGACGCCGATACTATTTGCATAGGGTATCACGCCAATAATTC AACTGACACCGTCGATACCGTTCTTGAGAAGAACGTGACCGTCACACATTCAGTTAACCTTCTTGAGGACTCTCATAATGGTAAATT GTGTCGCCTCAAGGGTATCGCACCGCTACAACTAGGCAAGTGTAATATAGCAGGGTGGTTGTTGGGAAACCCTGAGTGTGACCCG CTGCTGCCCGTTCGCAGTTGGTCTTACATCGTCGAGACACCAAACTCTGAGAACGGGATTTGCTATCCAGGGGATTTTATCGACTA TGAAGAGCTTCGCGAGCAGTTGAGCAGTGTTTCCTCTTTTGAGCGTTTTGAAATTTTCCCGAAAGAGAGTAGCTGGCCTAACCATA ACACAAACGGGGTCACCGCAGCTTGCTCTCATGAGGGGAAGTCAAGCTTCTATCGGAATCTGCTGTGGCTGACGGAAAAAGAAGG CTCGTACCCAAAGCTAAAAAATTCATACGTGAACAAGAAGGGGAAAGAGGTCCTGGTCCTGTGGGGGATCCACCATCCACCTAAT AGTAAAGAGCAGCAAAATTTGTACCAAAACGAGAATGCGTACGTGAGCGTAGTGACATCAAACTACAACCGCCGCTTCACTCCTGA GATTGCTGAACGGCCCAAGGTGCGTGATCAGGCCGGCAGGATGAACTATTATTGGACTTTGCTAAAACCTGGGGATACTATAATTT TTGAGGCGAATGGCAATCTAATCGCACCGATGTATGCCTTCGCCTTGAGCCGGGGCTTCGGCAGTGGGATTATTACCAGTAACGC CTCCATGCACGAATGTAACACAAAGTGTCAGACTCCATTAGGGGCAATCAACTCGTCACTTCCGTATCAAAATATTCACCCTGTGA CCATCGGCGAGTGTCCAAAATACGTTAGAAGTGCCAAGCTGAGAATGGTGACGGGTCTCCGGAACAATCCCTCAATCCAATCAAG GGGGTTATTTGGCGCTATTGCTGGTTTCATTGAGGGGGGATGGACCGGAATGATCGACGGCTGGTACGGATATCATCATCAGAAC GAACAGGGGTCCGGCTATGCTGCCGACCAGAAGTCCACTCAGAACGCGATCAATGGGATCACCAACAAGGTCAACACCGTCATC GAAAAGATGAATATTCAGTTTACAGCCGTCGGAAAGGAATTCAATAAGTTGGAGAAACGCATGGAAAACTTAAACAAGAAAGTGGA CGACGGATTCTTAGACATATGGACTTATAACGCAGAGCTGCTTGTACTCCTGGAGAATGAGCGAACCTTGGATTTTCATGATTCAA ACGTTAAGAACTTATATGAGAAGGTGAAGTCTCAACTCAAAAACAATGCCAAAGAAATCGGGAATGGGTGCTTCGAATTCTATCATA AATGTGATAACGAGTGTATGGAGAGCGTGCGGAATGGAACTTATGACTACCCCAAATACAGCGAAGAAAGCAAGTTGAACAGAGA GAAGGTGGATGGCGTGAAGCTGGAAAGCATGGGGATTTACCAGATTCTTGCAATCTATTCTACTGTTGCATCCAGTCTAGTTCTGC TCGTCTCTCTGGGGGCAATCAGCTTTTGGATGTGCTCTAATGGTTCCCTGCAGTGCCGTATCTGCATA

2692	ATGAAAGCAAACCTTCTGGTACTGCTCTGCGCCTTAGCCGCCGCTGATGCGGACACGATTTGTATTGGCTATCACGCCAATAATTC AACAGATACTGTCGACACAGTCTTGGAGAAAAATGTGACTGTAACCCATTCTGTCAATCTGCTCGAGGATTCACACAATGGAAAAC TGTGTCGGCTTAAAGGAATCGCTCCTCTGCAGTTGGGCAAGTGTAATATAGCTGGCTGGCTGTTAGGAAACCCAGAATGCGATCC TCTCTTACCTGTGCGATCCTGGTCTTACATCGTTGAGACACCGAACAGCGAGAATGGTATATGCTACCCGGGTGACTTTATCGATT ATGAAGAATTGAGAGAACAGCTGTCAAGCGTCAGCTCTTTCGAACGATTCGAGATCTTTCCCAAAGAGTCAAGCTGGCCTAACCAC AATACAAATGGAGTGACAGCTGCGTGTTCCCATGAAGGCAAGAGTTCTTTCTACCGAAACCTGCTTTGGCTGACCGAGAAGGAGG GATCATACCCTAAGTTAAAAAATTCATACGTCAACAAGAAGGGAAAAGAGGTACTAGTTCTTTGGGGTATTCACCACCCACCCAACT CCAAAGAGCAGCAGAATCTGTATCAAAATGAAAACGCGTACGTCTCCGTGGTGACCTCGAACTACAATCGACGATTCACTCCCGAA ATCGCTGAGAGGCCCAAAGTCCGGGATCAGGCCGGACGCATGAATTATTACTGGACTCTACTAAAGCCAGGTGATACAATTATCTT CGAAGCCAATGGCAATCTTATCGCTCCCATGTACGCATTCGCTCTGAGCCGTGGTTTTGGGTCTGGAATCATCACCTCCAACGCTT CTATGCATGAATGTAATACAAAATGCCAGACTCCGTTAGGGGCAATCAACAGTTCACTGCCATACCAAAACATCCACCCCGTGACT ATTGGGGAGTGCCCAAAGTATGTCCGGAGCGCCAAACTCCGGATGGTCACCGGCCTACGCAATAACCCCAGTATCCAGTCTAGG GGATTGTTCGGTGCTATCGCAGGTTTTATTGAGGGCGGCTGGACAGGAATGATCGACGGATGGTATGGGTATCATCACCAGAATG AACAAGGCTCCGGGTATGCGGCTGACCAAAAATCCACGCAGAACGCAATCAACGGAATCACCAATAAAGTGAATACTGTGATAGA AAAGATGAACATTCAGTTCACGGCTGTAGGCAAGGAGTTCAATAAGCTGGAGAAGAGAATGGAGAACCTAAATAAGAAGGTCGAT GACGGCTTCCTTGACATCTGGACTTACAATGCAGAACTGCTGGTACTGCTTGAGAACGAACGCACCCTGGATTTCCACGACTCTAA TGTGAAGAACCTATACGAAAAAGTGAAGTCCCAGCTGAAAAACAATGCTAAAGAGATAGGCAACGGCTGCTTCGAATTCTATCATA AATGCGACAACGAGTGCATGGAGTCAGTCCGAAATGGTACGTACGATTACCCAAAATATTCTGAAGAGTCCAAGCTAAATAGAGAA AAGGTGGACGGCGTAAAACTCGAGTCCATGGGCATCTACCAGATTCTGGCCATCTACAGTACGGTGGCTTCATCGCTTGTACTCC TGGTGAGCCTAGGGGCCATCTCTTTTTGGATGTGTTCTAACGGTTCCCTGCAGTGCCGCATTTGTATT
2693	ATGAAAGCTAACCTGCTGGTACTCTTATGCGCTCTGGCCGCTGCAGACGCCGATACCATCTGTATCGGGTATCATGCCAACAATAG CACCGACACCGTGGATACAGTCCTAGAAAAGAATGTGACCGTCACTCACTCAGTGAACCTATTAGAAGATTCCCACAACGGCAAGT TGTGTCGCCTCAAAGGCATCGCCCCACTGCAACTTGGGAAATGCAACATAGCCGGGTGGCTGCTGGGCAACCCAGAGTGCGATC CCCTGTTGCCAGTGCGCAGTTGGTCATATATCGTGGAGACTCCAAATTCAGAAAATGGGATTTGCTATCCAGGGGACTTCATCGAC TACGAAGAACTGCGCGAACAGCTGTCCAGCGTGAGTTCCTTCGAACGATTTGAAATTTTCCCAAAGGAGAGCAGCTGGCCTAACC ACAACACGAACGGAGTGACTGCCGCTTGCAGCCACGAGGGAAAGAGCTCTTTCTACCGCAACCTTCTGTGGTTGACAGAGAAGGA AGGGAGCTACCCCAAACTGAAGAACTCCTACGTTAATAAGAAAGGTAAGGAAGTGCTAGTGCTCTGGGGTATCCATCATCCACCAA ATAGCAAAGAGCAACAGAATCTGTATCAGAACGAGAACGCTTACGTGTCAGTTGTCACTTCCAACTATAACCGCCGGTTCACTCCA GAGATCGCTGAGCGCCCCAAGGTGCGCGATCAGGCTGGCCGCATGAACTATTACTGGACCTTGCTGAAGCCAGGGGACACAATT ATCTTTGAGGCCAATGGAAATCTCATCGCACCGATGTACGCCTTTGCCCTTTCCCGGGGCTTCGGGTCTGGCATCATTACCTCGAA TGCTTCGATGCACGAGTGCAACACCAAGTGCCAAACTCCCTTGGGAGCTATTAATTCTTCACTCCCATACCAGAATATTCACCCAG TTACAATCGGCGAGTGTCCAAAGTACGTGCGTTCCGCAAAACTGCGGATGGTTACAGGTCTGCGAAACAATCCAAGTATACAGAG TAGAGGACTTTTTGGAGCTATTGCCGGATTCATAGAAGGGGGCTGGACCGGAATGATTGACGGTTGGTACGGCTACCACCATCAA AACGAACAGGGGTCCGGATACGCCGCAGACCAAAAAAGTACCCAAAATGCTATTAATGGCATAACCAATAAAGTCAACACGGTCAT TGAAAAGATGAACATCCAGTTCACTGCTGTGGGAAAGGAGTTCAATAAGCTGGAGAAACGAATGGAGAACCTCAACAAAAAAGTAG ATGACGGCTTTTTGGACATCTGGACCTACAATGCAGAGCTTCTGGTGCTATTAGAGAACGAACGTACATTAGACTTTCACGACTCC AACGTCAAAAACCTGTATGAGAAGGTGAAATCCCAGCTTAAGAATAACGCAAAGGAAATCGGGAACGGCTGCTTTGAATTCTATCA CAAGTGTGACAATGAGTGTATGGAGAGCGTGCGCAATGGGACCTACGACTATCCTAAGTACTCAGAAGAGTCTAAACTGAACCGT GAAAAGGTCGACGGCGTTAAGCTGGAGTCTATGGGGATCTATCAGATCCTTGCAATTTATTCCACCGTGGCGTCTTCCTTGGTTTT GCTAGTGAGTTTAGGTGCTATCAGTTTCTGGATGTGCTCCAATGGCTCCCTGCAATGTAGGATCTGCATT

2694	ATGAAGGCCAACCTCCTGGTCTTGCTGTGCGCCCTGGCTGCTGCCGATGCAGATACCATTTGCATCGGCTATCACGCCAACAATT CTACAGATACCGTGGATACAGTACTCGAAAAAAATGTTACTGTGACGCACTCAGTCAACCTGTTGGAAGACTCGCACAATGGCAAA CTGTGCAGGCTGAAAGGCATTGCCCCCCTACAGCTGGGGAAGTGCAACATTGCAGGCTGGCTGTTGGGAAACCCTGAATGTGAT CCATTGCTCCCAGTGCGGTCCTGGTCTTACATCGTGGAGACACCAAACAGCGAGAATGGGATCTGTTACCCTGGCGATTTTATAG ACTACGAGGAGCTCAGGGAGCAGCTCAGCTCTGTGTCGTCTTTCGAGCGCTTTGAGATTTTTCCTAAGGAGTCCTCCTGGCCAAA CCACAACACGAATGGAGTGACAGCCGCCTGTAGCCACGAGGGCAAGTCCTCATTCTATAGAAACTTGCTTTGGCTAACTGAAAAG GAGGGCAGCTACCCTAAACTGAAAAATAGCTATGTGAACAAAAAAGGCAAAGAAGTGCTCGTCCTGTGGGGGATCCATCATCCTC CTAACTCAAAGGAGCAGCAGAACCTGTATCAGAATGAAAACGCTTACGTGTCTGTCGTGACTTCAAATTATAACCGGAGGTTCACT CCCGAAATTGCCGAGAGACCAAAGGTCAGGGACCAGGCAGGCCGCATGAATTACTATTGGACACTTCTGAAACCAGGCGACACAA TTATTTTCGAGGCAAACGGGAACCTTATAGCCCCAATGTACGCCTTCGCACTATCAAGAGGCTTCGGTTCCGGCATCATTACAAGC AACGCGAGCATGCATGAATGCAACACCAAGTGCCAAACCCCACTGGGTGCAATTAATAGTTCCCTGCCATATCAGAATATCCACCC GGTGACAATTGGAGAGTGCCCTAAGTACGTGAGGAGCGCTAAGTTAAGAATGGTGACGGGACTGAGAAACAACCCTTCCATTCAG TCCCGAGGACTTTTTGGAGCAATCGCCGGCTTTATTGAGGGGGGCTGGACGGGGATGATTGATGGCTGGTATGGGTATCACCAC CAGAATGAACAGGGAAGTGGGTACGCTGCTGATCAGAAGTCAACTCAGAATGCCATCAACGGGATAACAAATAAGGTAAATACTGT GATTGAAAAAATGAATATCCAGTTCACTGCCGTTGGGAAGGAGTTTAACAAGCTTGAGAAAAGGATGGAAAACCTGAATAAGAAGG TAGATGACGGATTCCTCGATATATGGACCTACAACGCTGAGCTGCTGGTTTTGCTAGAGAACGAAAGAACTTTGGATTTTCATGATT CTAACGTGAAGAACTTGTACGAAAAAGTGAAAAGTCAGCTTAAAAACAACGCAAAGGAGATTGGGAATGGCTGCTTTGAGTTTTAT CACAAGTGCGACAATGAATGTATGGAGAGCGTGAGGAACGGCACGTATGACTACCCCAAGTACAGTGAGGAATCCAAGCTGAACC GAGAAAAGGTCGACGGCGTGAAGCTGGAGTCTATGGGAATCTATCAGATTCTCGCAATTTACTCAACTGTCGCATCATCCCTAGTG CTCCTGGTGAGCCTGGGAGCCATTTCTTTCTGGATGTGTAGCAATGGTAGTTTACAATGCAGAATCTGTATT
2695	ATGAAGGCAAATCTGCTGGTCTTACTGTGCGCTCTGGCCGCTGCTGATGCTGATACAATCTGTATTGGCTACCACGCCAATAACTC CACTGATACAGTGGATACTGTGCTGGAAAAAAATGTTACAGTCACACATTCAGTGAACCTCCTAGAAGACAGCCATAACGGAAAAC TCTGCCGGCTGAAAGGTATCGCACCTTTGCAGCTAGGAAAGTGCAATATTGCGGGCTGGCTGCTTGGTAATCCCGAGTGCGACCC ACTCTTACCAGTGAGAAGTTGGAGCTATATCGTAGAGACCCCCAACTCTGAGAATGGAATCTGTTATCCAGGCGACTTTATTGATTA CGAGGAACTGCGCGAGCAACTCTCTTCTGTGTCTTCATTCGAGAGGTTTGAAATCTTCCCTAAGGAGTCGTCTTGGCCTAATCACA ATACCAACGGGGTGACCGCCGCTTGCAGCCACGAGGGGAAGAGTTCTTTTTACCGAAATCTTCTGTGGCTAACAGAAAAGGAAGG ATCATACCCAAAATTAAAGAATAGCTATGTAAACAAGAAGGGTAAGGAGGTTCTTGTCTTATGGGGCATACATCATCCCCCCAACTC TAAGGAACAACAGAATCTATATCAGAACGAGAATGCTTACGTGAGTGTGGTGACCAGTAATTATAATAGAAGATTCACTCCTGAGAT TGCCGAAAGGCCTAAAGTGCGCGACCAAGCGGGACGGATGAACTACTACTGGACCTTGCTCAAGCCGGGTGACACAATTATTTTT GAAGCAAATGGTAATCTCATTGCACCAATGTATGCGTTTGCCCTGAGTCGTGGATTTGGCAGCGGAATAATAACTTCAAACGCCAG CATGCACGAATGCAATACTAAATGTCAGACCCCACTGGGGGCAATCAACTCCTCACTGCCATATCAGAACATCCACCCCGTGACCA TTGGTGAGTGCCCTAAGTATGTTAGATCAGCGAAGCTTCGGATGGTAACAGGACTTAGAAATAATCCGAGCATTCAAAGCCGCGG GCTGTTTGGAGCCATTGCAGGGTTTATCGAAGGGGGTTGGACCGGCATGATTGATGGCTGGTATGGATACCACCATCAGAACGAA CAAGGCAGCGGGTATGCAGCTGACCAAAAGTCTACCCAGAACGCCATCAATGGAATCACGAACAAGGTTAATACAGTGATTGAGA AGATGAATATTCAGTTTACAGCCGTTGGCAAGGAGTTCAACAAGCTGGAGAAGCGAATGGAGAACTTAAATAAAAAGGTCGACGAC GGCTTCCTGGACATTTGGACATACAACGCTGAACTGTTGGTGCTTCTCGAGAATGAACGAACTTTAGATTTTCACGATAGCAACGT TAAAAACTTATATGAGAAAGTGAAGTCTCAGCTCAAGAATAATGCCAAGGAGATAGGTAATGGTTGCTTCGAATTCTATCACAAGTG TGACAACGAATGCATGGAGAGTGTTAGAAACGGTACATACGATTATCCCAAGTATAGTGAGGAGTCCAAACTCAATAGGGAAAAGG TGGACGGGGTGAAGCTCGAGAGCATGGGCATTTATCAGATCCTTGCAATATACTCAACCGTAGCATCTTCTCTCGTTCTTCTGGTG TCCCTGGGCGCTATTTCTTTTTGGATGTGCAGTAATGGCAGCCTTCAATGTAGAATTTGCATC

2696	ATGAAGGCCAATCTCCTGGTGCTTCTATGTGCACTGGCCGCGGCCGATGCCGACACCATTTGCATTGGCTATCACGCCAATAATT CCACTGACACAGTTGATACCGTGCTCGAAAAAAACGTAACCGTGACCCATTCCGTTAACCTGCTTGAGGATTCTCACAACGGGAAA CTCTGCAGACTGAAGGGCATCGCCCCCCTGCAGCTGGGAAAATGCAATATTGCCGGCTGGCTCTTAGGTAACCCAGAATGTGATC CATTACTGCCGGTCAGAAGCTGGAGCTACATCGTGGAAACCCCCAACTCTGAGAATGGAATCTGTTATCCCGGAGATTTTATTGAT TACGAGGAGCTCCGCGAACAGCTATCCTCTGTATCTTCTTTTGAGAGATTCGAGATTTTCCCCAAGGAGAGCAGCTGGCCAAACCA CAACACTAATGGTGTCACCGCCGCCTGCTCCCATGAAGGAAAGTCCAGCTTTTATCGGAACTTACTGTGGCTGACAGAGAAGGAG GGCTCTTACCCCAAGCTGAAGAACAGCTATGTCAATAAAAAGGGCAAGGAAGTCCTAGTGTTGTGGGGGATCCACCATCCACCTA ACAGCAAAGAGCAGCAGAACCTCTACCAGAATGAGAATGCGTACGTTTCTGTGGTGACTTCAAACTATAACCGCCGATTCACACCC GAGATCGCAGAGAGACCTAAAGTCAGGGACCAAGCGGGGAGAATGAACTACTACTGGACCCTCTTGAAGCCCGGCGATACCATT ATTTTTGAGGCTAACGGCAACCTGATCGCCCCCATGTATGCATTCGCTCTGAGCCGTGGCTTCGGATCTGGTATTATAACCTCAAA CGCAAGCATGCATGAATGTAACACCAAATGCCAGACCCCCCTGGGGGCCATCAACAGCAGTCTGCCATACCAGAACATCCACCCT GTCACCATTGGTGAATGCCCCAAGTATGTCCGATCAGCGAAACTGCGGATGGTCACGGGGCTGCGAAACAATCCTTCAATCCAGA GCCGAGGACTTTTTGGAGCTATCGCCGGATTCATTGAAGGCGGCTGGACAGGGATGATCGACGGTTGGTACGGGTATCACCACC AGAACGAGCAAGGCAGTGGCTACGCAGCTGACCAGAAATCCACCCAAAACGCCATTAACGGGATCACTAACAAAGTGAACACGGT CATCGAGAAGATGAATATTCAATTCACTGCCGTTGGCAAAGAGTTTAATAAGTTGGAAAAGCGCATGGAGAATCTAAACAAAAAGG TTGACGATGGTTTCTTGGATATCTGGACATATAATGCCGAGCTGTTAGTCCTGTTAGAAAACGAGAGAACCCTTGATTTTCATGATA GCAACGTCAAAAACTTATACGAGAAGGTGAAAAGCCAGCTTAAAAATAATGCCAAAGAGATAGGAAATGGCTGTTTTGAGTTCTATC ATAAGTGCGATAACGAGTGCATGGAGTCCGTTCGGAATGGTACCTACGATTACCCGAAGTACTCTGAGGAGAGCAAGTTGAATCG GGAGAAAGTTGATGGCGTCAAGCTCGAATCCATGGGTATCTACCAGATATTAGCTATTTACTCCACAGTCGCATCTAGCCTGGTCC TGCTGGTTTCGCTGGGCGCCATATCTTTTTGGATGTGTTCTAATGGCAGCTTACAATGCAGGATTTGTATC
2697	ATGAAAGCCAACCTCCTCGTGTTGCTGTGCGCTTTGGCCGCTGCCGACGCCGACACCATCTGTATAGGCTATCATGCGAATAATA GCACCGACACAGTAGATACCGTTCTGGAGAAGAACGTGACCGTGACTCACAGCGTGAATCTGTTGGAGGACTCCCATAATGGGAA GCTGTGTAGATTGAAAGGTATTGCACCGCTACAACTCGGGAAGTGCAATATCGCCGGGTGGCTGCTTGGTAACCCCGAGTGTGAC CCACTGCTGCCAGTTCGTTCTTGGAGTTACATCGTGGAGACTCCTAATAGCGAGAATGGTATCTGCTACCCTGGGGACTTCATCGA TTATGAGGAGCTCAGAGAGCAGTTGAGTTCCGTGTCATCATTTGAGCGGTTTGAGATTTTTCCAAAGGAAAGTTCGTGGCCGAACC ATAACACTAACGGAGTCACCGCCGCCTGTAGTCACGAGGGCAAATCTTCTTTTTACCGGAATTTGCTGTGGCTTACAGAAAAAGAG GGCTCCTATCCGAAGCTGAAGAACTCTTATGTGAATAAAAAAGGAAAGGAAGTCCTAGTGCTGTGGGGAATCCATCATCCTCCAAA TAGTAAGGAGCAGCAGAACCTATATCAGAACGAGAACGCATACGTGTCCGTGGTCACGTCAAACTACAACCGGAGGTTCACTCCC GAGATCGCCGAAAGGCCTAAGGTTCGGGATCAGGCTGGGCGCATGAACTACTACTGGACTTTACTGAAACCAGGAGACACCATTA TCTTCGAGGCTAACGGCAATCTCATAGCTCCGATGTACGCCTTTGCCTTGAGTCGCGGATTCGGAAGTGGTATCATTACTAGCAAT GCCTCAATGCATGAGTGTAACACAAAATGCCAGACCCCCTTGGGAGCAATTAACTCATCTCTACCATATCAGAACATCCACCCTGT CACTATAGGGGAGTGTCCCAAGTATGTCAGATCCGCCAAGTTGCGGATGGTAACCGGGCTGCGGAATAACCCCTCCATCCAGTCC CGTGGCCTTTTTGGCGCCATTGCGGGATTTATTGAGGGAGGCTGGACGGGCATGATTGATGGCTGGTATGGTTATCACCACCAGA ATGAACAGGGGTCTGGTTATGCTGCCGACCAGAAGAGTACCCAAAACGCCATCAATGGAATTACAAACAAGGTGAATACAGTGAT CGAAAAAATGAACATCCAGTTTACTGCAGTTGGTAAAGAATTCAACAAACTAGAAAAGCGGATGGAGAACCTCAACAAGAAGGTCG ATGACGGCTTTCTGGACATCTGGACATACAACGCCGAACTTCTAGTGCTTCTGGAGAATGAGCGCACACTGGATTTTCACGACTCA AATGTTAAGAACCTCTACGAAAAGGTGAAGTCCCAGCTCAAAAACAACGCCAAGGAGATTGGCAATGGCTGTTTTGAATTTTACCA CAAATGCGACAATGAATGTATGGAGTCCGTCCGGAATGGCACCTATGACTATCCAAAGTATTCTGAGGAATCCAAGCTGAATCGGG AAAAAGTGGATGGTGTGAAACTTGAGTCAATGGGAATTTATCAGATCCTGGCCATTTACAGTACGGTGGCCTCCTCACTGGTGCTA CTGGTAAGTCTGGGAGCAATTTCATTCTGGATGTGCTCCAATGGTTCACTGCAGTGCAGAATCTGCATA

2698	ATGAAGGCTAACCTGCTGGTACTTCTGTGTGCGCTCGCGGCAGCTGATGCCGACACGATCTGTATTGGATATCATGCGAACAATA GCACAGACACCGTGGATACAGTTCTGGAAAAGAACGTGACTGTCACGCACTCCGTCAACTTACTAGAGGACTCTCACAACGGGAA GCTCTGCAGACTGAAGGGAATTGCACCCTTGCAGCTGGGTAAGTGTAACATCGCTGGATGGCTACTTGGTAACCCCGAGTGTGAC CCTCTCCTGCCCGTGCGGTCCTGGAGCTACATAGTTGAGACACCAAACTCTGAGAACGGGATCTGCTATCCTGGGGACTTCATTG ACTACGAGGAGCTGCGGGAGCAGCTCTCCTCCGTCTCGTCTTTCGAAAGGTTCGAGATTTTCCCAAAAGAATCAAGTTGGCCCAA CCACAACACAAATGGTGTTACCGCCGCCTGTAGCCATGAAGGGAAATCAAGCTTCTACCGTAATCTGCTTTGGCTTACCGAAAAAG AAGGAAGCTACCCTAAGTTAAAAAACTCTTATGTCAACAAAAAAGGGAAAGAAGTTTTGGTGCTGTGGGGCATCCACCACCCTCCT AATTCGAAAGAACAGCAGAACTTGTATCAGAACGAAAATGCCTACGTGTCCGTGGTGACCAGTAACTACAACCGACGGTTCACCCC GGAGATAGCAGAGAGGCCAAAAGTCAGAGATCAGGCTGGCAGAATGAACTATTACTGGACCCTCCTAAAACCCGGAGACACTATT ATATTTGAGGCAAATGGTAATCTGATCGCTCCAATGTATGCTTTCGCCCTCTCCCGCGGATTCGGGTCAGGGATAATCACCTCTAA CGCTTCTATGCACGAATGTAACACCAAGTGCCAAACTCCCCTTGGAGCCATCAACAGTTCTCTACCATACCAGAATATCCATCCAG TGACGATCGGCGAATGCCCTAAATACGTCAGAAGTGCTAAGCTAAGAATGGTTACTGGGCTGAGAAATAATCCGTCCATTCAGAGC AGGGGGCTGTTTGGAGCGATCGCTGGATTTATTGAAGGAGGTTGGACCGGGATGATCGATGGCTGGTACGGTTATCACCATCAAA ATGAGCAGGGAAGCGGATACGCCGCAGACCAGAAAAGCACACAGAACGCTATTAACGGAATCACTAATAAGGTGAATACTGTCAT CGAGAAAATGAACATTCAGTTCACAGCAGTGGGCAAAGAGTTTAACAAGCTTGAAAAGAGGATGGAGAATCTCAACAAGAAGGTG GATGACGGCTTCCTCGACATCTGGACTTATAACGCCGAGCTCCTCGTCCTGCTCGAAAATGAAAGGACTTTGGACTTCCACGATTC AAACGTGAAGAATTTGTACGAAAAGGTGAAGTCCCAGCTGAAAAACAACGCTAAGGAGATAGGGAATGGCTGTTTCGAATTCTACC ACAAGTGCGACAATGAGTGCATGGAGAGTGTTAGAAATGGGACGTATGACTATCCTAAGTATTCGGAGGAATCAAAGTTGAATCGA GAGAAGGTGGACGGTGTGAAGTTAGAGTCGATGGGCATCTACCAGATTCTGGCTATCTATTCCACCGTGGCTAGCAGCTTGGTCC TCCTCGTATCATTGGGCGCAATTAGCTTCTGGATGTGCTCCAACGGTTCCCTTCAGTGCAGAATCTGTATC
2699	ATGAAAGCAAATCTGCTGGTGCTGTTGTGCGCGCTGGCGGCTGCTGACGCAGACACCATTTGCATCGGCTACCACGCCAACAACA GTACTGATACTGTGGACACAGTTTTAGAAAAGAATGTGACCGTAACACATAGCGTAAACTTGCTGGAAGACAGTCACAATGGAAAA CTCTGCCGGCTGAAGGGAATCGCCCCTCTACAGCTCGGCAAATGCAATATCGCCGGGTGGCTATTGGGGAATCCAGAGTGCGAC CCACTTCTGCCTGTGCGCAGTTGGTCATATATCGTGGAAACACCAAACTCCGAAAACGGCATTTGCTATCCTGGAGACTTTATTGA CTACGAAGAGCTAAGAGAGCAGCTGAGCTCCGTGTCCAGTTTTGAGCGATTTGAGATCTTCCCAAAGGAGAGCTCTTGGCCAAAT CACAATACCAACGGAGTGACCGCGGCTTGTTCTCACGAGGGTAAATCCTCCTTTTATAGGAACCTGCTGTGGCTGACCGAGAAAG AGGGATCTTACCCAAAACTTAAGAACAGCTATGTTAACAAGAAGGGAAAAGAAGTGCTTGTGCTGTGGGGTATTCACCACCCTCCT AATTCCAAGGAGCAGCAGAATCTGTATCAGAATGAGAATGCCTACGTCAGCGTGGTCACCTCAAATTATAACAGACGGTTCACCCC AGAGATTGCGGAGCGTCCCAAGGTAAGGGACCAAGCCGGACGCATGAACTATTATTGGACTCTGCTGAAACCTGGTGATACTATC ATCTTTGAGGCCAATGGCAATCTTATCGCCCCTATGTATGCATTTGCCCTATCTCGCGGATTTGGGAGCGGCATCATCACAAGTAA TGCCAGCATGCACGAATGTAATACCAAATGCCAGACACCTCTGGGCGCTATCAACAGCAGCCTGCCTTATCAGAATATACACCCTG TCACCATCGGGGAATGCCCTAAATATGTTCGCAGTGCCAAACTTCGTATGGTGACTGGGCTGCGCAACAACCCATCAATTCAGAG CAGGGGCCTGTTCGGAGCGATTGCCGGGTTTATCGAGGGAGGGTGGACGGGCATGATAGATGGCTGGTACGGATACCACCATCA AAATGAACAGGGCAGTGGCTACGCAGCGGACCAGAAATCCACCCAGAATGCTATAAACGGTATCACAAATAAGGTGAATACAGTC ATCGAAAAGATGAATATTCAGTTCACTGCGGTCGGTAAGGAGTTCAACAAACTGGAGAAGAGGATGGAAAATCTTAACAAAAAAGT GGACGACGGCTTCCTGGATATATGGACGTATAATGCAGAACTGTTAGTGTTACTCGAGAATGAGAGGACTCTCGACTTCCATGATT CCAACGTGAAGAATCTCTATGAGAAGGTGAAATCCCAACTGAAGAACAACGCCAAAGAGATAGGGAATGGATGCTTCGAATTCTAC CATAAGTGTGATAACGAGTGTATGGAGTCTGTGCGGAATGGCACTTACGACTATCCGAAATACAGTGAGGAGTCAAAGTTGAACC GGGAGAAGGTCGATGGCGTCAAGCTGGAGAGTATGGGAATCTATCAGATCCTCGCAATTTACTCAACCGTGGCTAGCTCCCTGGT ACTCCTGGTCTCACTTGGAGCCATCTCCTTCTGGATGTGCAGCAACGGCAGCCTTCAGTGCAGAATCTGCATT

2700	ATGAAGGCCAATCTTTTAGTGCTGTTGTGTGCACTGGCGGCAGCTGATGCTGACACCATCTGCATCGGATACCACGCCAATAACA GCACTGACACAGTTGACACTGTTCTGGAAAAAAACGTGACGGTGACCCACTCCGTCAATCTTCTGGAGGACTCCCACAACGGGAA GCTGTGTAGGTTGAAGGGTATCGCCCCGTTGCAGCTGGGCAAATGCAACATCGCGGGTTGGCTGCTTGGCAATCCTGAGTGCGA TCCACTGCTACCCGTACGGTCTTGGTCATATATCGTTGAGACCCCAAATAGCGAGAATGGCATCTGTTATCCTGGCGATTTCATCG ACTACGAAGAGCTGCGAGAACAACTGAGTTCGGTGTCTAGTTTCGAGAGGTTTGAGATCTTCCCAAAAGAAAGTTCCTGGCCTAAC CACAATACCAACGGGGTAACGGCCGCTTGTAGCCACGAGGGCAAATCAAGCTTTTACAGAAATCTGCTGTGGCTCACAGAAAAAG AAGGGAGCTATCCTAAATTAAAGAATTCATACGTGAACAAAAAAGGAAAGGAAGTACTCGTCCTGTGGGGAATTCACCACCCTCCA AATTCTAAAGAACAACAGAACTTGTACCAGAATGAAAACGCCTATGTATCCGTGGTTACCTCAAACTACAATCGGCGGTTCACCCC CGAAATAGCTGAGAGACCGAAGGTTCGGGACCAAGCGGGAAGAATGAACTATTACTGGACTCTCCTTAAGCCCGGTGATACTATC ATCTTTGAAGCCAACGGAAACCTTATCGCCCCAATGTATGCCTTCGCGTTATCCCGCGGGTTTGGAAGCGGTATTATTACTTCAAA CGCGAGCATGCACGAATGTAATACTAAATGTCAGACACCCCTTGGCGCCATCAATAGCAGTCTGCCATACCAAAATATTCACCCCG TGACAATTGGCGAGTGTCCAAAATATGTTAGAAGCGCCAAACTCAGAATGGTTACTGGCTTGCGGAATAACCCGTCCATTCAGTCC AGAGGGTTATTTGGGGCAATCGCCGGCTTTATAGAAGGGGGCTGGACTGGAATGATCGACGGTTGGTACGGATATCATCACCAGA ATGAACAGGGATCAGGATATGCCGCGGACCAAAAGTCTACCCAGAACGCCATCAACGGCATCACTAATAAGGTCAACACTGTGAT TGAGAAGATGAATATTCAGTTCACCGCAGTCGGCAAGGAATTCAACAAACTCGAGAAACGCATGGAAAACCTTAACAAAAAGGTGG ACGATGGGTTTCTGGACATTTGGACCTACAATGCCGAGCTGCTAGTGCTGCTGGAAAACGAGCGGACTCTTGATTTCCATGACTCT AATGTAAAAAACCTTTACGAAAAAGTTAAATCACAACTTAAAAATAATGCCAAAGAAATAGGGAACGGATGTTTCGAGTTCTACCAC AAATGTGATAACGAATGCATGGAGAGTGTTCGCAACGGAACATATGACTATCCTAAGTACAGCGAAGAGTCAAAACTAAATCGGGA GAAAGTCGACGGCGTGAAACTCGAATCCATGGGGATTTACCAGATCCTCGCAATCTATTCTACCGTGGCTAGCTCGCTGGTGCTG CTTGTTAGTCTGGGAGCTATCTCCTTTTGGATGTGTAGCAATGGAAGCCTGCAGTGTAGAATTTGCATA
2701	ATGAAGGCCAACCTGCTGGTCCTGCTGTGTGCACTCGCCGCCGCCGATGCCGACACTATTTGTATTGGCTACCACGCTAACAACT CCACAGATACAGTCGATACTGTGCTGGAAAAGAATGTTACCGTCACCCATAGCGTGAACTTGCTCGAGGATTCTCATAATGGGAAA CTCTGTCGGTTAAAGGGGATTGCACCTCTCCAACTGGGCAAGTGTAACATCGCGGGTTGGCTGCTTGGAAACCCTGAGTGCGATC CTCTACTTCCGGTCAGGTCGTGGAGTTACATTGTCGAGACCCCTAACTCAGAAAACGGTATCTGTTATCCGGGTGATTTTATCGAT TACGAAGAGCTCAGAGAACAGCTGTCCAGCGTGTCTTCATTCGAGCGATTCGAAATTTTCCCGAAGGAATCATCTTGGCCAAATCA TAATACTAACGGGGTGACGGCCGCATGTTCACATGAGGGCAAATCCAGCTTCTACCGAAACCTTTTGTGGTTGACGGAGAAGGAG GGCTCTTATCCCAAGCTGAAAAACAGCTATGTGAACAAGAAGGGGAAGGAGGTGCTCGTTCTGTGGGGGATCCATCACCCTCCTA ACAGCAAAGAGCAGCAAAACCTGTATCAGAATGAAAACGCCTATGTGTCCGTGGTGACGAGCAACTACAACCGGCGCTTTACCCC CGAAATTGCTGAACGGCCGAAGGTGAGGGATCAAGCCGGCCGGATGAATTATTACTGGACGCTGCTGAAACCTGGTGATACCATC ATCTTTGAAGCTAACGGAAACTTGATCGCTCCAATGTATGCATTCGCCCTGTCACGGGGATTCGGCAGCGGTATAATCACCAGTAA TGCCTCAATGCACGAATGCAATACCAAGTGCCAAACGCCCCTTGGTGCCATCAACTCATCTCTGCCCTATCAGAATATCCATCCTG TAACCATCGGAGAGTGTCCCAAGTATGTCAGATCAGCAAAACTGCGGATGGTCACAGGTCTCCGAAATAATCCCTCAATCCAGTCT CGCGGCCTGTTCGGCGCTATCGCCGGCTTTATTGAGGGCGGTTGGACCGGGATGATCGACGGCTGGTATGGATACCATCATCAG AACGAGCAGGGGAGCGGCTACGCGGCTGACCAGAAGTCAACCCAGAATGCTATCAATGGTATCACTAATAAGGTGAATACGGTGA TTGAAAAAATGAACATACAGTTCACCGCAGTGGGTAAAGAGTTCAATAAGCTGGAGAAGCGGATGGAGAATCTGAACAAAAAGGTC GATGATGGCTTCTTGGACATCTGGACCTATAACGCCGAGCTTCTTGTGCTGCTCGAGAACGAGAGAACACTGGACTTCCATGACT CTAACGTGAAGAACCTCTACGAGAAGGTGAAATCTCAGTTGAAGAACAACGCTAAGGAGATCGGCAATGGCTGCTTCGAATTCTAC CACAAGTGCGATAACGAATGCATGGAATCTGTTCGAAATGGAACTTACGATTACCCTAAATACAGCGAGGAAAGTAAGTTAAACAG AGAGAAAGTAGACGGGGTTAAGCTCGAAAGCATGGGGATCTATCAGATCCTAGCCATCTACTCGACTGTCGCGTCAAGTTTGGTG CTGCTGGTGTCTCTTGGAGCCATATCGTTTTGGATGTGCTCTAACGGCTCCCTGCAGTGCAGAATCTGCATA

2702	ATGAAGGCCAACTTGCTAGTGCTGCTGTGCGCGCTTGCAGCTGCGGATGCGGATACGATATGCATAGGATATCACGCCAATAATT CGACGGATACCGTGGACACAGTGCTGGAAAAGAATGTAACGGTCACACACAGCGTGAACCTGCTTGAAGATTCACACAACGGCAA ACTGTGCAGGCTCAAAGGAATCGCACCATTACAGCTGGGGAAATGCAACATTGCGGGGTGGCTGCTAGGAAACCCAGAGTGCGA TCCCCTGCTGCCCGTGCGGTCTTGGTCCTACATTGTAGAGACTCCGAACTCGGAGAACGGTATTTGCTACCCTGGAGATTTCATTG ACTATGAAGAACTCAGGGAGCAACTGTCCTCAGTTTCGAGCTTTGAAAGGTTCGAGATCTTTCCTAAGGAGTCCAGCTGGCCTAAC CACAATACAAACGGGGTGACCGCGGCCTGTTCACATGAGGGTAAATCTTCATTCTACCGCAACCTCCTGTGGTTAACCGAAAAAGA AGGAAGCTATCCTAAGCTGAAGAATTCTTACGTGAATAAAAAGGGTAAGGAAGTTCTGGTGCTGTGGGGCATCCACCACCCACCTA ATAGTAAGGAACAGCAGAACCTATACCAGAACGAAAATGCGTACGTGAGCGTGGTAACTTCAAACTATAATCGGAGGTTCACACCC GAAATCGCGGAGCGGCCCAAGGTGCGGGACCAGGCTGGGCGCATGAATTATTACTGGACACTGTTGAAACCTGGAGACACTATA ATCTTCGAGGCGAACGGAAACTTAATCGCTCCCATGTACGCATTCGCCCTGTCCCGTGGCTTTGGCAGCGGTATCATCACATCCA ACGCCTCTATGCACGAATGCAATACGAAGTGCCAAACCCCTCTCGGAGCCATTAATAGCTCCCTCCCATATCAGAACATTCACCCA GTGACCATAGGAGAGTGTCCTAAGTACGTTCGGAGCGCAAAACTCCGCATGGTGACGGGGTTGCGAAACAACCCTTCCATCCAGT CGAGGGGCCTTTTCGGGGCGATCGCTGGCTTCATCGAGGGAGGGTGGACCGGGATGATTGACGGCTGGTACGGGTATCACCAC CAGAACGAGCAGGGTAGCGGCTACGCCGCAGACCAGAAATCTACGCAGAATGCCATTAATGGGATTACAAACAAGGTTAATACCG TGATCGAGAAAATGAACATTCAGTTCACAGCCGTCGGCAAAGAATTTAACAAATTAGAGAAGCGGATGGAGAATCTGAATAAAAAA GTGGACGATGGATTCCTGGATATCTGGACGTATAATGCCGAGTTGCTCGTCCTTCTGGAAAACGAGAGAACCCTGGACTTTCACG ATTCAAATGTAAAGAACCTTTACGAAAAGGTGAAGTCTCAACTGAAAAACAACGCCAAGGAGATAGGGAATGGCTGCTTTGAATTC TACCATAAATGCGACAATGAATGTATGGAGTCTGTTAGGAATGGTACTTACGACTACCCGAAGTATTCCGAAGAGAGCAAGCTGAA TAGGGAGAAGGTGGATGGAGTGAAACTGGAATCTATGGGGATTTATCAGATCCTAGCTATTTATTCTACAGTAGCCAGCTCCCTCG TCCTTCTGGTCTCCCTGGGCGCAATTTCATTTTGGATGTGCAGCAACGGGTCCTTACAATGCCGCATCTGTATT
2703	ATGAAGGCTAACCTGCTTGTGTTATTGTGCGCCCTGGCGGCAGCAGACGCAGATACCATATGTATCGGGTATCACGCTAATAATAG CACTGACACCGTTGACACCGTCCTGGAAAAAAACGTGACCGTGACTCATTCAGTTAATCTTTTGGAGGACAGCCATAATGGCAAGC TTTGTAGACTCAAGGGCATCGCACCTTTACAGTTGGGTAAGTGCAACATCGCAGGATGGCTCCTTGGCAACCCTGAATGTGACCC GCTGCTCCCAGTCCGGAGTTGGTCATATATCGTAGAAACTCCCAATAGCGAAAACGGCATTTGCTACCCAGGCGACTTTATAGATT ACGAAGAGCTCCGGGAGCAGCTATCAAGTGTCTCGTCGTTTGAGCGTTTTGAGATTTTTCCGAAAGAATCCAGTTGGCCAAATCAC AACACAAACGGTGTAACAGCGGCTTGTTCACACGAGGGTAAAAGTTCTTTCTACAGAAATTTATTGTGGCTTACAGAGAAAGAAGG CTCTTACCCAAAGCTGAAGAATTCTTATGTGAACAAGAAGGGTAAGGAGGTGTTAGTACTGTGGGGAATACATCACCCTCCTAATA GCAAGGAGCAGCAGAACTTGTATCAGAACGAAAACGCATATGTGTCTGTGGTGACATCCAATTATAATAGGCGTTTTACTCCCGAA ATTGCAGAGCGACCTAAAGTGCGTGACCAGGCAGGTAGAATGAATTACTACTGGACCCTGTTGAAGCCCGGTGACACCATCATTT TCGAAGCAAATGGTAACCTGATCGCACCAATGTACGCATTTGCACTGTCCCGCGGATTTGGCTCCGGAATCATTACCAGTAATGCC TCTATGCATGAATGTAACACCAAATGCCAGACCCCCCTGGGCGCCATTAACTCCAGCCTGCCTTATCAGAATATACATCCTGTGAC AATAGGTGAGTGCCCCAAGTACGTGAGATCTGCAAAACTAAGAATGGTCACCGGACTCAGGAACAATCCCTCCATCCAGAGTAGG GGTCTGTTCGGAGCCATAGCAGGGTTCATTGAAGGTGGATGGACAGGGATGATCGATGGATGGTACGGCTACCACCACCAGAAT GAGCAGGGCTCTGGCTACGCCGCTGACCAGAAATCTACACAGAACGCCATTAATGGCATCACTAACAAAGTGAATACCGTCATTG AGAAGATGAATATTCAGTTCACTGCCGTTGGCAAGGAATTTAACAAACTGGAGAAGCGGATGGAGAATCTAAATAAGAAAGTAGAC GACGGGTTTCTCGATATTTGGACATATAACGCTGAGCTTCTAGTCCTACTCGAGAACGAGAGGACCCTCGACTTCCATGATAGCAA CGTGAAGAACCTATACGAGAAGGTCAAGTCTCAACTGAAGAACAACGCTAAGGAGATCGGAAATGGATGTTTCGAGTTTTATCACA AGTGTGATAACGAGTGTATGGAGTCTGTAAGGAACGGTACCTATGACTACCCAAAATACAGCGAGGAATCAAAGCTCAATCGTGAG AAAGTCGATGGAGTTAAGCTGGAATCTATGGGGATTTATCAGATCTTAGCTATCTATTCAACCGTTGCTTCCAGCTTAGTCCTCCTC GTGTCACTGGGGGCTATAAGTTTTTGGATGTGCAGTAATGGCAGTTTGCAGTGTAGGATCTGCATC

2704	ATGAAGGCTAATTTATTGGTGCTGTTGTGTGCACTGGCTGCCGCAGACGCCGACACCATCTGCATTGGATATCATGCCAACAATTC AACTGATACGGTCGATACGGTGCTTGAGAAAAATGTGACAGTTACACACTCTGTGAACCTTCTGGAAGACTCTCATAACGGCAAAC TTTGTCGGTTGAAGGGTATCGCCCCACTGCAGCTTGGTAAGTGCAACATCGCCGGTTGGCTCCTCGGAAATCCAGAGTGTGATCC ACTGTTACCTGTGCGTAGTTGGAGTTACATCGTGGAGACTCCTAATAGTGAAAACGGCATATGCTATCCAGGCGATTTCATCGATT ATGAAGAACTGCGTGAGCAGCTGTCCAGCGTGTCCTCATTCGAGCGCTTTGAGATCTTCCCGAAAGAGTCCAGTTGGCCAAACCA CAATACCAATGGGGTCACTGCCGCCTGTAGTCATGAGGGTAAAAGCTCCTTCTATCGGAATCTTCTCTGGCTGACCGAAAAAGAAG GGTCCTACCCCAAACTTAAGAATTCGTACGTCAACAAGAAGGGCAAAGAAGTACTTGTGCTGTGGGGTATCCACCACCCTCCCAA CTCTAAGGAGCAGCAGAATCTGTACCAGAATGAGAACGCTTACGTCAGCGTGGTGACCTCAAATTATAACCGAAGGTTCACTCCC GAGATAGCCGAGCGGCCTAAGGTCAGAGACCAAGCGGGTAGAATGAATTACTACTGGACACTACTAAAGCCCGGCGACACCATC ATCTTTGAGGCAAATGGAAATCTGATCGCCCCCATGTATGCATTTGCACTGAGCCGGGGGTTCGGATCTGGAATCATAACTTCAAA CGCCAGTATGCATGAATGCAATACAAAGTGTCAGACTCCCTTGGGTGCCATCAATTCCAGCCTGCCTTATCAGAATATCCACCCTG TGACAATTGGAGAGTGCCCAAAATACGTTCGTTCAGCTAAACTGAGAATGGTCACCGGCCTCCGGAACAATCCTAGTATACAGTCA CGCGGCCTTTTCGGCGCCATTGCCGGATTTATTGAAGGAGGCTGGACAGGCATGATAGACGGATGGTACGGATATCACCACCAG AACGAACAAGGGTCTGGCTATGCCGCAGACCAGAAATCCACCCAGAACGCTATCAATGGTATCACGAATAAAGTCAATACCGTCAT CGAGAAGATGAATATCCAGTTTACAGCTGTGGGAAAAGAGTTCAATAAGCTTGAAAAGAGGATGGAGAACCTGAATAAAAAAGTTG ATGACGGATTTCTCGACATCTGGACGTATAACGCTGAACTGCTTGTGCTGCTCGAAAACGAGAGGACATTGGATTTTCACGACTCC AATGTTAAGAATCTGTACGAGAAGGTGAAATCTCAGCTCAAGAATAATGCTAAGGAAATAGGCAACGGATGTTTCGAATTTTATCAT AAGTGCGACAATGAATGCATGGAATCAGTGCGCAACGGGACCTATGACTACCCCAAGTACAGCGAGGAGTCCAAGCTGAATAGAG AAAAGGTGGACGGCGTCAAACTGGAAAGTATGGGCATTTACCAAATTCTCGCCATCTACTCTACCGTTGCCAGCAGTCTCGTGCTG TTAGTATCACTCGGTGCCATCTCTTTTTGGATGTGTAGTAATGGAAGCCTGCAGTGCCGGATCTGTATC
2705	ATGAAGGCCAACCTGCTCGTGTTGCTGTGTGCACTAGCTGCCGCCGACGCAGACACCATCTGTATCGGCTACCATGCCAACAACA GCACCGACACAGTGGATACGGTGCTCGAGAAGAACGTGACTGTCACCCATAGTGTGAATCTCCTAGAGGATTCTCATAATGGGAA GCTATGCCGACTCAAGGGGATTGCACCTCTGCAACTGGGTAAGTGCAATATCGCCGGTTGGCTTCTTGGGAACCCCGAATGTGAT CCCCTGCTCCCCGTGCGATCTTGGTCATATATCGTTGAAACACCCAATAGCGAAAATGGCATATGCTACCCTGGAGATTTCATTGA CTATGAGGAACTCCGAGAACAGCTGTCAAGCGTGTCTAGTTTCGAACGGTTTGAAATATTCCCTAAGGAGAGTTCCTGGCCCAACC ATAACACAAACGGAGTGACCGCAGCCTGCTCTCATGAGGGGAAGTCCTCATTTTATCGCAATCTCCTGTGGCTCACTGAAAAGGA GGGAAGTTACCCAAAATTGAAAAACTCATACGTGAATAAGAAAGGCAAGGAGGTCCTCGTGCTGTGGGGGATTCATCACCCCCCT AACTCAAAAGAACAGCAGAACCTGTATCAGAACGAAAACGCCTACGTGAGTGTGGTGACCTCTAACTATAACCGGAGGTTCACGC CAGAGATTGCTGAAAGACCTAAAGTCAGGGACCAAGCCGGCAGAATGAATTACTATTGGACCCTGTTGAAGCCCGGCGATACCAT TATATTCGAGGCAAACGGCAACCTGATTGCCCCCATGTACGCGTTCGCCCTGAGTCGAGGCTTCGGCAGTGGGATTATTACCTCT AACGCCTCGATGCATGAGTGCAACACAAAATGTCAAACTCCACTTGGGGCCATCAACTCGTCTCTTCCTTACCAAAATATCCACCC GGTTACAATTGGGGAATGTCCCAAATACGTGCGGTCCGCAAAGCTTCGCATGGTCACGGGCCTGAGGAACAACCCCTCTATCCAG TCTCGCGGTCTATTTGGCGCTATTGCTGGCTTTATCGAGGGGGGGTGGACAGGGATGATTGATGGCTGGTACGGGTACCATCACC AAAATGAACAGGGGTCCGGCTATGCTGCTGACCAAAAATCTACTCAAAACGCCATTAATGGAATCACAAATAAAGTGAATACGGTC ATAGAAAAGATGAACATTCAATTTACAGCCGTGGGTAAGGAGTTCAACAAACTGGAGAAGCGAATGGAGAACTTAAACAAAAAGGT AGATGATGGTTTCCTCGATATCTGGACATATAATGCTGAGTTGTTGGTGCTCTTGGAGAATGAGCGGACCTTAGATTTCCACGACA GCAATGTTAAAAACCTCTATGAAAAGGTGAAAAGTCAACTGAAGAACAATGCGAAAGAGATCGGTAACGGGTGCTTCGAGTTCTAC CACAAATGCGACAACGAGTGTATGGAGAGTGTAAGGAATGGTACATATGATTATCCTAAATACTCTGAAGAGTCGAAGCTGAACAG AGAGAAAGTCGACGGCGTAAAGCTGGAGAGTATGGGTATCTACCAAATCTTGGCCATCTATTCTACCGTGGCGTCATCACTAGTTC TTCTGGTGTCCTTGGGGGCCATATCATTCTGGATGTGTTCAAACGGGTCCCTCCAGTGCCGCATCTGCATC

2706	ATGAAGGCAAACCTGTTGGTATTGTTGTGTGCCCTGGCAGCTGCGGATGCAGACACTATTTGCATCGGCTATCATGCAAATAATTC CACGGACACAGTTGACACCGTTCTGGAGAAAAATGTGACGGTCACTCATTCTGTGAACCTGCTGGAAGACTCTCATAACGGAAAG CTGTGCCGGTTGAAGGGAATTGCACCACTACAACTGGGAAAATGCAACATCGCCGGCTGGTTGCTAGGTAACCCTGAGTGCGATC CACTCCTTCCAGTGAGATCCTGGTCCTACATCGTCGAGACCCCAAACTCAGAGAACGGGATCTGTTATCCAGGAGATTTCATTGAC TATGAAGAGTTACGTGAACAGCTCTCTTCCGTGTCGTCATTCGAACGGTTTGAAATCTTCCCCAAAGAGTCATCCTGGCCAAATCAT AACACTAACGGCGTGACTGCAGCTTGTTCCCACGAAGGGAAATCATCTTTCTACAGAAATCTGCTGTGGCTCACCGAAAAAGAGG GGAGTTATCCCAAGCTGAAAAACTCCTACGTTAATAAAAAGGGAAAAGAAGTGCTTGTCTTATGGGGTATCCATCACCCCCCAAAT TCTAAAGAACAGCAGAACCTGTACCAGAACGAGAACGCGTATGTGTCAGTGGTGACCTCTAACTATAATCGCAGATTTACACCAGA GATCGCAGAGCGCCCTAAAGTACGGGACCAGGCCGGTCGGATGAACTACTATTGGACCCTACTGAAACCAGGCGATACCATCATC TTCGAGGCTAATGGGAACTTAATTGCCCCCATGTACGCTTTCGCACTTAGCCGTGGCTTTGGCAGCGGGATTATCACGTCTAATGC TTCCATGCACGAGTGTAATACGAAGTGTCAGACACCTCTTGGTGCCATTAACTCAAGTCTTCCCTATCAGAACATCCACCCGGTGA CCATCGGCGAGTGTCCCAAGTACGTTCGAAGCGCCAAACTGCGGATGGTCACAGGTCTAAGGAATAACCCCTCCATCCAGAGCC GCGGCTTGTTTGGTGCGATTGCCGGGTTTATCGAGGGAGGGTGGACTGGCATGATAGACGGATGGTACGGATACCACCACCAAA ATGAACAAGGCTCCGGATACGCCGCCGACCAAAAATCCACGCAGAACGCCATAAACGGAATTACAAACAAGGTAAACACAGTCAT AGAAAAAATGAATATTCAGTTTACAGCCGTCGGGAAAGAATTTAACAAGCTGGAAAAGAGAATGGAAAACCTGAATAAGAAGGTCG ACGATGGTTTTCTCGATATATGGACTTACAATGCTGAACTTCTGGTCCTTCTGGAAAACGAACGAACCTTGGATTTCCATGACTCTA ATGTGAAGAACTTGTACGAAAAGGTGAAATCACAGTTAAAGAACAACGCCAAAGAAATAGGAAACGGGTGCTTCGAATTTTACCAT AAATGTGATAACGAGTGTATGGAGAGCGTTCGTAATGGTACGTACGACTATCCTAAATATAGTGAAGAGAGCAAACTCAACCGTGA AAAAGTAGACGGCGTGAAACTGGAATCCATGGGGATCTACCAAATCCTGGCCATCTACTCGACGGTGGCAAGTTCTCTCGTGCTT CTTGTAAGCCTGGGTGCTATCTCTTTTTGGATGTGTAGTAACGGATCCCTGCAGTGTCGAATTTGCATT
2707	ATGAAAGCAAATCTTTTAGTTCTTCTGTGTGCGTTGGCTGCAGCCGACGCAGACACTATCTGCATTGGCTACCATGCTAATAATAG CACTGACACTGTAGATACAGTGTTGGAAAAGAACGTTACAGTGACTCACTCTGTCAATCTGCTGGAGGATTCCCATAACGGCAAAT TGTGCCGTCTAAAGGGCATTGCGCCACTCCAGCTGGGAAAGTGCAATATCGCAGGCTGGCTCCTGGGCAACCCAGAATGCGACC CGCTGCTGCCTGTTCGCTCATGGTCATACATCGTCGAAACCCCTAACAGTGAGAACGGGATTTGCTATCCCGGGGATTTCATCGAT TACGAAGAATTACGCGAGCAACTGAGTTCCGTGTCGTCTTTTGAGAGGTTTGAGATCTTCCCGAAGGAATCTTCATGGCCCAACCA TAACACCAATGGTGTGACTGCTGCCTGTTCACACGAGGGCAAAAGCTCTTTCTACCGAAATCTACTGTGGCTCACAGAGAAAGAG GGATCGTACCCCAAGCTGAAAAATTCCTATGTGAACAAGAAAGGCAAAGAGGTGCTGGTGCTTTGGGGAATTCATCACCCACCCA ATTCTAAGGAGCAGCAGAATCTGTATCAGAACGAAAATGCCTATGTTTCCGTGGTAACATCTAATTATAATAGGCGATTCACACCTG AGATTGCAGAGCGGCCGAAAGTCCGAGACCAGGCAGGCCGCATGAACTATTACTGGACGCTTCTCAAACCAGGCGACACCATCA TCTTCGAGGCCAATGGGAACTTGATCGCACCAATGTACGCGTTTGCCTTGAGTCGGGGCTTTGGGAGCGGTATTATCACCTCGAA CGCCTCTATGCATGAATGTAATACGAAGTGTCAGACCCCTCTGGGGGCAATAAATAGCTCACTGCCCTATCAGAACATCCACCCCG TGACCATCGGTGAGTGTCCCAAATACGTCAGGTCAGCTAAACTCCGTATGGTGACCGGCCTAAGGAATAATCCGTCCATTCAGTC CAGAGGTCTGTTCGGTGCAATCGCCGGGTTCATTGAGGGAGGATGGACGGGTATGATCGACGGCTGGTATGGCTATCACCACCA GAACGAACAGGGTAGTGGCTACGCCGCTGATCAAAAGTCCACTCAAAATGCTATCAACGGAATCACGAACAAAGTGAACACAGTG ATAGAGAAGATGAACATTCAGTTCACCGCTGTGGGTAAAGAGTTTAATAAGCTGGAGAAAAGGATGGAAAATCTCAATAAGAAAGT TGACGATGGATTCCTTGACATTTGGACTTACAATGCCGAGCTGCTGGTGCTTCTGGAGAATGAGAGGACCTTAGACTTTCACGACA GCAACGTTAAAAATCTCTATGAGAAAGTGAAATCACAGCTGAAGAATAATGCAAAAGAGATTGGGAATGGGTGCTTCGAGTTTTAT CACAAATGCGATAACGAATGTATGGAGTCCGTGAGAAATGGAACGTACGACTATCCTAAGTATAGCGAGGAAAGCAAACTCAACAG GGAAAAGGTGGACGGTGTTAAACTCGAATCCATGGGGATTTATCAGATATTAGCGATCTATTCGACCGTGGCATCATCTCTCGTTC TTCTTGTGTCTTTGGGCGCCATCAGCTTTTGGATGTGTTCAAATGGCTCTCTGCAGTGTCGCATCTGTATT

2708	ATGAAAGCCAATCTACTGGTCCTCCTCTGCGCCTTGGCTGCCGCAGATGCTGACACGATCTGTATTGGCTACCATGCGAACAATA GCACGGATACAGTGGACACCGTTCTGGAGAAAAATGTCACCGTGACACATAGCGTAAACCTGCTCGAAGATTCCCACAACGGTAA GCTCTGTAGACTCAAAGGTATTGCTCCCCTGCAGTTGGGCAAATGTAATATAGCCGGCTGGCTTCTCGGCAATCCCGAATGCGAT CCCCTGTTGCCCGTGCGGTCTTGGTCTTATATCGTCGAAACCCCAAATTCAGAAAACGGAATTTGTTATCCTGGCGACTTCATCGA TTACGAAGAGTTGAGGGAGCAGCTTTCAAGCGTTAGCAGTTTCGAGAGATTTGAGATATTTCCAAAGGAAAGTTCCTGGCCCAATC ATAACACCAACGGCGTGACCGCCGCTTGCTCTCACGAAGGCAAAAGTAGTTTCTATCGAAATCTGCTCTGGCTGACGGAGAAGGA AGGAAGTTACCCCAAGCTGAAAAACTCCTACGTCAACAAGAAGGGCAAAGAAGTGCTCGTACTGTGGGGCATCCATCACCCTCCC AATTCCAAAGAGCAGCAGAACCTGTATCAGAATGAGAACGCTTATGTATCCGTTGTGACTTCAAACTATAATCGCCGGTTCACTCCT GAGATCGCAGAGCGGCCCAAAGTGAGAGACCAGGCCGGGAGGATGAATTATTATTGGACACTCCTCAAGCCAGGCGACACAATT ATTTTTGAGGCTAACGGGAACTTGATCGCTCCCATGTATGCGTTCGCCTTGTCAAGAGGATTCGGATCCGGGATTATCACCTCTAA TGCTTCGATGCACGAATGTAACACAAAGTGTCAAACTCCACTGGGAGCCATAAATTCGTCCTTGCCATATCAGAATATTCACCCTGT TACAATTGGCGAGTGTCCCAAGTATGTCCGCTCTGCCAAGTTACGGATGGTGACCGGGTTGCGCAACAACCCAAGTATACAGTCA CGCGGACTGTTCGGAGCTATCGCGGGATTCATTGAGGGAGGCTGGACAGGGATGATCGATGGGTGGTACGGGTATCATCACCAG AACGAACAGGGATCGGGATATGCCGCGGACCAAAAGTCAACACAAAACGCCATCAACGGTATTACCAATAAGGTCAACACAGTGA TCGAAAAGATGAATATCCAGTTCACTGCTGTGGGCAAGGAGTTTAATAAACTGGAGAAGCGCATGGAGAACCTGAACAAGAAGGTT GACGACGGGTTCCTGGACATATGGACGTACAATGCCGAGCTGCTTGTCCTCCTGGAGAATGAGAGAACCCTGGATTTTCACGATT CTAATGTGAAGAATCTGTATGAAAAAGTGAAGTCCCAACTCAAAAACAACGCAAAAGAGATCGGCAACGGATGCTTTGAATTCTATC ATAAGTGCGATAATGAATGTATGGAAAGTGTGAGGAATGGGACTTACGATTACCCGAAATATTCCGAAGAGAGCAAACTGAATCGC GAGAAAGTTGACGGTGTAAAGCTGGAGTCTATGGGAATTTACCAAATTCTGGCCATTTATTCTACCGTGGCATCAAGCCTGGTTCT GCTGGTGTCCCTTGGCGCCATTTCATTCTGGATGTGTTCCAATGGCAGCCTCCAGTGCCGGATATGTATC
2709	ATGAAGGCTAACCTTTTGGTCCTTTTGTGCGCGCTTGCCGCGGCGGATGCAGACACCATTTGCATAGGATACCACGCCAATAATAGCACA GACACAGTGGATACAGTCCTTGAGAAAAATGTCACTGTAACTCATAGTGTTAACTTGCTAGAAGACTCTCATAATGGTAAATTATGTCGGC TTAAGGGAATCGCGCCACTCCAATTGGGCAAATGCAACATTGCTGGCTGGTTGTTAGGTAATCCCGAGTGCGATCCCCTGCTCCCTGTCC GGTCTTGGTCTTACATCGTTGAAACACCCAACAGTGAGAATGGCATCTGTTACCCCGGAGACTTCATCGACTACGAAGAGCTGCGAGAG CAGCTTTCTTCCGTCTCCTCCTTTGAACGATTCGAAATCTTCCCCAAAGAGAGCTCATGGCCGAACCATAACACTAATGGAGTTACAGCCG CCTGCTCCCATGAGGGGAAAAGCAGTTTTTATAGGAACCTGCTGTGGCTGACAGAAAAGGAGGGCAGCTACCCCAAGCTGAAGAATTCT TACGTGAACAAGAAGGGCAAAGAAGTATTGGTGCTGTGGGGCATTCATCATCCTCCGAACTCCAAGGAGCAGCAGAACCTTTACCAGAA CGAGAACGCCTACGTATCAGTTGTTACTAGCAACTATAACAGACGATTCACTCCTGAAATCGCCGAGCGGCCTAAGGTTCGGGATCAGG CTGGGAGAATGAACTATTACTGGACCCTGTTGAAGCCTGGAGACACTATTATCTTTGAGGCAAATGGAAACCTGATCGCACCTATGTATG CTTTCGCACTCAGCCGGGGCTTTGGCTCCGGGATCATCACTAGTAATGCTAGCATGCATGAGTGTAACACGAAATGTCAGACCCCACTG GGTGCTATAAACAGTTCCCTGCCTTATCAGAACATCCACCCTGTGACTATTGGAGAGTGTCCCAAGTACGTCAGGAGCGCAAAGCTCCG GATGGTCACAGGGCTGAGGAATAACCCAAGCATTCAGAGCCGGGGTCTATTTGGGGCCATCGCCGGATTCATCGAGGGAGGCTGGACT GGCATGATCGATGGCTGGTATGGCTATCATCACCAAAATGAACAGGGATCCGGCTACGCCGCAGACCAGAAATCTACACAGAATGCAAT CAACGGAATCACAAATAAAGTGAATACCGTCATCGAGAAGATGAACATACAATTTACCGCTGTGGGTAAAGAGTTTAATAAACTAGAGAAA CGGATGGAAAATCTTAATAAAAAGGTGGACGACGGGTTCCTCGATATTTGGACATATAATGCCGAGCTATTGGTGCTCCTGGAAAATGAA AGAACGCTTGATTTTCACGACTCTAACGTGAAAAATCTGTACGAAAAGGTTAAGAGTCAGCTAAAGAACAATGCCAAAGAAATCGGAAATG GTTGCTTCGAGTTCTACCACAAATGTGATAATGAATGCATGGAATCAGTCCGAAACGGTACCTACGATTATCCTAAATATTCTGAAGAATC AAAGCTAAACCGAGAGAAGGTCGATGGCGTCAAGCTTGAGAGCATGGGTATCTACCAGATACTGGCCATCTATAGTACTGTGGCATCTTC CCTGGTGCTTCTCGTGTCCCTGGGGGCGATCTCCTTTTGGATGTGTTCAAACGGATCCCTGCAGTGTCGGATATGCATC

2710	ATGAAGGCCAACCTGCTGGTTCTGCTTTGCGCTCTCGCCGCCGCCGATGCTGATACTATTTGCATAGGTTATCATGCTAACAACTCCACT GATACTGTTGATACAGTTCTCGAAAAAAACGTAACGGTCACTCACTCTGTAAACCTATTGGAAGACAGCCACAATGGGAAGTTGTGTAGG CTTAAAGGCATCGCTCCACTGCAGCTGGGCAAATGTAACATCGCAGGCTGGCTCCTGGGCAACCCGGAGTGTGATCCTTTACTGCCCGT CCGCTCCTGGTCTTACATTGTTGAGACACCCAACAGTGAAAACGGTATCTGTTATCCCGGCGATTTTATTGACTACGAGGAGTTGCGGGA ACAACTGTCTTCTGTGAGCAGTTTCGAGCGCTTCGAGATTTTTCCTAAGGAGTCTTCTTGGCCCAACCATAACACAAACGGGGCTGACCGC TGCCTGCTCACACGAGGGCAAGAGTAGCTTCTATAGAAATCTTCTCTGGCTTACTGAAAAAGAGGGATCTTATCCAAAGTTGAAAAACAGT TATGTGAACAAGAAGGGAAAAGAAGTGCTGGTCCTTTGGGGGATCCATCACCCACCCAACTCCAAGGAGCAGCAGAACCTGTACCAAAA CGAGAACGCATATGTTAGCGTCGTGACCAGCAACTACAATCGCAGGTTCACTCCCGAGATCGCCGAGAGGCCTAAAGTTCGCGACCAGG CTGGACGGATGAACTACTATTGGACACTATTAAAGCCAGGCGATACCATAATATTTGAAGCAAATGGGAACTTAATTGCACCGATGTACGC CTTCGCACTGTCCCGGGGGTTCGGAAGCGGGATTATTACCTCTAACGCCTCCATGCATGAGTGCAACACCAAGTGTCAAACACCCCTGG GAGCTATTAACTCTAGCTTGCCCTATCAGAACATCCACCCAGTCACTATCGGCGAGTGTCCTAAGTACGTTAGATCAGCCAAGCTGCGCA TGGTTACGGGGCTTCGGAACAACCCCTCAATCCAATCTCGGGGGCTTTTCGGTGCAATCGCCGGCTTTATAGAAGGCGGTTGGACCGGA ATGATTGATGGCTGGTATGGCTACCACCACCAAAACGAACAGGGAAGCGGGTATGCTGCAGACCAGAAGAGCACACAGAATGCTATCAA TGGGATCACCAACAAAGTAAACACGGTGATTGAGAAAATGAATATTCAGTTCACAGCGGTGGGCAAGGAGTTCAACAAACTGGAGAAGC GTATGGAAAATCTGAATAAAAAAGTTGACGACGGTTTCCTAGACATCTGGACATATAACGCCGAGTTGTTGGTGCTCCTCGAAAACGAGC GCACACTCGACTTCCACGATTCTAACGTTAAGAATCTTTACGAAAAAGTTAAATCGCAGCTGAAAAACAACGCGAAGGAGATCGGTAACG GCTGTTTCGAGTTTTATCATAAGTGTGATAATGAATGCATGGAATCCGTGCGGAACGGTACCTACGATTATCCTAAATATTCAGAGGAGAG TAAACTCAATCGCCAGAAGGTCGATGGAGTCAAGCTTGAGAGTATGGGAATTTATCAGATTCTGGCAATCTATAGCACCGTCGCATCCAG TCTCGTCCTCCTTGTTTCACTGGGAGCTATTTCATTCTGGATGTGCTCTAACGGGTCGCTGCAGTGCAGAATCTGCATT
2711	ATGAAGGCAAATCTTCTGGTACTTCTCTGCGCCCTCGCTGCAGCCGACGCAGATACCATTTGTATCGGCTACCACGCGAATAACTCAACG GATACCGTGGACACCGTGTTGGAGAAGAACGTTACCGTCACACATAGTGTGAATCTGCTTGAAGATTCACATAACGGAAAACTTTGTAGG CTCAAAGGCATCGCGCCACTGCAGCTGGGGAAATGCAATATTGCAGGTTGGTTACTGGGTAACCCAGAATGCGATCCTCTGTTACCTGT GCGTAGTTGGAGCTATATCGTTGAGACCCCAAATTCGGAGAATGGGATCTGCTATCCAGGGGACTTCATTGACTATGAAGAGCTGCGAG AACAGCTCAGCAGCGTATCTAGCTTCGAGAGATTTGAAATTTTCCCGAAGGAATCCAGCTGGCCAAACCACAATACCAATGGCGTTACGG CCGCCTGTAGCCACGAAGGTAAGTCTAGCTTCTACAGAAACCTGCTGTGGCTGACAGAAAAGGAGGGGTCTTACCCGAAGTTGAAGAAT AGCTACGTCAATAAGAAAGGAAAAGAGGTCCTTGTCCTCTGGGGCATCCACCACCCTCCAAACTCGAAGGAACAGCAGAACCTTTATCAG AACGAGAACGCTTATGTCTCCGTGGTGACATCCAATTATAACAGGCGTTTCACACCAGAGATAGCCGAAAGGCCCAAGGTCCGCGACCA AGCTGGTAGGATGAATTACTATTGGACACTCCTCAAACCAGGAGACACAATTATTTTTGAAGCAAATGGGAACCTGATTGCCCCCATGTAT GCCTTTGCGCTAAGCCGCGGTTTCGGAAGCGGCATTATCACGAGTAATGCATCAATGCACGAGTGTAACACCAAATGTCAAACCCCCTTG GGCGCAATTAACTCTTCCCTCCCTTACCAGAACATCCACCCCGTCACCATCGGTGAGTGCCCTAAATACGTACGGTCTGCTAAGTTGCGG ATGGTCACCGGCCTGCGAAATAACCCTAGCATTCAGTCCAGAGGGTTATTTGGCGCGATCGCAGGATTCATCGAGGGAGGGTGGACCG GCATGATAGACGGGTGGTATGGATACCATCATCAGAACGAGCAAGGCAGCGGCTACGCTGCTGACCAAAAGAGCACGCAGAACGCGAT CAATGGAATCACGAATAAGGTTAACACGGTCATCGAAAAGATGAACATACAGTTCACAGCTGTGGGCAAGGAGTTCAACAAGCTTGAAAA AAGAATGGAGAACTTGAACAAAAAAGTAGACGACGGCTTTCTCGATATTTGGACTTACAACGCCGAATTGCTAGTCTTGTTGGAGAATGAA CGCACCTTGGACTTCCACGACTCCAACGTCAAGAATTTATACGAAAAGGTCAAAAGCCAGCTTAAGAATAACGCCAAGGAGATCGGCAAT GGCTGTTTCGAATTTTACCATAAATGCGATAATGAATGTATGGAAAGTGTTAGGAACGGCACATACGACTACCCTAAATACTCCGAAGAGA GCAAACTCAATCGGGAGAAGGTTGATGGAGTGAAGTTAGAGAGCATGGGAATTTACCAAATACTCGCCATCTATAGCACCGTTGCCTCAA GCCTGGTCCTCCTAGTGTCCCTGGGCGCCATTAGTTTCTGGATGTGTTCCAATGGGTCTTTGCAATGTCGCATCTGTATA
2712	ATGAAAGCTAACCTGCTGGTCCTCCTATGTGCTCTCGCTGCCGCAGATGCCGATACCATATGCATTGGATATCACGCGAACAACAGCACC GATACGGTGGATACAGTGCTGGAAAAAAACGTGACTGTGACACATTCTGTCAATCTCCTGGAAGACTCCCACAATGGCAAGCTGTGTCGT TTCAAGGGGATCGCCCCCCTGCAGCTCGGAAAGTGCAACATAGCAGGGTGGCTGCTCGGGAATCCAGAGTGCGATCCGCTGCTCCCGG TACGGAGCTGGTCTTACATTGTTGAGACACCTAACAGCGAGAATGGCATTTGTTATCCTGGAGACTTCATCGACTACGAGGAACTCCGGG AACAGCTTAGCTCCGTAAGCAGCTTCGAAAGATTTGAGATTTTCCCTAAGGAGTCATCCTGGCCCAACCATAATACAAACGGAGTGACCG CTGCCTGTTCTCATGAAGGGAAAAGTTCCTTTTACAGAAATCTCCTGTGGCTGACTGAGAAAGAAGGAAGCTATCCTAAGCTGAAGAATTC TTACGTTAATAAGAAAGGAAAAGAAGTGCTGGTCCTCTGGGGGATACATCACCCCCAAATAGCAAGGAGCAACAGAATTTATATCAGAA TGAGAACGCATACGTCTCGGTTGTGACTTCGAACTACAACAGGCGGTTTACACCCGAGATCGCAGAGCGCCCCAAGGTTAGAGACCAGG CTGGGAGGATGAATTACTATTGGACGCTTCTCAAGCCAGGGGATACCATTATTTTTGAGGCTAATGGCAATCTGATTGCCCCTATGTACG CCTTCGCTCTGTCAAGGGGATTTGGAAGCGGCATCATCACCTCCAACGCTTCCATGCATGAGTGCAACACCAAGTGTCAGACACCCTTAG GCGCAATCAACAGCAGTCTGCCCTATCAAAACATTCATCCCGTGACAATCGGCGAGTGTCCGAAATACGTTCGATCCGCGAAGTTAAGGA TGGTTACGGGCCTACGTAATAACCCTTCAATCCAGTCTCGTGGCCTCTTCGGTGCTATCGCTGGTTTCATCGAAGGAGGCTGGACAGGA ATGATCGACGGATGGTATGGATATCACCATCAGAATGAGCAGGGCAGTGGGTATGCCGCAGACCAGAAATCCACCCAGAATGCCATAAA TGGAATCACCAACAAGGTGAACACAGTTATCGAAAAGATGAACATCCAGTTCACCGCCGTGGGAAAGGAGTTTAACAAACTTGAGAAGAG AATGGAGAATCTTAATAAAAAGGTAGATGATGGGTTCCTTGATATCTGGACATATAACGCCGAGCTTCTAGTGCTCCTGGAGAACGAGAG GACTCTGGACTTTCACGACTCCAACGTCAAAAACCTATACGAAAAGGTCAAGAGTCAACTGAAAAATAACGCAAAAGAGATAGGTAACGG ATGTTTTGAGTTCTACCACAAATGCGATAACGAGTGCATGGAATCCGTGAGGAATGGTACCTATGACTATCCAAAATATTCTGAGGAGTCC AAGCTCAACCGAGAAAAGGTGGACGGCGTAAAGCTCGAAAGCATGGGGATTTATCAGATACTTGCAATTTATTCCACTGTGGCTTCTTCC CTGGTGCTTTTGGTCAGCTTAGGGGCTATCAGCTTCTGGATGTGTTCCAACGGCTCCCTTCAGTGTAGAATTTGCATC

2713	ATGAAGGCAAACTTGTTGGTGCTACTCTGCGCATTAGCAGCTGCTGACGCAGATACAATTTGTATCGGGTACCACGCCAATAACTCTACA GATACAGTAGATACCGTCCTTGAAAAAAATGTCACCGTGACCCATAGTGTAAACCTCCTAGAAGATAGCCATAACGGCAAACTGTGTCGG CTTAAGGGCATCGCGCCTCTACAGCTGGGCAAGTGTAACATTGCTGGCTGGCTCCTTGGCAATCCTGAGTGCGACCCCTTGCTGCCAGT GAGGTCCTGGAGCTACATCGTGGAAACTCCCAATTCTGAGAATGGAATATGTTATCCGGGGGATTTCATCGACTACGAAGAGCTGCGCG AACAACTCTCATCAGTCAGCTCCTTTGAAAGATTCGAAATCTTCCCCAAGGAATCTAGCTGGCCAAACCATAACACTAATGGAGTAACAGC TGCATGTTCCCACGAGGGCAAGAGCTCTTTTTACAGGAACCTCCTATGGCTGACGGAGAAAGAAGGGTCCTACCCCAAGCTGAAAAATA GCTACGTTAACAAGAAAGGAAAAGAGGTGTTGGTTCTGTGGGGAATTCATCATCCTCCCAACAGTAAGGAGCAGCAGAATCTCTATCAAA ATGAAAACGCTTACGTCAGTGTGGTGACGTCTAATTACAACAGGAGATTTACACCAGAGATTGCCGAGCGGCCGAAAGTGAGAGATCAG GCAGGCAGGATGAATTATTACTGGACACTTCTGAAACCTGGAGACACAATCATTTTCGAAGCTAATGGAAACCTCATCGCACCGATGTAC GCATTCGCCTTGTCCCGGGGATTCGGCTCCGGCATCATCACCTCCAACGCTTCAATGCATGAATGTAATACTAAATGCCAGACGCCGCTT GGCGCTATTAACTCATCCTTGCCTTATCAGAATATTCATCCCGTTACCATTGGAGAGTGTCCCAAATACGTGCGTTCAGCCAAACTGCGGA TGGTCACTGGCCTGCGAAATAACCCTTCAATTCAATCTCGCGGACTGTTCGGAGCTATCGCCGGCTTCATAGAAGGGGGCTGGACCGGC ATGATTGACGGATGGTACGGTTACCATCACCAAAACGAGCAGGGTTCCGGCTATGCCGCCGACCAGAAATCAACCCAGAACGCCATAAA TGGTATTACCAACAAGGTTAATACGGTCATCGAGAAAATGAACATCCAGTTCACAGCAGTGGGAAAAGAATTTAATAAGCTCGAGAAAAGA ATGGAAAACCTGAACAAAAAAGTGGACGATGGTTTCCTCGACATATGGACCTATAATGCTGAGCTACTCGTCCTTCTGGAAAATGAGCGA ACGCTTGACTTCCACGATTCGAATGTGAAAAACCTATACGAAAAAGTGAAGTCCCAGCTCAAGAACAATGCGAAGGAGATCGGGAACGG CTGCTTTGAGTTTTACCACAAATGTGATAACGAATGCATGGAGTCTGTGCGAAATGGGACATACGACTATCCTAAGTATTCAGAAGAGTCA AAACTCAACCGGGAAAAGCTTGACGGAGTTAAGCTTGAGTCAATGGGAATTTACCAAATCCTGGCAATCTACTCTACTGTTGCCTCTTCAC TCGTCCTCTTAGTTTCACTAGGGGCTATCAGTTTCTGGATGTGTTCTAATGGTAGCCTGCAGTGTCGAATTTGTATA
2714	ATGAAAGCTAATCTCTTAGTTCTGCTCTGCGCCTTAGCGGCTGCAGATGCCGACACCATATGCATTGGCTACCACGCTAATAACTCCACC GATACAGTGGACACCGTGCTGGAAAAGAACGTCACTGTGACCCACTCAGTGAACCTGTTGGAGGATTCACACAACGGCAAGTTATGCAG ACTCAAGGGAATTGCACCACTGCAGCTGGGTAAGTGTAATATCGCTGGCTGGCTTCTCGGAAATCCCGAGTGCGACCCTCTGTTACCTG TCAGGAGCTGGAGTTATATCGTGGAAACTCCTAATAGTGAAAATGGAATATGCTATCCCGGCGACTTCATCGACTACGAGGAGCTAAGGG AACAGCTGTCAAGCGTCTCCTCCTTCGAAAGATTCGAGATCTTCCCGAAGGAGAGTTCGTGGCCAAACCATAACACTAACGGGGTCACC GCGGCCTGTTCTCACGAGGGGAAGTCATCCTTCTACCGAAACCTTCTGTGGCTGACGGAGAAAGAAGGCAGCTATCCCAAACTGAAGAA TAGTTACGTGAATAAGAAGGGAAAGGAGGTACTGGTCCTCTGGGGCATCCATCATCCCCCCAACAGTAAGGAGCAGCAGAATCTTTACC AGAACGAGAACGCTTATGTGAGCGTCGTAACCTCAAACTACAACCGCAGGTTCACTCCTGAAATTGCCGAACGGCCGAAAGTGAGGGAC CAAGCCGGCCGCATGAACTACTATTGGACGCTTCTCAAGCCCGGTGATACTATTATTTTCGAGGCAAACGGCAACTTAATCGCTCCTATG TACGCATTTGCACTGAGTCGGGGTTTTGGTTCGGGCATCATCACCTCAAATGCTTCCATGCACGAATGCAATACAAAATGCCAGACCCCA CTTGGAGCCATCAACTCCTCACTGCCTTATCAGAATATCCACCCAGTCACTATCGGGGAATGCCCGAAATATGTCAGAAGCGCTAAGCTT CGGATGGTGACCGGTCTGCGCAATAATCCGTCTATACAATCCAGGGGACTGTTTGGCGCGATAGCCGGATTTATTGAGGGGGGATGGAC TGGCATGATCGATGGGTGGTACGGGTACCATCACCAGAATGAACAGGGTTCCGGCTACGCCGCAGATCAAAAGTCCACCCAGAACGCTA TTAACGGAATCACCAACAAGGTCAATACTGTCATTGAGAAAATGAATATCCAGTTCACTGCCGTGGGCAAGGAGTTCAATAAGCTAGAGA AGCGTATGGAAAACCTTAACAAGAAGGTGGACGATGGATTTTTGGATATTTGGACCTATAATGCTGAGTTGTTGGTCCTCCTGGAGAATG AAAGAACTCTGGACTTCCATGATAGCAATGTGAAAAACCTCTACGAAAAAGTCAAATCACAGCTTAAGAACAACGCCAAAGAAATTGGGAA TGGATGCTTTGAGTTTTACCATAAATGCGACAACGAGTGTATGGAGTCTGTCAGAAACGCCACATACGATTACCCCAAGTATTCCGAAGA ATCAAAACTGAATCGGGAAAAGGTTGATGGAGTGAAATTAGAAAGCATGGGAATCTATCAAATTCTTGCAATTTATAGCACCGTTGCTAGC TCACTGGTCCTGCTCGTTAGTTTGGGCGCCATTAGCTTCTGGATGTGTAGCAATGGGTCTCTCCAATGCAGAATTTGCATT
2715	ATGAAAGCGAATCTGCTGGTCCTGCTGTGCGCCCTGGCGGCTGCTGACGCGGATACTATTTGCATCGGCTATCACGCTAATAACTCGAC AGACACCGTGGATACAGTCTTGGAGAAAAATGTGACTGTAACCCACTCGGTTAACCTTCTCGAGGACTCGCATAACGGGAAACTGTGTAG GCTGAAGGCAATCGCACCACTCCAGCTGGGCAAGTGCAATATAGCTGCCTGGCTGCTCGGGAATCCTGAGTGCGACCCTCTCCTGCCC GTCCGGTCCTGGAGCTATATAGTGGAAACACCTAACAGCGAGAATGGTATCTGCTATCCGGGGGATTTTATCGACTATGAGGAACTACGA GAACAGCTCTCCTCCGTTAGCAGCTTCGAGCGATTTGAAATCTTTCCTAAGGAGAGCAGTTGGCCTAACCACAACACTAATGGGGTGACA GCCGCCTGTAGCCATGAGGGAAAGTCTTCTTTTTATAGAAACTTGCTGTGGTTGACTGAGAAAGAGGGGAGCTACCCGAAACTAAAGAAC TCCTACGTGAACAAGAAAGGCAAAGAGGTTTTGGTTCTGTGGGGCATCCACCATCCTCCCAATAGCAAGGAACAGCAGAATCTCTATCAG AACGAAAACGCCTACGTGTCAGTGGTGACCAGTAATTATAATCGACGCTTTACCCCCGAAATTGCAGAGCGGCCGAAAGTCCGCGACCA GGCCGGACGCATGAATTACTATTGGACCCTGCTAAAGCCCGGGGACACTATCATATTTGAAGCCAACGGGAACCTGATTGCACCCATGT ATGCGTTTGCTCTGTCACGCGGGTTTGGTAGCGGGATTATCACTTCCAACGCTTCAATGCATGAATGTAACACTAAGTGTCAGACCCCAC TGGGCGCCATTAACAGCTCGCTGCCATACCAGAATATACACCCAGTTACTATTGGTGAATGTCCTAAGTATGTACGGTCCGCCAAGCTGC GGATGGTTACTGGCCTCAGGAACAATCCTTCAATTCAGTCTAGGGGGCTGTTCGGAGCCATAGCTGGCTTTATTGAGGGAGGCTGGACA GGCATGATTGACGGCTGGTACGGCTACCACCATCAGAACGAGCAAGGCTCCGGCTACGCCGCGGACCAGAAGTCAACTCAAAATGCGA TAAACGGAATCACCAACAAGGTGAACACCGTCATAGAAAAAATGAATATACAATTCACAGCCGTGGGAAAGGAGTTCAATAAGCTCGAGA AGCGTATGGAGAATCTCAATAAAAAAGTGGATGATGGGTTCCTCGATATCTGGACCTATAACGCTGAACTCTTAGTCCTCCTCGAAAACGA GCGAACGCTCGACTTCCATGATTCTAATGTAAAAAATCTTTACGAAAAGGTTAAGTCACAGCTGAAGAACAACGCCAACGAGATAGGAAA CGGCTGTTTCGAGTTTTATCATAAGTGTGACAACGAGTGTATGGAATCTGTCCGCAATGGCACTTATGATTACCCCAAGTACTCCGAGGA AAGTAAACTTAATAGAGAAAAAGTTGATGGCGTTAAGCTGGAGTCAATGGGGATTTATCAGATCCTAGCTATATATTCTACTGTGGCTTCT AGTCTCGTTTTGCTGGTGAGCCTGGGAGCCATAAGCTTTTGGATGTGTAGCAATGGGTCCCTCCAGTGTCGAATCTGTATC

2716	ATGAAAGCCAACTTGCTGGTGTTGCTCTGCGCCCTTGCCGCCGCCGACGCAGATACCATTTGTATTGGCTACCATGCCAATAATTCTACG GACACTGTGGATACCGTCTTGGAGAAAAATGTAACTGTAACTCACTCCGTTAATCTTCTCGAAGATTCACACAATGGCAAGCTGTGCCGC CTTAAAGGCATTGCCCCCTTGCAGCTGGGCAAGTGTAATATTGCTGGATGGCTCCTCGGCAATCCTGAGTGTGACCCCCTCCTGCCCGT TAGGTCGTGGAGCTACATTGTGGAGACCCCAAATTCGGAAAATGGGATATGTTACCCAGGCGACTTTATCGACTATGAGGAACTACGTGA GCAGCTGAGTTCTGTTTCCAGTTTTGAAAGATTTGAGATCTTTCCAAAGGAGAGCTCTTGGCCGAACCATAACACCAATGGGCTCACCGC CGCTTGCAGTCATGAAGGAAAATCAAGCTTCTACCGAAACTTGCTGTGGCTCACGGAAAAGGAGGGCTCCTACCCCAAGTTGAAGAACT CCTATGTGAACAAAAAGGGCAAAGAGGTTCTCGTGCTCTGGGGAATTCACCACCCTCCCAACAGCAAGGAGCAGCAGAACTTGTATCAG AACGAAAATGCTTACGTTAGCGTGGTGACTTCCAATTATAATCGACGCTTTACACCGGAGATTGCTGAACGCCCGAAAGTTCGGGATCAA GCCGGCCGGATGAACTATTATTGGACCCTCCTGAAACCCGGTGATACTATTATTTTTGAGGCAAACGGAAACCTGATTGCTCCCATGTAC GCCTTCGCCCTGTCCCGGGGGTTTGGCAGCGGAATTATAACTAGCAACGCTAGCATGCACGAGTGTAATACTAAGTGTCAGACCCCTCT GGGGGCAATAAACTCCAGCTTACCTTATCAGAACATCCACCCTGTTACCATAGGCGAGTGCCCGAAGTACGTTAGGTCAGCTAAGCTAAG AATGGTCACTGGTTTGAGGAATAACCCCAGCATTCAGAGTAGGGGGCTGTTTGGGGCAATTGCAGGTTTTATCGAGGGCGGTTGGACTG GTATGATCGACGGCTGGTATGGATATCATCATCAGAATGAGCAGGGGTCTGGGTATGCTGCGGACCAGAAGTCCACGCAAAATGCCATA AACGGAATCACGAATAAGGTTAATACCGTCATCGAAAAAATGAACATTCAGTTTACTGCGGTGGGTAAGGAATTCAATAAACTCGAAAAGA GAATGGAGAACTTAAATAAGAAAGTCGACGACGGATTTTTGGATATCTGGACCTATAACGCCGAACTCCTCGTTCTGTTGGAGAACGAGC GCACCCTCGACTTCCACGATAGTAACGTGAAAAATCTGTATGAGAAGGTGAAGTCCCAGCTCAAGAACAATGCCAAGGAAATCGGAAACG GATGTTTTGAGTTCTACCACAAGTGCGACAATGAATGCATGGAGAGTGTGCGCAATGGGACGTACGACTACCCTAAATATAGTGAAGAGT CTAAGCTCAACCGGGAGAACGTGGACGGAGTGAAGCTGGAGAGTATGGGAATTTACCAGATACTGGCTATCTACAGTACAGTTGCCAGC TCCCTCGTTCTTCTGGTGTCTCTGGGGGCCATCTCTTTCTGGATGTGTTCCAATGGCAGCCTCCAATGCCGGATCTGCATT
2717	ATGAAAGCAAATCTTCTGGTGTTACTATGCGCATTAGCTGCCGCTGACGCCGACACCATTTGCATTGGCTATCACGCCAATAATAGCACT GACACTGTCGACACAGTTCTTGAAAAGAACGTGACTGTTACCCACTCCGTGAATCTACTCGAGGATTCTCATAACGGAAAGCTCTGTCGC CTTAAGGGTATTGCCCCTCTCCAGCTTGGAAAATGCAACATTGCTGGCTGGCTGCTTGGAAACCCTGAATGTGACCCACTCCTGCCCGTC CGGAGCTGGAGTTACATAGTGGAGACTCCAAATAGCGAGAACGGTATATGTTACCCAGGTGATTTCATTGACTATGAGGAGTTGAGAGAG CAGTTGAGCTCCGTTTCGTCTTTTGAAAGGTTCGAGATCTTCCCAAAAGAGTCAAGCTGGCCTAACCATAACACCAATGGTGTTACGGCC GCATGTAGCCATGAGGGCAAGTCGTCCTTTTATCGAAATCTCCTATGGCTCACAGAGAAGGAAGGATCTTACCCAAAGCTCAAGAACTCG TACGTAAACAAGAAGGGGAAGGAAGTCCTTGTTCTGTGGGGTATACATCACCCACCAAATTCCAAGGAACAGCAGAATTTATATCAGAAC GAAAACGCATATGTTTCTGTCGTTACATCCAACTATAACCGGAGATTCACCCCAGAGATCGCCGAAAGACCGAAGGTGCGGGACCAGGC AGGGCGAATGAACTATTACTGGACATTGCTCAAGCCCGGAGACACCATAATTTTCGAGGCTAATGGCAACCTCATTGCCCCGATGTACGC ATTCGCTCTTTCTCGGGGATTTGGCAGTGGGATTATCACCTCGAACGCCTCCATGCATGAATGTAATACTAAATGTCAAACACCACTGGG AGCCATTAACAGTAGCCTGCCCTATCAAAATATCCACCCGGTCACTATTGGGGAGTGCCCAAAGTATGTGCGGTCTGCAAAGTTGAGAAT GGTTACCGGGCTGCGCAACAATCCAAGTATTCAGAGTAGGGGGCTATTTGGTGCAATAGCAGGCTTTATCGAAGGGGGCTGGACTGGAA TGATAGATGGCTGGTATGGATATCACCACCAGAACGAGCAGGGTAGCGGCTACGCCGCCGATCAGAAGAGCACTCAGAATGCTATCAAC GGAATCACAAATAAAGTTAATACCGTAATAGAAAAGATGAATATCCAGTTCACAGCTGTGGGTAAAGAGTTCAACAAGCTGGAAAAACGGA TGGAAAACCTGAATAAAAAGGTGGATGATGGTTTCCTCGATATATGGACTTATAACGCTGAGCTTCTGGTTCTGCTGGAAAATGAACGCAC ACTGGACTTCCACGATAGTAATGTAAAAAATCTGTATGAAAAGGTTAAATCACAACTGAAAAATAATGCTAAGGAAATTGGGAACGGGTGT TTCGAATTCTATCACAAGTGCGATAATGAGTGTATGGAGTCAGTTCGGAACGGGACCTACGATTATCCTAAGTATTCAGAAGAGTCTAAAC TGAACCGCGAAAAAGTGGATGGAGTCAAACTGGAGTCGATGGGCATCTATCAGATCCTCGCCATTTATAGCACCGTGGCATCGTCCCTG GTGCTTTTGGTTAGCCTGGGCGCCATTAGTTTCTGGATGTGTAGTAACGGGTCTCTGCAGTGTAGGATTTGTATT
2718	ATGAAGGCAAATCTGCTCGTGTTACTCTGCGCTTTGGCCGCCGCTGACGCCGACACTATTTGCATCGGTTACCACGCCAACAACAGCAC CGATACAGTGGACACGGTATTGGAAAAAAACGTAACTGTGACACATTCAGTAAACCTGCTCGAGGACAGCCACAACGGTAAGCTGTGCC GGTTGAAAGGTATCGCACCACTGCAGCTTGGAAAATGTAATATCGCCGGGTGGCTGCTGGGCAATCCTGAGTGTGATCCTCTGCTCCCT GTACGGAGTTGGTCGTATATTGTGGAAACCCCTAACAGCGAAAACGGGATTTGTTATCCTGGCGACTTTATCGACTATGAGGAGCTCCGG GAGCAGCTGTCCAGCGTAAGTTCATTCGAGAGGTTCGAGATTTTCCCCAAAGAGAGTTCCTGGCCTAACCACAACACAAACGGTGTGACA GCTGCCTGCAGCCACGAGGGGAAGTCCTCATTCTATAGGAATTTGCTCTGGCTGACCGAAAAGGAAGGTTCGTACCCGAAGCTTAAAAA TTCTTACGTTAACAAAAAGGGGAAGGAGGTACTCGTTCTGTGGGGGATCCATCATCCCCCAAATTCTAAGGAACAGCAGAACCTATACCA GAATGAAAACGCATATGTCAGCGTGGTCACATCTAACTACAACAGACGCTTCACACCCGAAATCGCTGAGCGCCCCAAGGTACGAGATC AGGCTGGCAGGATGAACTACTATTGGACTTTACTGAAGCCGGCGATACAATAATTTTCGAGGCTAACGGAAATCTGATCGCCCCAATGT ATGCCTTTGCTCTTTCCCGCGGCTTTGGTTCAGGGATTATTACTTCTAACGCTTCGATGCACGAGTGTAATACCAAATGCCAGACCCCGCT GGGCGCCATTAATAGCTCATTGCCTTATCAGAATATCCATCCCGTTACCATTGGCGAATGTCCCAAGTATGTGAGGAGCGCAAAACTTCG GATGGTAACAGGTCTAAGGAATAACCCTAGCATTCAGTCACGGGGGCTGTTTGGGGCAATAGCAGGCTTTATTGAGGGCGGATGGACAG GCATGATCGATGGCTGGTATGGTTACCACCATCAGAACGAGCAGGGCTCTGGATATGCAGCCGACCAGAAGAGTACACAAAATGCAATC AATGGCATTACTAACAAAGTGAACACGGTCATTGAAAAAATGAATATCCAGTTCACAGCCGTGGGAAAGGAGTTTAACAAGCTGGAAAAA CGGATGGAAAATCTAAATAAGAAAGTCGACGACGGTTTTTTGGATATCTGGACCTACAACGCTGAACTACTGGTTCTTCTGGAGAACGAA AGAACCCTGGACTTTCACGACTCTAATGTGAAGAATCTGTACGAAAAAGTGAAGTCTCAGTTGAAGAACAACGCTAAAGAGATTGGGAAT GGCTGTTTCGAATTCTACCACAAATGCGATAATGAGTGCATGGAGTCTGTCAGGAATGGTACTTACGACTATCCCAAGTATTCTGAGGAG AGTAAGCTCAACAGGGAAAAAGTCGACGGAGTCAAACTGGAGTCGATGGGCATATACCAGATTCTAGCCATCTACTCCACCGTGGCCAG TAGTTTAGTCTTGCTCGTAAGCCTTGGCGCTATCTCCTTCTGGATGTGCTCCAATGGATCCCTGCAGTGTAGAATCTGTATC

2719	ATGAAGGCGAATCTACTGGTTTTATTGTGCGCCCTCGCTGCTGCCGACGCCGATACCATCTGCATCGGATACCATGCCAATAACTCCACA GATACCGTGGACACTGTGCTGGAAAAGAATGTTACAGTGACACATTCAGTGAACCTGCTGGAAGATAGTCATAATGGCAAACTGTGTAGA CTAAAGGGCATCGCCCCCCTGCAACTTGGTAAGTGTAACATCGCTGGTTGGCTTCTGGGGAACCCCGAATGCGATCCACTGCTGCCTGT CCGCAGTTGGTCCTATATTGTCGAAACCCCCAACTCTGAAAACGGGATTTGTTATCCAGGAGACTTCATTGACTATGAAGAATTGCGGGA ACAGCTTTCAAGTGTGAGCTCTTTCGAACGGTTTGAAATCTTTCCGAAGGAGTCCTCGTGGCCAAATCATAACACTAATGGCGTGACAGC GGCATGTAGTCACGAAGGAAAATCTTCCTTTTATAGAAACCTCCTGTGGCTGACAGAGAAGGAAGGCAGTTATCCCAAACTGAAGAACAG TTACGTCAATAAAAAGGGGAAAGAAGTACTGGTACTTTGGGGAATCCACCATCCCCCCAACAGCAAGGAGCAGCAGAATCTTTATCAAAA TGAGAACGCCTACGTCAGCGTGGTGACCTCGAATTACAACAGGCGGTTCACACCAGAAATCGCGGAAAGGCCGAAGGTGCGGGATCAG GCCGGCAGGATGAACTACTATTGGACTCTCCTAAAACCAGGCGACACTATTATATTTGAAGCCAACGGCAACCTTATCGCTCCAATGTAC GCCTTCGCTTTGAGCCGTGGGTTTGGCTCCGGAATTATAACATCAAATGCAAGTATGCACGAGTGCAACACAAAGTGCCAGACACCACTC GGAGCAATCAACAGCTCCTTGCCATACCAAAACATTCATCCTGTTACCATCGGCGAATGCCCGAAGTATGTGAGGAGCGCAAAACTGAG GATGGTGACGGGCTTGCGGAACAACCCTTCTATCCAGTCCCGGGGCCTGTTTGGAGCGATCGCAGGATTCATTGAAGGGGGATGGACG GGGATGATCGATGGATGGTACGGATATCATCACCAAAATGAACAAGGGTCAGGTTATGCTGCTGATCAAAAGTCAACACAGAATGCAATC AACGGCATAACCAATAAAGTTAACACAGTGATTGAAAAAATGAATATTCAGTTTACCGCCGTGGGAAAAGAGTTCAATAAACTGGAGAAGA GGATGGAGAATTTGAACAAGAAGGTAGATGACGGCTTTCTCGACATTTGGACTTATAATGCAGAGCTATTAGTGCTCCTCGAGAATGAGA GAACTCTTGACTTTCACGACAGTAACGTAAAAAACCTCTATGAAAAGGTCAAATCACAGCTTAAGAATAATGCAAAAGAGATCGGAAACGG ATGCTTCGAATTTTACCACAAATGCGACAACGAATGTATGGAATCCGTGAGGAACGGTACCTACGATTACCCCAAGTACAGCGAGGAAAG CAAGCTGAATCGAGAAAAAGTTGACGGTGTGAAGCTGGAGTCCATGGGCATCTACCAGATTCTTGCTATCTATTCTACAGTAGCTTCCTC CCTTGTACTTCTGGTGTCCCTGGGTGCCATTTCATTCTGGATGTGCAGCAACGGTTCCTTGCAGTGTCGGATCTGCATA
2720	ATGAAAGCTAATCTTCTCGTGTTGCTGTGCGCCTTAGCGGCCGCAGACGCGGACACAATCTGTATTGGGTACCATGCCAATAATTCCACT GACACCGTAGACACTGTACTGGAAAAGAATGTGACGGTAACCCACTCCGTTAATCTGCTGGAGGACTCTCACAACGGCAAACTCTGTCG GCTGAAAGGTATTGCACCACTTCAGCTGGGGAAGTGTAACATTGCCGGATGGCTACTAGGCAACCCTGAATGTGACCCCCTGCTGCCAG TGCGAAGCTGGAGTTACATTGTGGAAACTCCCAATAGTGAAAACGGTATCTGCTACCCTGGCGATTTCATTGATTATGAAGAGCTGAGGG AACAGCTTTCTAGTGTGTCGTCCTTCGAGAGGTTTGAGATCTTCCCCAAGGAGAGCTCATGGCCAAACCACAATACAAACGGGGTCACTG CGGCATGTAGCCATGAGGGCAAGTCATCCTTCTATCGAAACCTCCTGTGGCTGACCGAAAAGGAAGGTAGCTACCCCAAACTTAAAAACA GTTATGTGAACAAAAAGGGCAAGGAAGTCTTAGTGCTTTGGGGCATCCACCACCCCCCAAATAGCAAAGAGCAACAGAATTTGTACCAGA ACGAAAATGCTTATGTGTCTGTAGTAACCTCTAACTATAACAGGCGTTTCACTCCCGAGATTGCCGAGCGACCGAAAGTTAGAGACCAGG CCGGACGCATGAATTACTACTGGACCCTGTTGAAACCTGGAGATACGATAATCTTTGAGGCCAACGGAAACCTGATAGCTCCTATGTATG CGTTCGCACTCAGTCGCGGGTTCGGGTCCGGGATAATCACCAGTAACGCGTCCATGCACGAGTGTAATACGAAGTGCCAGACCCCACTA GGAGCGATTAACTCGTCTCTGCCATATCAGAATATCCATCCTGTGACAATCGGAGAGTGTCCTAAATACGTCCGCTCAGCCAAACTGCGA ATGGTGACGGGCCTTCGCAACAATCCCAGTATCCAGTCTCGGGGACTCTTTGGAGCCATCGCCGGTTTTATCGAAGGCGGATGGACTGG GATGATCGATGGCTGGTACGGCTACCACCACCAAAATGAGCAAGGCTCCGGGTACGCTGCCGACCAGAAGTCAACTCAGAATGCAATTA ACGGGATCACCAACAAGGTTAACACCGTAATCGAAAAGATGAACATCCAGTTCACGGCCGTGGGGAAGGAGTTTAATAAACTTGAGAAAC GGATGGAGAACCTCAACAAAAAGGTGGACGACGGCTTTCTGGACATCTGGACCTATAATGCTGAGCTTTTGGTGCTGCTGGAAAACGAG CGCACACTGGATTTCCACGATAGTAATGTGAAGAACCTGTACGAAAAGGTTAAGTCCCAACTCAAAAATAACGCGAAAGAAATCGGCAAT GGGTGCTTTGAATTTTATCACAAGTGTGACAACGAATGCATGGAATCTGTGCGGAATGGGACCTACGATTACCCAAAATATAGCGAAGAA TCCAAGCTGAACAGAGAGAAGGTGGACGGCGTGAAACTCGAAAGTATGGGTATCTACCAAATCCTAGCCATCTACTCTACCGTTGCCAGT AGTCTAGTGTTGCTGGTGTCGTTAGGTGCCATCTCTTTTTGGATGTGTAGCAATGGCTCTCTGCAATGTCGGATCTGTATC
2721	ATGAAGGCCAACCTGCTTGTGCTGCTTTGCGCACTAGCCGCTGCAGACGCCGACACAATTTGTATTGGATACCATGCTAATAACTCGACC GACACAGTTGACACAGTCCTTGAGAAGAACGTGACAGTTACTCACTCCGTGAACCTGCTCGAGGACTCGCATAATGGCAAACTCTGCCGT CTGAAGGGTATCGCCCCCCTCCAACTGGGGAAATGCAATATACCCGCCTGGCTGCTGGGAAATCCAGAGTGCGATCCACTGTTGCCTGT GCGCAGTTGGAGTTATATCGTGGAGACACCCAACTCAGAGAACGGTATCTGCTACCCTGGGGACTTCATAGATTATGAAGAGCTGCGCG AGCAGCTATCTTCCGTTAGCTCTTTCGAACGCTTTGAAATCTTTCCTAAAGAAAGCAGCTGGCCAAATCATAACACAAACGGAGTGACAGC TGCCTGCAGTCACGAGGGTAAAAGTTCTTTTTATAGAAATCTCCTGTGGTTGACGGAGAAAGAGGGCAGCTATCCAAAGCTCAAGAATAG CTACGTCAATAAAAAAGGTAAGGAGGTGCTAGTCTTATGGGGAATCCATCACCCACCCAATTCCAAGGAACAGCAGAACCTCTATCAGAA CGAGAACGCCTATGTGTCCGTAGTGACTAGCAACTATAATCGTAGATTCACACCTGAGATTGCAGAGAGACCCAAGGTTAGAGACCAAGC CGGCCGCATGAATTATTACTGGACACTGCTTAAACCCGGGGATACTATCATCTTCGAAGCGAATGGAAACCTGATCGCTCCAATGTATGC TTTTGCTCTGTCCAGGGGTTTTGGCTCGGGGATTATCACAAGCAACGCATCTATGCACGAGTGTAATACCAAGTGTCAGACCCCCCTTGG CGCTATCAACTCTAGTCTACCGTACCAGAACATCCACCCTGTTACCATTGGTGAATGCCCGAAATACGTGCGGAGTGCCAAGCTGAGAAT GGTGACCGGCCTGAGGAACAATCCTTCCATCCAGTCCCGGGGACTTTTCGGAGCTATCGCTGGCTTCATTGAAGGTGGGTGGACCGGC ATGATCGACGGGTGGTATGGATATCACCATCAGAACGAGCAAGGAAGCGGGTATGCTGCAGACCAGAAGTCCACACAGAACGCAATAAA CGGCATCACTAACAAAGTGAACACCGTGATTGAAAAAATGAATATACAGTTTACCGCCGTAGGGAAGGAATTTAATAAACTGGAAAAGCG CATGGAAAATTTGAACAAAAAAGTCGACGATGGTTTTTTAGACATTTGGACCTATAATGCGGAACTGCTAGTACTGCTCGAGAACGAACGG ACCCTCGACTTTCACGATTCAAACGTCAAAAATCTATACGAGAAGGTGAAGTCGCAGCTCAAGAATAATGCCAAAGAAATCGGCAACGGA TGTTTCGAATTCTACCACAAATGCGACAATGAATGTATGGAATCTGTTAGAAACGGCACGTATGATTACCCCAAGTACAGCGAAGAATCTA AACTCAACCGGGAGAAAGTCGACGGAGTTAAGCTTGAGTCCATGGGAATCTATCAGATCTTGGCGATCTATTCCACCGTGGCAAGTTCCC TCGTATTGCTGGTGTCCCTTGGAGCAATTTCATTCTGGATGTGCAGTAACGGCAGCCTTCAATGCCGCATCTGTATA

2722	ATGAAGGCAAACCTTCTTGTCCTGCTGTGCGCACTCGCCGCAGCCGACGCCGACACAATCTGCATTGGGTATCACGCTAATAATTCAACT GACACTGTGGATACCGTACTGGAAAAGAATGTTACTGTGACTCATAGTGTAAATCTGCTCGAGGACAGTCACAATGGAAAACTTTGCCGG TTAAAGGGCATCGCACCCCTGCAGCTGGGTAAGTGCAACATCGCGGGATGGCTGCTGGGCAACCCCGAGTGTGATCCCCTGCTTCCAG TGCGGTCTTGGTCTTACATTGTCGAAACACCCAACTCTGAAAACGGAATTTGTTACCCTGGCGACTTCATCGACTACGAAGAGCTTAGAG AGCAGCTATCGAGCGTGAGTTCTTTTGAAAGATTTGAGATCTTCCCAAAGGAAAGCTCTTGGCCCAACCATAATACTAATGGCGTCACAG CCGCCTGCTCACATGAAGGGAAGTCTTCTTTCTACAGGAATCTGTTGTGGTTGACGGAGAAGGAGGGAAGTTATCCAAAGCTTAAAAATT CCTACGTGAACAAGAAGGGTAAAGAGGTCCTGGTGCTCTGGGGCATCCACCATCCTCCGAACTCCAAAGAACAGCAGAATCTGTATCAG AACGAAAATGCATATGTCAGTGTGGTGACAAGTAATTATAACCGCCGCTTTACCCCCGAGATCGCCGAACGCCCGAAGGTACGGGACCA AGCGGGTCGGATGAACTACTATTGGACCCTGCTAAAGCCTGGGGACACTATCATCTTCGAGGCCAACGGTAACCTGATCGCTCCCATGT ATGCGTTTGCCCTGAGCAGAGGATTCGGCTCTGGAATCATAACGAGCAACGCCTCAATGCATGAATGCAACACTAAGTGTCAAACTCCCT TGGGCGCAATAAACTCATCCTTGCCATATCAGAACATCCACCCAGTCACCATCGGGGAGTGCCCTAAATATGTGCGTTCAGCCAAGTTAC GCATGGTAACGGGACTCCGGAATAACCCCAGCATTCAATCTCGAGGGCTTTTTGGAGCTATCGCAGGATTCATTGAAGGGGCTGGACT GGGATGATCGATGGATGGTATGGGTACCATCATCAGAACGAGCAGGGCTCCGGCTACGCAGCCGATCAGAAGAGCACTCAGAACGCCA TCAATGGCATCACAAACAAAGTCAACACTGTGATTGAAAAAATGAATATACAGTTTACCGCAGTGGGTAAGGAATTCAACAAGCTGGAGAA ACGGATGGAAAATCTAAACAAGAAAGTGGATGATGGCTTTCTCGACATCTGGACCTACAACGCCGAACTGCTGGTCCTGCTGGAAAATGA GCGCACACTGGATTTCCACGACTCGAACGTCAAGAATCTATACGAGAAAGTGAAATCCCAGCTCAAGAATAATGCCAAGGAGATCGGCAA CGGATGCTTCGAATTTTACCATAAATGCGACAACGAGTGTATGGAGTCCGTCCGAAACGGGACATACGACTATCCCAAGTATTCAGAAGA GTCAAAGCTTAACCGCCAAAAGGTGGATGGCGTTAAGCTGGAATCTATGGGTATTTACCAGATACTGGCCATTTATAGCACTGTGGCTAG CTCATTAGTGCTGCTGGTGTCGCTGGGAGCGATTTCCTTTTGGATGTGTAGCAACGGGAGTTTACAGTGCCGGATTTGCATC
2723	ATGAAGGCTAACCTCCTGGTGCTGCTGTGTGCCCTGGCTGCAGCGGACGCCGACACGATCTGTATCGGATACCATGCTAATAATTCTAC CGACACTGTAGACACCGTGCTGGAAAAGAATGTTACAGTGACCCATAGTGTTAATCTCCTCGAAGATTCACACAACGGCAAGCTCTGCCG TCTCAAAGGAATAGCACCTCTTCAGCTGGGCAAGTGTAACATCGCCGGGTGGCTTCTGGGCAACCCCGAATGTGATCCCCTTCTGCCAG TCCGCTCATGGTCCTATATTGTGGAAACCCCTAATTCCGAGAACGGAATTTGTTATCCCGGAGACTTTATTGATTATGAGGAACTTAGAGA ACAACTGTCTTCGGTCTCGTCCTTTGAGAGGTTCGAGATTTTCCCTAAGGAGAGCTCCTGGCCCAATCACAACACTAACGGTGTGACCGC CGCCTGCTCGCATGAGGGGAAGTCCTCTTTCTACCGCAACCTCCTGTGGCTGACAGAGAAGGAGGGGTCCTATCCCAAACTAAAAAACT CATACGTGAACAAAAAGGGAAAAGAGGTCTTGGTTCTCTGGGGCATCCATCACCCACCAAACAGTAAGGAGCAGCAAAATCTGTACCAGA ATGAAAACGCATATGTTAGCGTAGTGACCTCTAACTATAACAGGCGGTTTACTCCAGAGATTGCCGAGCGGCCCAAAGTGCGAGACCAG GCAGGAAGGATGAACTATTATTGGACTTTGTTAAAACCCGGAGACACAATTATCTTCGAGGCAAACGGGAACTTGATCGCACCCATGTAC GCCTTCGCACTGAGCAGGGGTTTCGGCTCCGGCATCATCACTTCTAATGCCTCTATGCACGAGTGTAATACGAAGTGTCAGACCCCATTG GGAGCCATCAACTCCAGTCTGCCCTATCAGAACATCCATCCGGTGACTATAGGCGAATGCCCCAAGTACGTCAGGTCCGCTAAGCTCCG CATGGTCACCGGACTTAGGAACAACCCGAGTATTCAGAGCCGAGGCCTGTTCGGAGCCATCGCAGGGTTCATAGAGGGAGGGTGGACC GGGATGATAGATGGATGGTATGGCTACCATCATCAAAACGAACAGGGCAGCGGATATGCAGCTGATCAGAAGTCCACCCAGAACGCCAT AAACGGAATCACCAACAAAGTGAACACCGTCATAGAAAAAATGAACATACAGTTCACCGCCGTGGGCAAAGAATTCAATAAACTTGAAAAA CGGATGGAAAATTTGAATAAGAAGGTGGATGACGGCTTCCTGGATATCTGGACTTACAATGCGGAGCTGCTCGTGCTTCTGGAGAACGA GAGAACGCTCGACTTCCACGATTCTAACGTTAAGAACCTGTATGAGAAAGTTAAAAGTCAGCTCAAAAATAACGCGAAGGAAATCGGGAA CGGCTGCTTTGAATTTTACCATAAATGCGATAATGAGTGCATGGAATCCGTGCGAAACGGCACGTACGACTATCCAAAGTATTCAGAGGA ATCAAAATTGAACAGAGAAAAGGTCGATGGTGTGAAACTGGAGAGTATGGGGATATATCAAATCCTTGCGATCTACTCTACTGTGGCTTCT TCCCTGGTCCTGCTGGTGTCCTTGGGGGCAATTTCATTTTGGATGTGTTCTAATGGGAGTTTACAGTGCAGGATCTGCATA
2724	ATGAAAGCCAATCTTCTGGTTCTGCTCTGCGCTCTAGCTGCTGCCGATGCTGACACTATTTGTATAGGCTACCACGCAAATAATTCGACG GATACTGTCGACACGGTCCTGGAAAAAAATGTCACAGTGACTCATTCAGTTAATCTCTTGGAGGATTCTCACAATGGAAAGTTGTGTAGAC TGAACGGCATTGCTCCATTGCAGCTTGGGAAGTGTAATATCGCGGGCTGGTTACTGGGCAACCCTGAATGCGACCCCCTCTTGCCTGTC CGGTCCTGGTCTTATATCGTGGAGACACCAAATTCTGAGAATGGAATCTGCTATCCGGGTGACTTTATCGATTACGAGGAGCTTCGGGAG CAGCTGTCTTCAGTGTCCAGTTTTGAAAGATTTGAGATATTTCCTAAGGAGAGCTCCTGGCCAAATCATAATACCAATGGCGTGACAGCC GCATGTTCACACGAAGGAAAAAGCTCCTTCTATCGGAACCTGCTCTGGCTTACAGAAAAGGAAGGCTCTTACCCCAAACTCAAGAACAGT TACGTGAATAAGAAAGGTAAAGAGGTTCTAGTACTCTGGGGCATCCACCACCCGCCAAACAGCAAGGAGCAACAGAACCTCTATCAGAA CGAGAATGCCTACGTGTCCGTGGTCACATCAAACTATAACCGCAGGTTTACACCAGAAATTGCAGAGCGGCCTAAAGTCCGTGATCAGG CAGGCAGGATGAACTATTACTGGACTTTGCTGAAGCCGGGAGACACCATTATCTTCGAGGCGAATGGGAACCTCATTGCCCCAATGTAC GCCTTCGCGCTCTCTCGCGGCTTTGGCTCAGGAATTATCACGTCAAATGCTTCTATGCACGAGTGTAATACAAAATGTCAGACCCCTCTG GGCGCCATAAACTCTTCCCTGCCTTATCAAAATATTCACCCAGTGACTATTGGGGAGTGCCCCAAATACGTTAGAAGCGCTAAACTGCGT ATGGTAACGGGTTTGCGGAACAACCCAAGTATCCAGAGCCGCGGCCTGTTTGGCGCCATTGCTGGTTTTATTGAGGGGGGTTGGACAGG AATGATCGATGGATGGTACGGCTACCACCATCAGAATGAACAGGGGAGTGGGTACGCCGCCGACCAAAAGTCCACACAGAATGCGATCA ACGGGATCACTAACAAGGTGAACACCGTGATTGAGAAGATGAACATACAGTTTACCGCAGTCGGAAAGGAATTCAACAAACTTGAAAAAC GCATGGAGAATCTGAATAAAAAAGTGGACGACGGATTTTTAGATATATGGACCTATAATGCCGAATTGCTGGTGCTCTTAGAAAACGAACG AACGCTGGACTTCCACGATTCTAACGTTAAAAACCTTTATGAGAAGGTTAAGAGCCAGTTGAAGAACAACGCGAAGGAAATCGGAAACGG ATGCTTCGAATTTTACCACAAGTGTGACAACGAGTGTATGGAATCTGTTAGAAATGGCACCTACGATTATCCAAAGTACTCAGAAGAAAGC AAACTTAACAGGGAAAAGGTGGATGGTGTGAAACTGGAGTCCATGGGTATCTACCAGATACTTGCAATCTACTCTACGGTTGCTTCAAGC TTGGTGCTGCTTGTGAGCTTGGGAGCCATCTCATTTTGGATGTGTTCAAATGGCAGCTTACAGTGTAGGATCTGCATA

Table 17: Codon Optimized Sequences Encoding H7 Hemagglutinin

SEQ ID NO.	Nucleic Acid Sequence
2725	ATGAATACACAGATCCTTGTATTTGCCCTGATCGCAATCATTCCTACTAATGCTGACAAGATTTGCCTCGGCCACCATGCGGTTAGTAATG GGACCAAAGTGAACACACTGACGGAGAGAGGAGTAGAGGTAGTTAACGCCACCGAGACTGTGGAGAGAACTAACATCCCACGGATTTGT TCCAAGGGTAAACGTACTGTGGATCTGGGGCAGTGCGGCTTGTTGGGTACCATCACTGGACCGCCACAGTGCGACCAGTTTCTGGAGTT TTCCGCCGACCTGATTATAGAACGCCGAGAAGGCAGCGATGTCTGTTACCCCGGTAAATTCGTGAATGAAGAAGCCCTCCGCCAAATCCT GCGTGAGAGTGGGGGCATTGACAAAGAGGCGATGGGCTTTACATACAGCGGGATACGCACCAACGGCGCGACGTCAGCCTGCCGCAG GAGTGGAAGTTCTTTCTATGCCGAGATGAAGTGGCTGTTGTCTAATACTGATAATGCAGCCTTCCCCCAGATGACAAAGTCGTACAAGAA CACCAGAAAGTCACCCGCACTCATTGTGTGGGGTATTCACCATTCTGTCTCTACTGCTGAGCAGACGAAACTGTATGGTTCTGGGAACAA GCTCGTGACCGTGGGGTCCAGCAACTATCAGCAGAGCTTTGTGCCCTCCCCAGGAGAGAGGCCACAGGTGAATGGGTTGTCAGGCCGT ATCGATTTCCACTGGCTTATGTTGAATCCAAACGACACAGTGACATTTTCTTTCAACGGCGCCTTCATTGCACCAGACAGAGCTAGCTTTC TGCGAGGGAAGAGTATGGGTATCCAGTCTGGTGTGCAGGTCGACGCTAACTGTGAGGGGGATTGCTACCATAGCGGGGGAACAATTATT TCAAACCTGCCCTTCCAGAATATTGACAGCAGGGCAGTGGGAAAGTGCCCCCGCTATGTGAAACAGAGATCTTTGTTGCTGGCCACCGG CATGAAGAACGTTCCCGAAATTCCTAAGGGCCGGGGGCTTTTTGGGGCGATAGCCGGGTTCATTGAAAACGGGTGGGAGGGGCTAATC GATGGTTGGTACGGTTTTAGGCATCAAAACGCCCAGGGCGAAGGCACTGCGGCCGACTACAAATCCACACAGAGCGCCATCGACCAAAT CACTGGGAAACTTAACCGACTTATTGAGAAAACAAATCAACAGTTTGAGCTGATTGATAACGAATTTAATGAGGTAGAGAAGCAGATCGGC AATGTAATAAACTGGACAAGGGACAGCATTACTGAAGTCTGGAGCTACAACGCCGAGTTGCTCGTCGCGATGGAGAATCAGCATACCATC GATTTAGCCGACTCAGAGATGGATAAGCTGTATGAACGCGTAAAGAGGCAGCTGCGCGAGAACGCTGAAGAGGACGGGACTGGCTGCT TCGAAATTTTCCACAAATGCGATGATGATTGTATGGCTTCCATCCGGAACAATACCTACGACCATTCTAAGTACCGTGAAGAGGCCATGCA GAATAGGATCCAGATTGACCCAGTGAAGCTGAGCTCGGGTTATAAAGACGTGATTTTGTGGTTCAGTTTTGGCGCCTCTTGTTTTATCCTT CTGGCAATTGTTATGGGGTTGGTTTTCATCTGCGTCAAGAACGGCAACATGCGCTGTACTATTTGTATC

2726	ATGAATACCCAGATTCTCGTCTTCGCACTGATAGCTATAATTCCAACCAATGCCGACAAGATATGCTTGGGACACCATGCGGTGTCTAAC GGCACAAAGGTGAATACTCTGACAGAGAGAGGAGTGGAGGTGGTTAACGCTACAGAGACTGTGGAAAGGACGAATATACCAAGGATATG CTCCAAGGGCAAGAAGACCGTCGACCTCGGCCAGTGTGGCCTGCTTGGCACTATCACAGGTCCCCCTCAATGCGATCAATTTCTGGAAT TCTCTGCTGATCTGATAATCGAGCGCCGCGAGGGGTCTGACGTATGCTATCCCGGGAAATTCGTGAATGAGGAGGCTCTGAGACAGATA TTGCCGGCAGTCGGGGGGCATCGATAAAGAAGCTATGGGATTCACTTACTCGGGGATCAGAACAAACGGAGCTACATCTGCGTCTGGC GGTCTGGCAGCTCGTTTTATGCCGAGATGAAGTGGCTGCTCTCCAATACAGACAATGCCGCATTCCCACAAATGACGAAAAGCTATAAAA ATACTCGCAAGTCACCCGCACTAATCGTGTGGGGCATCCACCACAGCGTGTCCACCGCAGAGCAGACCAAATTGTATGGTAGCGGCAAT AAGCTGGTTACAGTGGGATCCTCCAACTACCAACAGTCCTTCGTCCCAAGCCCAGGGGCCCGACCGCAAGTTAACGGTCAGTCCGGACG CATTGATTTTCACTGGCTGATGCTTAACCCAAACGACACAGTTACCTTTTCGTTTAACGGCGCTTTCATTGCCCCAGATCGGGCCTCTTTT CTGCGGGGTAAGAGCATGGGAATCCAGTCGGGGGTACAGGTCGACGCGAATTGTGAAGGCGATTGTTACCATTCTGGGGGCACTATTAT ATCAAATCTTCCATTCCAGAATATTGACTCGCGCGCTGTGGGCAAATGCCCCAGGTACGTTAAGCAGCGATCACTGCTCCTCGCCACAGG TATGAAAAATGTTCCAGAGATTCCCAAGGGCAGGGGGCTGTTCGGTGCAATCGCCGGCTTTATAGAAAATGGGTGGGAGGGACTCATCG ATGGTTGGTACGGTTTTAGGCATCAAAACGCCCAGGGGGAGGGGACGGCTGCCGATTACAAGTCTACCCAGTCAGCTATAGACCAGATC ACTGGGAAACTAAACAGGCTGATCGAGAAAACCAACCAGCAGTTCGAACTGATTGACAACGAGTTCAACGAGGTTGAAAAGCAGATTGGT AACGTGATTAACTGGACCAGAGACTCTATCACCGAGGTCTGGAGCTACAATGCAGAACTACTTGTGGCTATGGAAAACCAGCATACAATT GATCTTGCTGACTCTGAGATGGACAAGCTTTATGAGAGGGTGAAGAGACAGCTTCGGGAGAACGCTGAGGAGGATGGAACCGGTTGTTT CGAGATCTTTCACAAATGCGATGACGATTGCATGGCTAGCATTAGGAATAACACCTATGACCACTCTAAATACAGAGAGGAGGCTATGCA GAATCGCATCCAGATCCACCCAGTGAAGCTGTCCTCTGGATACAAGGACGTAATCCTGTGGTTCTCATTTGGGGCTTCGTGCTTCATCTT GCTGGCGATTGTTATGGGCTTAGTGTTTATCTGCGTTAAAAACGGGAACATGCGGTGTACAATCTGTATC
2727	ATGAACACACAGATTCTGGTCTTTGCCTTGATCGCAATAATTCCAACTAATGCCGACAAAATTTGTTTAGGTCATCACGCCGTTTCAAATG GAACTAAGGTCAACACCCTGACCGAGAGGGGCGTCGAGGTCGTGAATGCCACAGAAACTGTGGAGAGAACCAACATCCCTCGCATCTGT AGCAAGGGCAAGAAGACTGTCGATCTTGGACAGTGCGGTCTCCTGGGAACGATTACAGGCCCCCCACAGTGCGACCAATTCCTAGAATT CTCTGCCGACTTAATCATCGAAAGGCGTGAGGGATCTGACGTGTGCTATCCGGGAAAGTTCGTGAATGAGGAGGCCCTACGGCAGATTC TGCGGAAATCCGGAGGTATAGACAAAGAGGCAATGGGCTTCACCTACAGTGGGATACGCACAAATGGGGCCACATCCACTTGTCGGAGA AGCGGCTCTTCCTTTTATGCCGAGATGAAGTGGCTCTTATCCAACACAGACAATGCCGCCTTCCCTCAGATGACGAAGAGCTACAAAAAC ACCAGGAAGTCTCCAGCAATCATTGTTTGGGGCATCCACCATAGCGTCTCCACCGCTGAGCAAACCAAATTATATGGGTCCGGGAATAAA TTGGTCACAGTCGGTAGCTCCAACTACCAGCAGTCATTTGTCCCCTCACCGGGGGCTCGCCCACAAGTAAATGGTTTGTCTGGCAGAATC GACTTTCATTGGCTCATGCTCAATCCAAACGACACGGTGACTTTCAGCTTTAACGGGGCATTTATTGCCCCCGACCGAGCATGTTTTCTTA GAGGCAAAAGCATGGGGATTCAGAGCGGAGTCCAGGTCGACGCGGATTGCGAGGGTGACTGCTACCACTCAGGGGGTACCATTATCTC CAACCTTCCTTTTCAGAATATCGATTCGAGAGCTGTGGGCAAGTGCCCTCGGTATGTTAAACAGCGATCCCTCCTTCTGGCCACTGGTAT GAAGAACGTGCCTGAGATACCCAAAGGGAGGGGCCTGTTTGGGGCTATCGCTGGATTTATTGAGAATGGTTGGGAGGGCCTTATCGACG GCTGGTACGGCTTTAGGCACCAGAACGCGCAGGGCGAAGGCACAGCTGCCGACTACAAAAGCACCCAGAGCGCGATTGACCAGATCAC CGGCAAATTGAACAGGCTTATCGAAAAAACGAACCAACAGTTTGAGCTAATTGACAATGAGTTTAACGAGGTCGAGAGGCAGATTGGGAA CGTCATCAACTGGACAAGAGACAGCATTACCGAGGTGTGGTCTTACAATGCAGAGTTGCTAGTGGCCATGGAAAACCAACATACGATCGA CCTCGCGGATTCTGAGATGGATAAGCTCTATGAAAGGGTGAAAAGGCAGCTGCGGGAAAATGCTGAGGAAGACGGAACAGGATGTTTTG AAATCTTCCACAAATGTGATGATGACTGTATGCCCTCTATAAGGAACAACACCTATGATCATTCCAAGTATCGAGAAGAGGCCATGCAGAA CCGTATCCAGATTGACCCTGTAAAACTCAGTTCAGGGTACAAGGATGTGATCCTGTGGTTCAGCTTTGGCGCAAGCTGTTTCATTCTTCTC GCGATTGTGATGGGGCTAGTGTTCATTTGTGTGAAAAATGGAAATATGAGATGTACCATTTGTATC
2728	ATGAACACCCAGATTCTGGTCTTCGCTCTGATTGCTATTATCCCAACCAATGCAGACAAGATCTGTCTGGGGCACCATGCTGTGTCGAAC GGCACGAAAGTGAACACACTGACCGAAAGAGGAGTCGAGGTGGTGAACGCGACCGAGACAGTGGAGCGGACAAACATCCCACGTATCT GCTCCAAGGGCAAGCGTACAGTGGACTTGGGGCAATGTGGCCTCCTTGGAACGATCACAGGACCACCGCAGTGCGACCAGTTTCTGGA GTTTTCTGCCGATTTAATCATCGAAAGGCGGGAGGGTTCCGACGTTTGTTACCCGGGGAAGTTCGTTAACGAGGAAGCCCTGAGACAGA TCCTACGTGAGAGCGGCGGTATCGACAAGGAGGCTATGGGCTTTACATATAGCGGAATTAGAACAAACGGCGCTACCAGCGCCTGCCG CCGCAGCGGTTCATCTTTTTATGCCGAAATGAAGTGGCTGCTCTCCAACACCGACAACGCCGCTTTTCCACAGATGACGAAATCCTATAA GAATACCAGGAAAAGTCCAGCTCTGATTGTGTGGGGGATACACCACTCTGTGTCAACAGCGGAACAGACAAAGCTCTATGGAAGCGGAA ACAAACTCGTAACGGTCGGCAGTTCCAACTATCAGCAATCCTTTGTGCCAAGTCCAGGCGCGCGTCCTCAGGTGAACGGCCTAAGTGGG AGAATCGATTTCCATTGGTTAATGCTTAATCCAAATGACACCGTGACCTTTAGCTTTAACGGCGCCTTTATTGCTCCCGACCGAGCTTCTTT CCTTAGAGGTAAGTCAATGGGTATCCAGAGCGGCGTGCAGGTTGACGCAAACTGTGAAGGCGATTGTTACCACTCCGGAGGCACTATCA TTAGCAACTTGCCATTTCAGAACATAGATTCACGCGCTGTGGGAAAGTGCCCTAGGTACGTGAAGCAGAGGTCCCTGTTGCTCGCTACG GGCATGAAGAACGTGCCAGAGATTCCCAAAGGCAGAGGGTTGTTCGGTGCCATTGCTGGTTTCATTGAGAATGGTTGGGAGGGCCTGAT CGACGGGTGGTACGGTTTCCGTCACCAGAATGCTCAAGGGGAGGGTACAGCAGCCGACTATAAATCCACCCAGAGCGCAATCGATCAAA TTACCGGCAAGCTAAATAGGCTGATTGAAAAAACTAATCAGCAGTTCGAGCTCATAGACAATGAGTTCAACGAAGTGGAAAAGCAGATTG GCAACGTCATCAATTGGACCAGAGATTCCATCACCGAGGTGTGGTCTTATAATGCCGAGCTGTTGGTTGCCATGGAAAATCAGCATACAA TAGACCTGGCCGACTCCGAAATGGATAAACTGTACGAGAGGGTTAAACGCCAACTCCGTGAGAATGCAGAGGAAGACGGAACAGGTTCT TTCGAGATCTTTCACAAATGCGACGACGACTGCATGGCCTCCATAAGGAATAATACATATGATCATAGCAAATACAGGGAGGAGGCAATG CAAAACCGGATTCAGATTGATCCTGTGAAGCTGAGCAGCGGCTACAAGGACGTTATTCTTTGGTTCAGTTTCGGTGCATCATGCTTCATA CTCCTGGCAATAGTGATGGGTCTGGTGTTCATCTGCGTGAAAAACGGAAATATGAGGTGCACCATATGCATC

2729	ATGAATACCCAAATACTCGTCTTTGCTCTCATCGCTATCATTCCTACGAATGCCGACAAAATATGCCTAGGACATCACGCCGTGAGCAATG GGACAAAGGTGAATACCTTAACCGAAAGAGGCGTGGAGGTCGTGAACGCCACAGAGACTGTGGAGCGCACCAATATACCTAGAATCTGC TCAAAGGGCAAGAAAACAGTGGACCTAGGACAGTGTGGGCTTCTGGGGACGATCACTGGGCCACCACAGTGCGACCAGTTCCTGGAGT TCAGCGCGGACCTGATCATCGAACGCCGAGAGGGCTCTGATGTGTGCTACCCTGGCAAGTTTGTGAACGAGGAGGCTCTTAGGCAGATT CTCAGGCAATCCGGCGGAATTGACAAGGAAGCTATGGGCTTCACTTATAGTGGAATCCGCACCAATGGGGCAACGTCCGCTTGTAGGAG ATCCGGGAGCAGTTTTTATGCAGAAATGAAATGGCTGCTTAGTAACACGGACAACGCCGCTTTTCCGCAGATGACTAAGTCCTACAAGAA CACCCGCAAATCACCAGCTCTGATTGTGTGGGGGATTCACCATTCCGTGAGCACAGCCGAGCAGACAAAGTTGTATGGCTCTGGAAATA AACTCGTGACTGTGGGCAGTTCGAACTACCAGCAGTCTTTTGTGCCTTCTCCCGGAGCCCGTCCCCAGGTCAACGGGCAGTCCGGAAGA ATTGATTTCCACTGGTTGATGCTCAACCCGAATGATACAGTGACTTTTAGCTTCAACGGCGCTTTTATTGCCCCTGATCGTGCCAGCTTCC TTAGGGGCAAATCCATGGGGATTCAGTCAGGAGTTCAGGTAGATGCCAATTGTGAGGGAGATTGTTACCACTCAGGGGGTACGATCATC AGCAACTTGCCATTTCAGAACATCGACAGTCGGGCAGTGGGCAAATGCCCACGTTACGTGAAGCAACGGAGTCTCCTGTTAGCCACAGG GATGAAAAATGTTCCAGAGATCCCTAAGGGCCGGGGACTGTTCGGGGCCATTGCCGGATTTATCGAGAATGGATGGGAGGGACTTATTG ATGGTTGGTACGGCTTTCGGCATCAGAATGCCCAAGGGGAGGGCACAGCTGCCGACTATAAATCAACCCAGTCTGCAATAGATCAAATC ACCGGTAAGCTGAATCGCCTTATCGAAAAAACGAACCAACAATTCGAGCTGATCGATAACGAATTCAATGAAGTGGAGAAACAGATTGGG AATGTTATTAACTGGACAAGAGACAGCATAACCGAAGTCTGGTCCTATAATGCCGAGTTACTTGTGGCCATGGAAAACCAGCATACGATT GACTTAGCCGACTCCGAGATGGACAAGTTGTATGAGCGGGTGAAGAGACAGTTACGGGAGAACGCCGAAGAGGACGGGACTGGTTGTT TTGAAATCTTCCATAAGTGTGACGACGACTGCATGGCCAGTATTCGCAATAACACCTACGACCATAGCAAGTACAGGGAAGAAGCCATGC AGAATCGTATTCAGATCGATCCTGTCAAACTTAGTAGTGGGTACAAGGACGTTATCCTGTGGTTTTCTTTTGGGGCCTCCTGTTTCATTCT TCTGGCTATTGTAATGGGACTGGTTTTCATATGTGTGAAAAATGGTAACATGCGGTGCACTATATGTATC
2730	ATGAATACTCAGATTCTAGTGTTCGCTTTGATAGCTATAATCCCAACCAACGCCGACAAAATATGCCTGGGGCACCACGCAGTATCCAAC GGGACCAAGGTCAATACCCTGACCGAACGGGGTGTGGAAGTCGTGAACGCCACAGAGACCGTTGAGAGAACAAACATCCCTAGAATTTG TAGCAAAGGGAAGAGGACTGTCGACTTGGGGCAGTGCGGGTTACTGGGAACAATCACTGGCCCACCCCAGTGTGACCAGTTCCTCGAG TTCTCTGCTGATCTTATCATTGAGAGAAGAGAGGGCAGTGACGTGTGTTATCCTGGAAAATTCGTAAACGAGGAGGCGCTTAGGCAGATT CTGCGCGAATCAGGAGGTATCGATAAAGAGGCAATGGGGTTTACTTACAGCGGTATCCGAACCAACGGCGCTACATCTGCATGCCGCAG GTCTGGCTCATCTTTCTATGCAGAGATGAAGTGGTTGCTCTCTAACACCGATAATGCGGCATTTCCTCAGATGACCAAGAGCTACAAGAA CACCAGAAAATCCCCCGCGTTGATCGTGTGGGGCATCCACCATTCAGTGTCCACCGCTGAGCAGACAAAACTCTACGGATCTGGGAACA AGTTGGTGACTGTGGGGTCCTCTAACTACCAGCAATCGTTTGTGCCATCCCCAGGGGCTAGACCCCAGGTCAACGGTCTCTCTGGCCGG ATCGACTTCCACTGGCTTATGCTTAACCCAAACGATACCGTCACCTTTAGCTTTAATGGTGCTTTCATTGCCCCCGACCGGGCAAGCTTCT TACGAGGCAAGAGTATGGGAATACAGAGTGGGGTTCAAGTGGATGCCAACTGCGAGGGCGACTGTTACCACAGCGGCGGCACCATAAT TTCAAATCTGCCGTTCCAGAATATAGACTCTCGAGCTGTTGGGAAGTGCCCCCGATACGTAAAGCAGCGCTCTCTGTTACTGGCCACCGG CATGAAAAATGTTCCTGAGATACCGAAGGGACGTGGCCTCTTCGGGGCCATCGCCGGGTTCATCGAGAACGGCTGGGAAGGACTGATC GATGGATGGTACGGATTCAGGCATCAGAATGCTCAGGGAGAAGGGACGGCTGCAGATTATAAGTCTACACAGTCCGCCATCGATCAGAT CACCGGTAAGCTTAATAGAATCATAGAGAAGACGAACCAGCAGTTCGAGCTGATCGATAACGAGTTTAATGAGGTCGAAAAGCAGATTGG CAACGTGATAAATTGGACAAGAGACAGTATTACTGAAGTGTGGTCCTACAACGCCGAGCTTTTGGTGGCCATGGAAAACCAACACACCAT CGACTTAGCAGATTCCGAGATGGATAAACTTTACGAGCGAGTAAAAAGACAGCTCAGGGAGAATGCTGAAGAAGACGGCACAGGATGTT TCGACATTTTCCACAAGTGTGATGACCGATTGCATGGCATCAATTCGCAACAACACTTACGATCATAGCAAGTATCGCGAGGACCCCATGC AGAACCGCATCCAAATTGATCCTGTCAAGCTAAGTTCCGGGTATAAAGACGTGATTTTGTGGTTTTCTTTTGGCGCTAGCTGCTTTATTCT ACTGGCTATTGTCATGGGCTTGGTTTTCATCTGCGTGAAAAATGGCAATATGCGCTGCACTATATGTATC
2731	ATGAATACACAAATTCTGGTCTTCGCCTTGATTGCCATTATACCAACTAATGCCGATAAGATTTGCCTCGGGCACCATGCCGTCTCCAATG GCACAAAAGTCAACACACTGACTGAACGAGGGGTCGAGGTGGTCAACGCCACTGAAACCGTGGAAAGGACCAACATCCCCCGCATTTGC AGCAAGCGGTAAAAAAACGGTAGACCTGGGACAGTGTGGGCTGCTAGGTACTATTACCGGCCCCCCACAATGTCACCAATTCTTAGAGTT CTCCGCTGACTTGATAATTGAACGGAGAGAGGGTTCTGACGTGTGCTACCCTGGGAAGTTTGTAAATGAAGAAGCCCTTCGACAGATTCT CAGGGAGTCCGGCGGTATTGATAAGGAGGCCATGGGATTTACGTACAGCGGAATTCGGACTAATGGGGCCACTTCGGCCTGCCGGCGC TCTGGTTCCTCCTTTTATGCAGAGATGAAATGGCTCCTATCCAACACGGATAACGCCGCCTTCCCACAGATGACAAAATCCTACAAGAACA CTCGGAAGAGCCCAGCGCTGATTGTGTGGGGTATTCACCATAGCGTGTCCACTGCCGAGCAGACGAAGCTGTACGGAAGTGGCAACAA ATTGGTCACTGTAGGCTCATCTAATTACCAGCAGAGCTTTGTTCCTTCCCCTGGGGCAAGGCCACAAGTGAATGGCCAAAGTGGTAGAAT CGATTTCCACTGGCTCATGTTGAATCCTAATGACACGGTAACTTTCAGTTTCAACGGCGCTTTCATTGCACCCGACCGGGCTTCATTTCTC CGGGGGAAGTCCATGGGGATCCAGTCAGGAGTCCAGGTCGATGCTAATTGTGAAGGCGATTGTTACCACTCCGGTGGGACCATCATTAG CAATCTGCCATTTCAGAACATCGATTCCAGAGCAGTCGGCAAGTGCCCTAGATATGTGAAGCAGAGGAGCCTGCTGCTAGCTACGGGTA TGAAAAATGTGCCAGAGATCCCAAAGGGGCGGGGTCTTTTCGGTGCCATAGCTGGATTTATCGAGAATGGCTGGGAGGGTCTGATCGAC GGATGGTACGGGTTCAGGCATCAGAATGCCCAGGGGGAGGGCACTGCCGCCGACTATAAGTCTACACAAAGCGCTATTGACCAAATTAC CGGTAAGCTGAACAGACTGATCGAGAAGACAAACCAGCAGTTCGAGCTCATCGATAATGAATTCAACGAGGTAGAAAAGCAAATCGGCAA CGTGATTAACTGGACCAGGGATTCTATCACTGAGGTGTGGTCATACAACGCGGAGCTGCTCGTTGCCATGGAAAATCAGCACACTATTGA CCTGGCCGACAGCGAAATGGACAAGCTGTACGAAAGAGTGAAACGACACGCTGCGGGAGAACGCCGAAGAGGACGGGACTGGTTGTTTC GAAATCTTCCATAAGTGCGATGACGACTGTATGGCTTCAATCAGGAACAACACATACGACCACTCTAAGTACAGAGAGGAGGCTATGCAG AATCGTATCCAGATTGACCCTGTAAAACTCTCCTCTGGGTACAAAGACGTGATCCTGTGGTTTTCGTTCGGCGCATCCTGCTTCATCCTGC TCGCGATCGTCATGGGATTAGTGTTCATTTGCGTCAAAAATGGCAATATGAGGTGTACCATTTGCATC

2732	ATGAATACTCAGATCCTAGTGTTTGCTCTCATTGCTATAATCCCCACAAATGCTGACAAGATCTGCTTGGGCCATCACGCCGTGAGTAACG GCACTAAGGTGAATACGCTGACGGAACGGGGTGTAGAAGTTGTAAACGCCACCGAAACAGTCGAAAGGACAAACATCCCGCGCATCTGC AGTAAAGGTAAAAAGACCGTTGACCTCGGCCAATGCGGACTGCTGGGGACTATCACCGGACCCCCCCAATGTGATCAATTTCTGGAGTT CAGTGCGGACTTGATAATCGAAAGACGGGAAGGTTCCGATGTGTGCTATCCAGGCAAGTTCGTTAATGAGGAAGCACTCCGGCAAATTC TTCGGCGAAAGCGGGGGCATAGACAAGGAAGCCATGGGCTTTACTTACAGCGGAATCCGCACAAACGGAGCAACATCCGCTTGCAGGAG AAGTGGTAGTTCTTTTTATGCCGAGATGAAATGGTTGCTGAGCAATACCGACAACGCAGCCTTTCCACAGATGACCAAGAGCTACAAAAA TACAAGAAAGTCCCCAGCTCTGATTGTGTGGGGAATACACCATAGCGTTAGCACAGCCGAACAAACAAAGCTGTACGGTTCAGGAAATAA ACTTGTGACGGTAGGATCCAGCAATTATCAGCAATCTTTCGTGCCATCACCAGGCGCTCGGCCTCAGGTCAACGGGCAGTCGGGCCGTA TAGATTTTCATTGGCTGATGCTGAATCCTAATGATACTGTGACCTTTAGCTTTAACGGGGCGTTCATAGCCCCGATCGGGCATCTTTTCT CAGGGGTAAAAGCATGGGTATTCAGAGTGGAGTGCAGGTTGACGCCAATTGCGAGGGAGATTGCTATCATTCCGGCGGGACCATTATTT CCAACCTCCCCTTCCAGAACATCGACAGCCGAGCTGTCGGCAAATGCCCCAGGTATGTGAAACAACGCAGCCTGTTGTTAGCAACAGGG ATGAAGAATGTTCCTGAAATTCCAAAAGGGCGAGGTCTGTTCGGCGCTATCGCAGGGTTCATTGAGAACGGTTGGGAGGGCCTGATCGA TGGCTGGTATGGCTTTAGGCACCAAAACGCACAGGGCGAAGGGACTGCCGCAGACTACAAATCTACACAGTCAGCGATTGATCAAATCA CTGGCAAACTGAACCGCCTCATTGAGAAGACCAACCAGCAGTTCGAACTGATAGACAATGAGTTCAATGAGGTTGAAAAGCAGATAGGTA ATGTGATAAATTGGACCAGAGATAGTATAACTGAAGTCTGGTCCTATAACGCCGAGTTACTTGTCGCTATGGAGAACCAGCACACAATTGA CCTGGCGGACAGCGAGATGGATAAGCTGTACGAACGCGTAAAAAGACAACTGCGCGAGAATGCGGAGGAGGACGGCACCGGGTGTTTC GAGATCTTTCACAAGTGTGATGATGATTGTATGGCCTCCATTAGAAACAACACCTATGATCACTCCAAATACCGGGAGGAAGCAATGCAG AACCGCATACAGATTGACCCCGTCAAGTTGAGTTCCGGGTATAAAGACGTGATACTCTGGTTCAGTTTTGGTGCCTCTTGCTTTATCCTGC TTGCCATCGTGATGGGACTGGTGTTTATTTGCGTTAAAAACGGGAATATGCGTTGTACGATTTGTATA
2733	ATGAATACCCAGATTCTTGTGTTCGCCCTCATTGCCATCATCCCCACCAATGCCGACAAGATTTGCTTGGGACATCACGCTGTGAGTAATG GCACAAAAGTAAATACGTTAACCGAACGCGGGGTTGAAGTGGTGAATGCTACAGAGACGGTAGAGCGCACAAACATCCCGAGAATTTGC TCCAAAGGCAAAAAGACTGTTGACCTTGGCCAGTGCGGGCTGCTGGGTACGATTACTGGGCCACCTCAGTGCGATCAGTTCCTAGAATT TTCAGCAGATTTAATCATCGAAAGACGTGAAGGATCAGATGTGTGTTATCCCGGAAAGTTCGTGAATGAGGACGCTTTGAGACAGATCTT GCGTGAATCTGGGGGCATAGATAAGGAAGCCATGGGGTTTACTTATTCTGGAATCCGCACTAATGGGGCCACCAGCGCATGTAGGCGCT CCGGATCCTCATTCTATGCCGAGATGAAGTGGCTTCTCAGTAACACAGATAATGCCGCCTTCCCCCAAATGACAAAGAGCTACAAGAATA CACGGAAAAGCCCCGCCCTGATCGTATGGGGGATTCATCATTCTGTTAGCACTGCCGAGCAAACCAAACTGTACGGCTCAGGAAATAAG CTCGTGACGGTAGGGTCATCAAACTACCAGCAGAGCTTCGTACCTTCCCCCGGCGCTAGACCTCAGGTCAACGGTCAGTCCGGACGAAT TGATTTCCACTGGTTAATGCTCAACCCAAATGACACCGTCACATTCAGTTTTAACGGTGCCTTTATTGCTCCTGACAGAGCCTCTTTTCTGA GGGGCAAGTCCATGGGCATCCAGAGCGGGGTACAGGTCGACGCCAATTGCGAGGGCGACTGCTACCATTCAGGGGGGACGATCATTA GTAACTTACCCTTTCAAAACATTGATTCACGGGCCGTAGGCAAGTGTCCGAGATACGTCAAGCAGCGGAGCTTGCTGTTGGCAACTGGC ATGAAAAACGTTCCCGAAATCCCTAAGGGCCGAGGTCTTTTTGGGGCAATTGCGGGCTTCATAGAGAATGGTTGGGAGGGACTGATCGA TGGGTGGTATGGCTTCAGACATCAGAACGCACAGGGTGAGGGTACCGCTGCTGACTATAAGAGCACCCAGAGCGCAATAGATCAGATCA CTGGCAAGCTTAACCGACTTATCGAAAAGACCAACCAGCAGTTTGAATTGATTGACAACGAATTTAACGAAGTTGAGAAGCAAATTGGGAA CGTTATTAATTGGACAAGGGATTCCATCACAGAGGTTTGGAGTTACAATGCGGAGTTATTAGTGGCAATGGAGAACCAACATACTATTGAC CTTGCGGATAGCGAGATGGACAAGCTGTACGAGAGGGTCAAGCGCCAGTTACGGGAAAACGCGGAAGAGGATGGAACTGGATGTTTCG AGATCTTCCACAAATGCGATGACGATTGCATGGCGTCGATCCGAAACAATACCTACGATCACAGTAAATACAGGGAAGAGGCAATGCAGA ACCGAATTCAGATTGATCCCGTGAAGCTCAGCTCAGGGTACAAGGACGTGATTCTGTGGTTTTCATTTGGAGCTAGCTGCTTTATACTGCT GGCTATTGTTATGGGCTTGGTATTCATCTGTGTGAAGAATGGTAACATGAGGTGTACAATTTGCATT
2734	ATGAATACTCAGATCCTGGTGTTTGCTCTGATAGCTATCATCCCCACCAACGCCGATAAGATTTGCCTGGGCCACCACGCAGTGAGTAAC GGGACAAAAGTCAACACTCTAACTGAGCGCGGCGTTGAAGTTGTAAATGCCACTGAAACTGTTGAGAGAACTAATATACCCCGCATTTGC TCAAAAGGCAAGAGGACTGTGGATCTCGGCCAGTGTGGGCTTCTGGGGACCATAACTGGTCCGCCCCAATGTGACCAGTTCTTGGAGTT TTCCGCAGACCTAATTATCGAGCGTCGCGAAGGCTCCGATGTGTGCTACCCCGGCAAGTTTGTCAATGAAGAAGCCCTGAGGCAGATTC TGCGAGAAAGCGGGGGTATAGACAAAGAAGCTATGGGCTTTACTTATAGCGGAATTCGTACAAACGGCGCTACATCCGCCTGTCGACGG TCTGGTTCTAGCTTCTATGCTGAGATGAAATGGTTGCTTAGCAACACAGACAATGCCGCTTTCCCCCAGATGACTAAAAGTTATAAAAATA CCCGGAAGAGCCCCGCACTTATCGTATGGGGGATCCATCACAGTGTGTCGACTGCCGAACAGACCAAGCTCTACGGGAGCGGCTCCAA GCTGGTGACAGTGGGGTCTTCTAATTACCAACAATCCTTTGTCCCTTCCCCTGGCGCCCGCCCCCAAGTGAATGGGCTAAGCGGAAGGA TTGACTTTCATTGGCTGATGCTGAATCCCAACGATACAGTGACGTTCAGCTTTAACGGGGCGTTTATTGCCCCTGACCGTGCAAGTTTCT CCGCGGGAAATCCATGGGCATCCAAAGCGGCGTCCAAGTGGATGCAAACTGCGAAGGAGACTGCTATCACTCTGGAGGAACTATTATTT CTAACCTGCCTTTCCAGAACATCGACTCAAGGGCCGTGGGGAAATGCCCGCGCTATGTTAAACAGAGAAGCCTGTTGCTTGCCACAGGT ATGAAGAACGTGCCCGAGATCCCTAAGGGCAGAGGGCTGTTCGGCGCAATCGCAGGGTTCATTGAAAACGGATGGGAGGGCCTCATCG ACGGATGGTACGGTTTTCGGCACCAAAACGCACAGGGGGAGGGCACAGCGGCGGATTATAAGAGCACTCAGAGTGCCATAGATCAGAT TACGGGGAAGCTCAATAGACTTATAGAGAAGACCAATCAGCAATTCGAGCTGATCGATAACGAGTTCAATGAGGTAGAAAAGCAGATTGG GAATGTCATCAACTGGACAAGAGATAGTATAACAGAGGTGTGGAGCTACAACGCCGAGTTACTTGTGGCCATGGAAAACCAGCACACAAT CGACCTGGCTGATAGCGAAATGGACAAGCTGTATGAAAGAGTCAAGAGACAGCTTCGAGAAAACGCAGAGGAGGACGCAACCGGCTGC TTCGAGATTTTTCACAAATGTGACGATGATTGCATGGCGTCAATCCGGAACAACACCTATGACCACTCTAAGTATCGAGAGGAGGCGATG CAGAACCGGATACAGATCGATCCTGTTAAGCTTTCATCAGGTTACAAGGACGTAATCCTGTGGTTCTCGTTCGGCGCATCCTGCTTTATCC TTCTGGCAATTGTGATGGGGCTCGTATTCATCTGCGTGAAAAATGGCAATATGAGATGTACTATTTGCATT

2735	ATGAATACACAGATCCTTGTGTTCGCACTGATCGCGATTATCCCCACCAACGCAGACAAGATCTGTTTAGGCCACCATGCTGTGAGCAAT GGGACAAAAGTCAACACGCTGACAGAGCGGGGAGTGGAGGTGGTAAACGCTACTGAAACGGTCGAGAGAACTAACATTCCTAGAATTTG TTCAAAAGGGAAAAGGACCGTGGATCTGGGCCAGTGTGGGCTGCTGGGAACAATCACTGGGCCACCACAATGCGATCAGTTTTTAGAGT TCTCGGCCGACCTAATCATAGAGCGCAGAGAGGGCTCCGATGTATGCTATCCCGGGAAGTTTGTGAATGAAGAAGCTTTAAGGCAGATT CTTAGAGAATCCGGTGGGATTGACAAGGAGGCAATGGGGTTTACATACTCCGGTATTAGAACAAACGGCGCCACCTCCGCGTGCAGGAG AAGCGGCTCCTCTTTTTATGCTGAGATGAAATGGCTGCTGTCAAACACCGACAACGCCGCATTTCCCCAAATCACCAAGTCCTACAAGAA TACAAGAAAGTCTCCAGCCCTGATTGTTTGGGGGATACACCACTCAGTGTCCACCGCTGAACAGACCAAGTTGTATGGTAGCGGAAACAA GCTGGTGACCGTCGGGTCCTCAAATTATCAGCAGTCATTTGTCCCATCTCCGGGCGCTAGGCCTCAGGTGAATGGGCTGAGTGGACGCA TTGACTTCCACTGGCTTATGCTGAATCCTAATGACACAGTTACATTTAGTTTCAACGGGGCATTTATTGCGCCAGATAGAGCCAGTTTCTT GCGCGGCAAATCTATGGGAATCCAGAGCGGCGTTCAAGTGGACGCCAACTGTGAGGGGGACTGCTACCACAGCGGCGGGACAATTATC AGCAACCTGCCATTCCAGAATATTGATTCACGGGCTGTGGGAAAGTGTCCTAGGTACGTCAAACAGCGGTCTCTGCTGTTAGCAACCGG AATGAAGAATGTACCCGAGATCCCTAAAGGCCGAGGGTTGTTCGGCGCAATTGCGGGCTTCATTGAGAATGGTTGGGAGGGTCTCATCG ATGGTTGGTACGGCTTCAGACACCAGAATGCCCAAGGCGAAGGAACTGCAGCCGACTATAAGTCAACTCAAAGCGCAATCGACCAGATC ACAGGGAAACTGAATCGCTTGATCGAAAAGACCAACCAGCAATTCGAGCTGATCGACAATGAGTTTAACGAGGTCGAGAAGCAGATCGG AAATGTGATTAATTGGACTCGAGACTCAATTACCGAGGTTTGGAGTTACAATGCCGAACTGCTGGTGGCCATGGAGAATCAACATACCAT AGATCTCGCCGATAGCGAGATGGATAAACTGTATGAACGAGTTAAGCGACAACTGAGAGAAAACGCCGAAGAAGACGGAACCGGGTGCT TTGAAATCTTCCACAAATGCGATGATGATTGTATGGCTTCAATTCGCAACAACACCTACGATCACAGCAAGTACCGAGAAGAGGCCATGC AAAACCGGATTCAAATCGATCCAGTCAAGCTGAGTTCTGGATATAAGCATGTGATTCTGTGGTTTTCATTTGGGGCTAGCTGTTTCATTCT TCTTGCCATCGTGATGGGTCTCGTGTTTATTTGTGTCAAGAATGGGAACATGCGCTGCACGATATGCATC
2736	ATGAACACCCAAATTCTAGTTTTCGCCCTTATCGCCATCATACCGACAAACGCAGACAAAATATGTCTCGGACACCACGCCGTGAGTAAT GGCACCAAGGTGAACACTCTTACCGAGCGAGGGGTGGAGGTGGTGAATGCAACAGAAACCGTGGAGCGCACCAACATCCCCCGGATTT GTTCAAAGGGAAAAAAGACAGTCGACCTGGGGCAGTGTGGGCTCCTGGGCACTATCACTGGGCCCCCGCAGTGCGACCAATTCCTGGA GTTTAGCGCAGACCTGATCATCGAAAGAAGGGAGGGGTCAGACGTGTGTTACCCGGGAAAGTTCGTGAACGAAGAGGCACTTCGCCAG ATATTAAGAGAGAGCGGCGGGATTGATAAAGAAGCTATGGGCTTCACCTATTCTGGCATTAGGACCAATGGCGCCACAAGTGCCTGTAG GAGATCAGGCTCGAGCTTCTACGCTGAGATGAAATGGCTCCTGTCGAACACAGACAACGCCGCTTTTCCTCAGATGACGAAATCCTATAA AAACACGCGGAAGTCTCCTGCGCTCATTGTGTGGGGAATTCATCACAGTGTGTCTACCGCTGAGCAGACAAAGCTTTACGGTTCCGGAA ATAAATTAGTGACAGTTGGCTCTAGTAACTATCAGCAAAGTTTCGTACCATCCCCGGGAGCGAGACCGCAGGTAAATGGACAGTCCGGG CGGATAGATTTCCATTGGCTCATGCTGAATCCTAATGACACCGTCACATTCAGCTTTAACGGGGCATTCATCGCACCCGATCGAGCTAGT TTTCTTCGCGGCAAGTCTATGGGAATTCAATCTGGCGTACAGGTGGATGCCAATTGCGAGGGTGATTGCTACCACTCTGGCGGTACCATC ATCTCGAACCTCCCTTTTCAAAACATCGATAGCAGAGCCGTGGGCAAGTGCCCTCGATATGTTAAGCAGAGAAGCCTGCTGCTCGCAACC GGCATGAAGAACGTGCCTGAGATTCCTAAGGGCCGCGGCTTATTCGGCGCCATTGCTGGCTTCATCGAAAATGGCTGGGAAGGTTTAAT CGATGGATGGTATGGGTTCCGCCATCAGAACGCTCAAGGTGAGGGCACTGCCGCCGATTACAAAAGCACCCAGAGTGCAATCGACCAG ATTACTGGTAAGCTTAATCGACTAATTGAGAAAACCAACCAGCAGTTCGAACTGATTGATAACGAGTTTAATGAAGTCGAGAAGCAGATTG GAAACGTTATCAATTGGACCCGAGACAGTATAACTGAGGTGTGGAGCTACAACGCCGAGCTCCTCGTGGCTATGGAGAACCAACATACA ATCGATCTTGCAGATTCTGAAATGGATAAGCTATACGAAAGAGTAAAGCGACAGCTCCGGGAAAACGCAGAGGAAGACGGAACTGGATG TTTCGAGATTTTTCACAAGTGCGACGACGATTGCATGGCAAGCATTAGAAATAACACATACGACCACTCAAAGTACCGCGACGAGGCGAT GCAGAATAGGATACAGATCGACCCTGTAAAGCTGAGCTCTGGGTACAAGGACGTGATCTTGTGGTTTTCTTTCGGCGCATCATGCTTCAT CTTGCTGGCAATCGTGATGGGCCTGGTCTTTATTTGCGTAAAGAACGGCAATATGCGCTGTACAATCTGTATC
2737	ATGAATACTCAGATACTGGTGTTCGCACTCATCGCAATTATACCCACTAATGCCGATAAGATTTGTTTAGGACATCACGCCGTCAGTAACG GAACTAAGGTTAACACACTAACAGAGCGAGGCGTCGAAGTGGTTAATGCAACTGAGACGGTCGAGCGCACAAACATCCCTCGTATTTGC AGCAAAGGAAAGAAAACCGTCGATCTGGGACAGTGCGGACTCCTGGGGACCATAACCGGCCCTCCCCAGTGCGATCAGTTTCTGGAGTT CTCAGCAGATCTTATAATCGAGCGACGGGAAGGTTCCGACGTATGTTATCCCGGTAAATTCGTCAACGAAGAAGCCCTGAGACAGATCCT GCGGGAGAGTGGAGGAATTGACAAGGAGGCCATGGGATTTACCTACAGCGGCATTAGAACGAATGGGGCCACATCTGCCTGCCGACGC AGCGGCAGCTCATTCTACGCCGAGATGAAATGGTTACTGTCCAATACCGACAACGCCGCGTTCCCGCAGATGACCAAATCTTACAAGAAC ACTCGAAAATCCCCCGCACTTATTGTGTGGGGCATCCACCATAGTGTGTCAACCGCTGAGCAGACTAAACTCTACGGTTCTGGTAACAAA CTCGTGACTGTGGGAAGCTCAAACTACCAGCAAAGCTTTGTGCCCTCTCCCGGGGCCAGGCCACAGGTGAACGGACAGTCCGGACGCA TAGACTTCCACTGGCTGATGCTAAATCCGAACGACACCGTCACCTTCAGTTTTAATGGGGCTTTCATCGCCCCTGACAGAGCTAGCTTCC TGCGCGGTAAGAGCATGGGTATTCAGAGCGGGGTGCAGGTCGACGCCAACTGCGAAGGTGATTGCTATCATTCAGGGGGCACAATTAT CTCCAACCTCCCATTTCAGAACATCGATAGCCGCGCAGTCGGTAAGTGTCCACGATACGTTAAACAAAGGTCACTGCTCCTTGCCACTGG CATGAAAAACGTCCCGGAGATCCCGAAGGGCCGCGGACTCTTTGGGGCCATCGCTGGGTTCATTGAAAATGGTTGGGAGGGGCTCATT GACGGATGGTATGGCTTTAGGCACCAGAACGCCCAAGGAGAGGGGACCGCAGCCGATTATAAGAGCACACAGTCTGCCATTGACCAGA TAACCGGTAAACTTAACCGGTTAATCGAAAAAACTAACCAGCAGTTCGAATTAATCGACAACGAATTTAATGAAGTCGAGAAACAGATCGG CAACGTCATCAACTGGACCCGAGATAGCATAACTGAAGTGTGGTCCTACAACGCCGAGCTGCTCGTGGCTATGGAGAATCAGCATACCA TCGACTTGGCCGACAGTGAAATGGATAAGCTGTACGAGAGAGTTAAGCGCCAGCTGAGACAGAACGCAGAAGAGGACGGGACCGGCTG CTTTGAGATTTTCCATAAGTGCGATGACGACTGTATGGCAAGCATTCGCAATAATACTTACGATCACTCCAAATATAGGGAGGAGGCCATG CAAAACCGCATTCAGATTGATCCGGTGAAGTTGTCAAGTGGATATAAGGATGTGATTCTCTGGTTCTCCTTCGGGGCCTCCTGTTTTATTC TTCTCGCAATCGTTATGGGCTTGGTCTTCATCTGCGTAAAAAATGGGAATATGAGATGTACTATCTGTATC

2738	ATGAACACACAAATTCTTGTATTTGCCCTTATAGCAATCATTCCAACCAACGCAGACAAGATATGCCTGGGGCACCACGCCGTCTCTAACG GTACAAAGGTAAACACACTAACCGAACGCGGTGTCGAGGTAGTGAATGCAACTGAGACCGTGGAGCGGACCAACATACCAAGAATCTGC TCCAAAGGGAAGAGAACGGTGGATCTAGGACAGTGCGGGCTGCTCGGGACTATCACTGGACCACCTCAGTGTGATCAATTTTTAGAGTT CTCGGCAGACTTAATCATCGAACGGCGGGAAGGATCTGATGTTTGCTACCCTGGCAAGTTTGTTAATGAAGAGGCGCTGCGACAGATTTT GCGCGAATCCGGTGGTATCGATAAAGAAGCAATGGGCTTCACTTATAGTGGCATTCGGACCAATGGAGCCACAAGCGCATGCCGAAGGA GCGGGTCCAGCTTTTACGCTGAAATGAAATGGCTCTTGAGTAATACAGATAACGCCGCCTTTCCTCAAATGACAAAAAGCTATAAGAATAC GCGAAAAAGCCCTGCTCTAATCGTCTGGGGTATCCACCACTCTGTGTCTACAGCCGAACAAACCAAGCTGTATGGCAGCGGCTCAAAGC TGGTAACAGTGGGATCAAGCAATTACCAGCAGAGTTTTGTGCCTAGCCCTGGAGCTAGACCGCAGGTGAACGGCCTTTCAGGTCGGATA GACTTTCACTGGCTTATGTTAAATCCAAATGACACGGTGACCTTCTCTTTCAACGGGGCATTTATCGCACCAGATAGGGCATCTTTTCTGC GGGGGAAGAGTATGGGCATCCAATCCGGAGTGCAGGTTGATGCAAATTGCGAGGGGGATTGTTATCACTCCGGCGGTACAATAATTAGT AACCTCCCATTTCAGAATATTGATAGTCGTGCCGTCGGAAAATGTCCTCGCTATGTCAAGCAGAGATCGCTGTTACTTGCCACCGGCATG AAAAATGTGCCCGAAATCCCAAAGGGCCGAGGTCTGTTCGGGGCTATCGCAGGCTTCATTGAAAATGGATGGGAAGGCCTGATCGACGG GTGGTACGGCTTCCGTCATCAGAACGCACAGGGGGAAGGGACTGCAGCGGATTATAAGTCCACCCAAAGCGCTATTGACCAGATCACC GGCAAGCTCAATCGGCTCATAGAAAAAACTAATCAACAGTTTGAGTTGATAGATAATGAGTTTAACGAGGTCGAGAAGCAGATTGGGAAT GTGATCAATTGGACCAGAGACTCAATAACTGAGGTGTGGTCATACAATCGCGGAGTTGTTGGTCGCTATGGAAAATCAACACACAATCGAC CTGGCTGACTCAGAAATGGATAAGCTGTACGAGCGGGTCAAGAGACAGCTGCGCGAGAATGCCGAGGAAGATGGAACAGGTTGCTTTG AAATTTTCCATAAATGCGATGATGACTGTATGGCAAGTATTCGGAACAATACTTACGATCACAGTAAGTATAGAGAAGAAGCAATGCAAAA CAGAATTCAGATTGATCCGGTGAAGCTGAGCTCAGGATACAAAGATGTTATCCTATGGTTTTCTTTTGGGGCCTCATGCTTTATCTTATTG GCCATAGTTATGGGGCTAGTATTTATCTGCGTAAAATCAAGGAATATGAGGTGCACCATCTGCATT
2739	ATGAATACACAGATCTTGGTTTTTGCCCTCATCGCCATTATCCCCACAAATGCCGACAAAATTTGCTTGGGCCACCACGCCGTGTCCAATG GTACAAAAGTCAATACACTGACCGAGAGGGGAGTGGAGGTGGTGAATGCCACAGAAACCGTTGAAAGAACGAATATTCCTAGAATTTGTT CTAAAGGGAAAAGAACCGTTGATCTCGGGCAATGCGGTCTGTTAGGCACAATCACAGGCCCTCCCCAGTGTGATCAATTTCTTGAGTTCA GTGCCGGATCTTATCATTGAGAGACGGGAAGGGTCTGATGTCTGTTACCCCGGCAAGTTTGTGAACGAGGAAGCCCTGCGCCAGATCCTC AGAGAATCTGGGGGTATCGATAAGGAAGCTATGGGGTTTACTTATTCCGGGATCAGGACTAATGGGGCCACCAGCGCCTGCCGGCGGT CCGGCAGCTCATTCTACGCCGAGATGAAGTGGCTGCTGTCCAATACCGACAATGCGGCCTTTCCGCAGATGACCAAGAGCTATAAAAAC ACGCGCAAATCACCCGCTTTAATCGTCTGGGGGATACATCATTCAGTGAGCACTGCCGAGCAAACCAAGCTGTATGGGAGCGGCAGTAA ATTGGTTACTGTAGGTAGCTCTAACTACCAGCAGTCTTTCGTCCCAAGCCCAGGAGCCCGTCCGCAGGTGAATGGACTCAGTGGAAGGA TTGATTTTCATTGGCTAATGCTCAACCCGAACGATACTGTGACGTTTTCATTTAATGGCGCATTTATCGCGCCCGATCGAGCCAGCTTCCT CCGGGGTAAGAGTATGGGGATCCAATCAGGAGTGCAGGTCGATGCTAACTGCGAGGGAGACTGTTACCATTCCGGCGGCACTATTATCA GCAATCTGCCCTTCCAGAATATCGACTCCAGAGCCGTGGGAAAGTGTCCAAGGTACGTGAAACAGAGGAGCTTACTGCTTGCCACAGGG ATGAAGAATGTCCCCGAGATTCCCAAAGGAAGGGGGCTGTTCGGAGCCATCGCAGGGTTCATTGAGAACGGGTGGGAAGGCTTAATCG ATGGCTGGTACGGCTTCAGACACCAAAATGCACAGGGCGAAGGAACCGCCGCCGATTACAAGTCAACGCAATCCGCTATAGATCAGATA ACCGGAAAGCTGAACAGACTCATCGAAAAAACAAACCAACAGTTCGAACTTATAGACAACGAATTCAACGAGGTGGAAAAACAGATCGGG AACGTTATCAATTGGACACGAGACTCAATCACAGAGGTATGGTCCTACAACGCTGAACTCCTGGTTGCGATGGAGAATCAGCACACCATT GACTTAGCCGACTCTGAGATGGATAAACTGTACGAGAGAGTGAAAAGGCAGCTCCGAGAGAACGCAGAAGAGGATGGTACAGGATGCTT CGAAATCTTTCACAAATGCCACGATGATTGTATGCCCAGCATCAGAAATAATACATATGACCATTCCAAATATCGGGAAGAAGCAATCCAA AACCGGATCCAGATCGATCCTGTTAAACTCAGCAGCGGTTACAAGGACGTGATTCTGTGGTTCTCCTTCGGCGCATCATGTTTTATTCTTC TTGCTATTGTGATGGGACTAGTATTTATATGCGTGAAGAATGGCAACATGAGATGTACCATTTGCATC
2740	ATGAATACCCAAATCCTTGTATTCGCTCTGATCGCAATCATTCCAACCAATGCTGATAAAATCTGCTTAGGACACCACGCTGTGAGCAATG GTACAAAGGTCAATACACTGACAGAGCGGGGCGTAGAAGTGGTCAACGCCACCGAGACGGTGGAAAGGACCAATATCCCTAGGATTTGC TCGAAGGGCAAGAGAACAGTAGACCTGGGGCAATGTGGATTGTTGGGGACTATTACGGGACCGCCCCAGTGTGATCAGTTTCTGGAGTT CAGCGCTGATCTCATCATCGAGCGCCGAGAGGGATCCGACGTGTGCTACCCAGGAAAATTCGTTAACGAGGAAGCATTAAGACAGATTT TACGTGAATCGGGTGGCATTGATAAAGAAGCCATGGGGTTTACTTACAGTGGAATTAGAACCAATGGAGCCACCTCGGCTTGCCGGAGA TCTGGGTCTAGCTTTTACGCAGAAATGAAGTGGCTGCTGTCTAACACGGATAATGCCGCGTTCCCGCAGATGACCAAAAGTTATAAAAAT ACCAGAAAGAGCCCTGCATTAATTGTCTGGGGCATTCACCACTCTGTTTCCACGGCCGAACAGACAAAATTGTATGGAAGCGGAAACAAG CTGGTCACAGTCGGATCTAGCAACTACCAGCAGAGCTTTGTGCCCAGTCCAGGGGCCCGCCCTCAGGTCAATGGGCTTTCTGGGAGAAT TGATTTTCACTGGCTGATGTTGAACCCCAACGACACTGTGACCTTTAGCTTCAATGGCGCGTTTATTGCGCCCGACAGAGCCTCCTTTTTA CGGGGGAAAAGCATGGGAATACAATCTGGTGTGCAGGTGGATGCAAATTGCGAAGGAGACTGCTATCACTCCGGGGGTACCATTATTAG CAATCTCCCATTCCAGAACATCGACTCCAGAGCCGTTGGTAAATGCCCCAGGTATGTCAAGCAGCGTAGTCTCTTGTTGGCCACAGGAAT GAAAAACGTGCCTGAGATCCCCAAAGGGCGCGGCCTATTTGGGGCAATTGCCGGCTTTATCGAGAACGGATGGGAGGGCCTGATCGAC GGTTGGTATGGCTTTCGCCATCAGAATGCTCAGGGGGAAGGCACAGCTGCAGACTACAAGAGCACCCAGAGCGCCATCGATCAAATCAC CGGCAAGCTCAACCGGATCATCGAGAAGACAAATCAGCAATTTGAATTAATCGACAACGAGTTCAATGAGGTAGAAAAGCAGATAGGAAA CGTGATCAACTGGACCAGGGATAGCATTACTGAAGTCTGGTCATACAATGCCGAACTCCTGGTGGCAATGGAGAACCAGCACACTATTGA TCTTGCCGACTCTGACATGGACAAACTGTACGAGCGCGTGAAGCGACAGCTGAGGGAAAACGCTGAGGAAGACGGTACTGGCTGCTTT GAAATCTTCCACAAATGCGACGACGATTGCATGGCATCCATACGGAACAACACTTATGATCACTCAAAGTATCGGGAGGAAGCAATGCAG AATAGAATTCAAATCGATCCCGTTAAACTGAGCTCCGGCTACAAGGATGTGATTCTCTGGTTTAGCTTCGGAGCATCATGCTTTATCCTCC TAGCAATTGTAATGGGTCTGGTGTTCATATGCGTAAAGAATGGCAATATGCGCTGTACTATCTGTATT

2725	ATGAACACGCAAATTCTAGTGTTTGCTCTAATTGCCATCATACCCACAAACGCCGATAAAATCTGCCTTGGGCATCACGCAGTAAGTAACG GTACCAAGGTGAACACGCTCACAGAGCGCGGGGTCGAGGTCGTGAATGCCACCGAAACAGTGGAACGCACTAACATCCCTCGAATCTG TTCAAAAGGTAAGCGGACTGTGGATCTGGGACAGTGTGGCCTGCTTGGTACGATCACCGGTCCTCCCCAATGCGATCAGTTCCTGGAGT TCTCAGCAGACTTAATCATCGAGCGCAGAGAGGGTAGCGATGTCTGTTACCCCGGAAAGTTTGTGAAGGAAGAAGCCCTTCGCCAGATT CTCACAGAGTCAGGAGGCATTGATAAGGAAGCCATGGGGTTTACCTACAGCGGTATCCGTACCAACGGAGCCACCTCTGCCTGCCGAC GCTCAGGTTCTAGCTTCTACGCTGAAATGAAATGGTTACTGAGCAATACCGATAACGCCGCATTCCCCCAGATGACAAAAAGCTACAAGA ATACCCGCAAAAGCCCTGCCCTTATAGTTTGGGGGATACATCACTCCGTGTCCACCGCAGAACAGACCAAGCTGTACGGCAGCGGCAAC AAATTGGTGACCGTGGGGAGTAGTAATTATCAACAGAGCTTTGTGCCCTCGCCTGGGGCCAGACCTCAGGTCAATGGACTTAGTGGAAG GATTGATTTCCACTGGCTTATGCTCAATCCTAATGATACCGTCACATTCAGCTTCAACGGGGCGTTCATCGCACCCGATCGCGCAAGCTT CCTGCGTGGGAAGAGTATGGGCATCCAATCTGGGGTTCAAGTGGATGCAAATTGTGAGGGCGATTGTTACCACAGCGGCGGGACGATC ATCTCCAACCTCCCCTTCCAGAATATTGATTCTCGTGCCGTTGGGAAATGTCCTAGATATGTGAAGCAGCGTTCCCTCCTGCTGGCTACA GGCATGAAGAACGTTCCGGAAATCCCCAAGGGCAGAGGACTGTTTGGCGCCATTGCAGGATTTATTGAGAATGGATGGGAGGGCCTCAT CGATGGTTGGTACGGATTCAGACACCAGAACGCTCAGGGGGAAGGAACAGCCGCCGATTATAAGTCTACCCAGTCGGCTATCGACCAAA TCACCGGAAAACTGAACAGGCTGATAGAGAAAACTAACCAGCAGTTCGAGTTGATCGACAACGAGTTCAACGAAGTGGAGAAACAGATC GGCAACGTCATTAACTGGACACGCGACTCAATTACAGAGGTGTGGTCCTACAACGCTGAGCTGCTGGTAGCAATGGAGAACCAGCATAC CATCGACCTGGCTGATTCTGAAATGGATAAGCTGTATGAACGGGTCAAGCGTCAACTGCGCGAAAACGCCGAAGAGGATGGAACAGGCT GCTTCGAAATTTTTCATAAATGCGATGACGACTGCATGGCAAGCATCCGGAACAACACATACGATCACTCAAAATACCGCGAGGAGGCTA TGCAGAATCGCATTCAGATTGATCCTGTGAAGTTATCCTCCGGCTACAAAGATGTCATCCTCTGGTTCAGTTTCGGCGCGTCTTGCTTCAT CCTCCTAGCAATCGTGATGGGTCTCGTTTTCATCTGTGTGAAGAACGGGAATATGAGATGCACTATTTGCATA
2726	ATGAATACCCAGATCCTTGTGTTTGCTCTGATCGCCATCATTCCCACAAACGCCGACAAAATTTGCCTCGGTCACCATGCTGTGTCAAACG GTACTAAAGTCAACACATTGACAGAGAGGGGCGTGGAGGTGGTGAATGCAACCGAGACCGTTGAGCGAACAAATATCCCTCGCATCTGC TCCAAGGGCAAGAAAACTGTGGATTTGGGCCAATGTGGGCTTTTGGGAACCATTACTGGGCCTCCTCAGTGTGATCAGTTCCTCGAGTTC TCTGCCGACCTGATTATTGAACGCCGACAAGGTAGCGATGTCTGTTACCCAGGCAAATTCGTCAACGAAGAGGCACTTCGCCAGATACT GAGAGAGAGTGGAGGCATTGATAAAGAAGCTATGGGGTTTACCTATTCTGGGATAAGGACCAACGGGGCCACCAGTGCTTGTAGACGTA GTGGCAGCAGCTTCTACGCGGAAATGAAATGGCTCCTGTCAAATACAGACAATGCTGCTTTTCCTCAGATGACTAAATCGTACAAAAACA CGCGGAAGAGCCCCGCCTTAATCGTGTGGGGAATCCACCACTCCGTGTCCACAGCGGAACAGACCAAACTTTACGGCTCCGGGAATAA GCTAGTGACCGTGGGCAGCTCCAACTATCAACAAAGTTTCGTCCCTTCCCCCGGTGCGAGACCGCAGGTCAACGGACAGTCAGGGAGG ATCGATTTCCACTGGCTGATGTTAAACCCTAATGACACTGTGACTTTCTCCTTCAATGGAGCGTTCATTGCACCGGATCGAGCGTCTTTTC TCCGGGGAAAGTCTATGGGTATCCAGTCTGGCGTGCAGGTGGATGCCAATTGCGAGGGAGATTGCTACCACTCCGGGGGCACTATCAT CTCGAATCTCCCCTTCCAGAACATCGACAGCCGGGCCGTGGGTAAATGCCCACGGTATGTCAAGCAAAGATCCTTACTGCTCGCCACAG GGATGAAGAACGTGCCTGAGATCCCTAAGGGCCGGGGGCTGTTTGGAGCCATCGCTGGGTTCATTGAAAACGGTTGGGAAGGCCTCAT TGACGGATGGTATGGCTTCAGGCACCAGAACGCTCAGGGGGAGGGGACCGCCGCCGACTACAAGAGTACGCAATCTGCTATAGATCAG ATCACTGGTAAACTCAACCGCTTGATCGAGAAAACCAATCAGCAGTTCGAGCTTATTGACAACGAGTTTAACGAGGTAGAGAAGCAGATT GGAAATGTCATCAATTGGACTAGAGACTCTATCACAGAGGTGTGGAGCTATAATGCCGAATTGTTGGTGGCAATGGAAAACCAGCACACC ATCGACCTGGCCGACAGCGAAATGGACAAGTTGTATGAACGCGTTAAACGTCAGTTGAGGGAGAACGCCGAGGAGGACGGGACGGGTT GCTTTGAAATCTTTCATAAGTGCGATGACGACTGTATGGCTTCTATCCGCAATAACACTTATGACCACAGTAACTACCGGGAAGAAGCTAT GCAAAACAGAATACAGATCGACCCAGTTAAATTATCAAGTGGCTATAAAGATGTGATCCTCTGGTTCAGCTTTGGAGCCTCCTGTTTCATT TTATTGGCCATAGTCATGGGCCTCGTATTTATCTGCGTGAAAAATGGCAACATGCGCTGCACCATCTGCATT
2727	ATGAACACTCAGATCCTAGTGTTTGCTCTGATAGCGATCATACCCACCAATGCAGACAAAATCTGTCTGGGCCATCATGCAGTCTCTAATG GAACCAAAGTGAACACCCTAACTGAGAGGGGCGTCGAGGTGGTGAATGCGACCGAAACTGTTGAAAGGACCAATATTCCCAGAATCTGC TCAAAGGGAAAAAAAACGGTGGACCTGGGGCAGTGTGGATTACTAGCCACGATCACCGGGCCTCCTCAGTGTGATCAATTTCTGGAGTT CAGCGCAGATCTTATTATTGAGAGAAGAGAGGGCTCTGACGTGTGCTACCCAGGCAAGTTCGTGAACGAGGAGGCTCTGCGGCAGATCC TACGTGAGAGCGGAGGCATAGACAAGGAGGCAATGGGATTTACCTATAGCGGGATACGGACAAATGGCGCAACGTCCGCATGTCGCCG GAGTGGGTCTTCCTTTTACGCGGAGATGAAGTGGCTGCTGAGCAACACCGACAATGCCGCGTTTCCTCAGATGACAAAATCATATAAGAA CACCCGAAAGTCCCCAGCACTCATTGTTTGGGGGATCCACCACAGTGTAAGCACCGCCGAGCAGACTAAACTATACGGCAGCGGAAATA AACTTGTCACCGTCGGCTCCTCCAATTACCAACAGTCCTTCGTCCCTTCACCGGGTGCCAGGCCCCAGGTGAACGGACAATCCGGGCGT ATTGACTTTCACTGGCTGATGTTGAATCCAAACGACACCGTGACGTTTTCGTTCAACGGCGCATTTATTGCCCCCGATCGGGCGAGCTTC CTGCGGGGAAAATCCATGGGCATCCAGAGTGGGGTCCAAGTGGACGCAAACTGTGAGGGCGACTGCTACCACTCCGGTGGAACAATCA TTAGCAATCTTCCCTTTCAGAACATTGACTCTCGGGCCGTAGGCAAGTGTCCGCGCTACGTGAAACAGCGCTCTCTGCTATTGGCTACAG GCATGAAGAATGTACCAGAGATACCTAAGGGACGAGGCCTCTTTGGGGCCATCGCCGGCTTCATCGAGAATGGGTGGGAAGGCCTGAT TGACGGTTGGTATGGCTTTCGCCACCAGAATGCACAGGGTGAGGGAACGGCAGCCGATTACAAGTCCACGCAGAGTGCCATCGATCAG ATTACCGGTAAATTAAACCGGCTCATTGAGAAAACTAACCAGCAGTTCGAACTGATCGATAATGAGTTTAATGAGGTGGAGAAGCAAATTG GCAACGTGATTAATTGGACCAGGGACAGTATAACGGAAGTGTGGTCCTATAATGCAGAATTGCTGGTGGCCATGGAAAACCAGCACACG ATCCATCTAGCTGATTCTGAGATCGCACAAACTGTATGAGCGTGTTAAGCGACAGCTTCGCGAAAATGCAGAGGAAGATGGCACAGGCTG CTTTGAGATATTCCACAAATGCGACGATGACTGTATGGCTTCAATACGTAATAATACATACGACCACTCAAAATACCGTGAAGAGGCCATG CAAAATCGCATCCAGATCGATCCTGTGAAACTATCATCAGGCTATAAAGACGTAATTCTTTGGTTTTCTTTCGGCGCAAGTTGCTTCATTCT TCTCGCCATTGTGATGGGACTTGTCTTCATATGTGTGAAAAACGGAAATATGCGCTGCACCATCTGTATC

2728	ATGAACACTCAAATTCTAGTCTTCGCCCTGATCGCAATCATCCCTACTAATGCCGACAAGATCTGTCTGGGACACCACGCAGTTTCAAACG GGACAAAGGTGAACACCCTGACCGAGCGGGGGGTAGAAGTGGTTAACGCAACTGAGACCGTGGAACGTACCAATATACCCAGGATATG TAGCAAGGGTAAAAAGACTGTCGATCTCGGTCAGTGTGGACTCCTGGGAACCATCACCGGACCTCCCCAGTGTGATCAGTTCCTGGAGT TCAGCGCAGATTTAATCATCGAACGTCGCGAAGGATCCGACGTCTGCTATCCGGGTAAGTTTGTAAACGAGGAGGCTCTGAGACAAATC CTCACAGAAAGTGGGGGCATACACAAGGAGGCAATGGGGTTCACTTATTCTGGGATTAGCACAAATGGCGCCACTAGCGCTTGCCGCC GAAGTGGCAGTTCCTTCTACGCCGAGATGAAGTGGTTGCTTAGCAATACTGATAATGCTGCTTTCCCCCAGATGACGAAGTCTTACAAAA ATACTCGTAAGAGCCCCGCGCTGATCGTGTGGGGGATCCACCACAGTGTGAGCACTGCCGAGCAAACCAAGCTGTATGGGTCTGGGAA TAAATTAGTGACCGTGGGCTCTAGTAACTACCAGCAGTCATTCGTTCCAAGTCCCGGCGCACGGCCTCAGGTTAATGGCCAGTCTGGCC GGATTGACTTTCATTGGCTCATGCTGAATCCCAATGACACGGTGACATTCTCCTTTAACGGAGCATTTATCGCACCAGACAGGGCTTCCTT CCTTCGCGGAAAATCCATGGGAATTCAGAGCGGGGTCCAGGTCGATGCTAATTGTGAAGGGGATTGTTACCACTCTGGAGGCACAATTA TTAGCAATCTGCCCTTCCAAAACATCGACTCCCGTGCTGTGGGCAAATGTCCACGCTACGTCAAGCAGAGATCATTACTTCTTGCTACAG GTATGAAAAACGTGCCGGAAATCCCTAAGGGGAGAGGACTTTTCGGTGCCATAGCTGGATTCATAGAGAACGGATGGGAAGGGCTCATT GACGGCTGGTATGGCTTCAGACATCAGAACGCTCAGGGAGAGGGAACTGCTGCTGACTACAAGAGTACCCAGTCAGCTATCGACCAGAT TACTGGGAAATTGAACCGCCTGATTGAAAAGACTAACCAGCAGTTCGAGCTCATCGACAATGAATTCAATGAAGTGGAGAAGCAGATCGG TAATGTCATCAATTGGACTCGGGATTCAATTACAGAGGTGTGGTCTTACAATGCAGAGCTGCTGGTTGCCATGGAAAACCAGCATACCAT TGACCTGGCAGACAGTGAAATGGATAAACTGTACGAGAGGGTTAAACGGCAATTGCGCGAAAATGCCGAGGAAGACGGGACAGGATGTT TTGAGATCTTTCATAAATGCGATGACGACTGTATGGCTTCGATTCGAAACAATACTTACGACCACTCAAAGTATAGAGAAGAAGCAATGCA GAACCGGATTCAAATCGACCCTGTTAAGCTGTCATCCGGCTATAAGGATGTAATTCTCTGGTTCAGCTTTGGTGCATCATGTTTTATACTC CTCGCCATCGTGATGGGCCTTGTATTCATCTGCGTCAAGAACGGGAATATGAGATGCACCATTTGTATT
2729	ATGAACACACAGATCCTCGTTTTTGCCTTGATCGCAATCATCCCAACAAACGCAGACAAAATCTGTCTGGGTCACCACGCCGTCAGTAAC GGGACTAAGGTGAACACCCTCACGGAGCGGGGAGTCGAGGTCGTGAATGCTACTGAGACCGTGGAGCGTACCAACATCCCCAGGATTT GCAGCAAAGGAAAGCGGACCGTAGACCTCGGGCAATGTGGGCTGTTAGGCACAATCACTGGCCCTCCCCAGTGTGACCAGTTTCTTGA GTTCTCTGCTGACTTAATCATCGAGCGTCGAGAAGGGTCAGATGTCTGCTATCCTGGCAAATTTGTGAATGAGGAAGCACTTCCGCAGAT TTTGAGAGAATCGGGCGGCATCGATAAGGAGGCTATGGGCTTCACCTACTCTGGGATTAGGACCAATGGGGCCACCAGTGCCTGTAGG CGCTCAGGCAGTTCCTTTTACGCTGAGATGAAGTGGCTACTGAGTAATACGGATAACGCCGCCTTCCCTCAGATGACTAAGAGCTACAAG AACACCAGAAAATCTCCTGCCTTGATAGTGTGGGGAATACACCACAGCGTATCCACCGCAGAACAAACCAAGTTGTATGGGAGCGGAAA CAAGTTGGTCACCGTTGGATCATCGAATTACCAGCAGAGCTTTGTGCCTTCTCCCGGCGCTCGGCCCCAGGTGAACGGGCTGAGTGGTC GTATCGACTTCCATTGGCTTATGCTTAACCCGAATGACACCGTAACGTTCAGCTTCAATGGCGCTTTTATTGCACCCGACCGCGCGTCATT CCTACGCGGAAAGTCCATGGGGATCCAGTCTGGTGTGCAGGTGGACGCCAATTGTGAGGGCGATTGCTATCACTCCGGAGGGACAATT ATCTCGAATCTCCCGTTCCAGAACATTGACAGCCGGGCTGTCGGGAAGTGCCCGCGCTATGTGAAACAGAGGTCCCTCCTTCTTGCAAC TGGCATGAAAAACGTTCCCGAGATCCCAAAGGGCCGTGGACTGTTTGGAGCCATCGCTGGCTTCATAGAGAACGGCTGGGAGGGCCTG ATTAATGGTTGGTACGGCTTTAGACACCAAAATGCCCAGGGAGAAGGCACCGCCGCAGACTACAAATCGACTCAGTCCGCAATTGATCA GATTACCGGGAAACTCAACCGGCTCATCGAAAAGACTAACCAGCAGTTTGAATTGATTGACAACGAGTTCAATGAGGTGGAGAAGCAGAT CGGGAATGTCATCAACTGGACACGGGACTCAATAACCGAAGTGTGGTCATACAATGCGGAGCTGCTTGTCGCCATGGAAAATCAGCACA CAATCGACCTAGCAGACTCTGAAATGGACAAGCTGTATGAGAGAGTGAAGCGACAATTACGTGAAAATGCGGAGGAGGACGGAACTGGC TGCTTCGAGATCTTCCACAAGTGTGATGACGATTGCATGGCCTCTATCCGCAACAACACTTACGATCATAGCAAATACACCCAGGAAGCC ATGCAGAACCGGATCCAGATTGATCCTGTGAAGCTGTCTAGTGGCTATAAAGACGTAATCCTCTGGTTCAGCTTCGGCGCAAGCTGCTTC ATCCTTCTCGCCATAGTGATGGGCTTAGTGTTTATATGCGTGAAGAACGGGAACATGCGCTGTACCATTTGTATC
2730	ATGAATACTCAAATCCTCGTCTTTGCACTCATAGCAATTATCCCTACCAACGCTGATAAGATTTGTTTGGGGCATCACGCCGTCTCCAACG GGACTAAGGTAAATACATTGACGGAAAGAGGGGTGGAAGTTGTGAATGCCACTGAGACTGTGGAGCGGACCAACATCCCCCGTATATGT TCAAAAGGAAAACGCACAGTCGACTTGGGTCAGTGCGGACTCCTGCGGACTATTACCGGGCCACCTCAGTGCGACCAATTTCTGCAATT CAGCGCGGACCTGATCATTGAGCGCCGAGAGGGAAGTGACGTATGCTACCCTGGGAAGTTTGTTAATGAGGAAGCACTAAGGCAAATAC TGCGGGAATCGGGCGGAATTGATAAGGAAGCTATGGGCTTCACGTACAGCGGAATTAGAACCAACGGGGCTACGAGTGCGTGTAGACG GTCCGGTAGTTCCTTTTACGCCGAAATGAAATGGCTCCTAAGCAATACGGACAACGCCGCATTTCCTCAGATCACAAAATCATACAAGAA CACTAGGAAGAGTCCCGCCCTTATTGTGTGGGGAATTCATCACAGTGTGAGTACTGCCGAACAGACTAAGCTCTATGGCTCAGGAAATAA GCTCGTCACCGTTGGGTCCAGTAACTATCAGCAGTCTTTTGTCCCGTCACCTGGGGCCAGACCTCAAGTCAACGGCTTGAGTGGACGTA TCGATTTTCATTGGCTGATGCTGAACCCAAATGACACTGTCACTTTTAGCTTCAATGGGGCTTTTATCGCTCCAGATCGGGCATCCTTCCT GCGTGGGAAATCCATGGGAATTCAATCAGGTGTCCAGGTGGATGCTAATTGCGAGGGGGACTGCTATCACTCAGGGGGCACTATTATAT CCAATCTGCCCTTTCAGAACATCGATTCCAGGGCGGTGGGCAAATGTCCTAGATATGTAAAACAGAGGTCTCTGCTTCTGGCGACAGGCA TGAAGAATGTTCCTGAAATCCCCAAGGGCAGGGGCCTGTTCGGGGCCATTGCAGGCTTTATCGAGAACGGCTGGGAGGGGCTAATCGA CGGCTGGTACGGCTTTCGACATCAGAACGCTCAGGGTGAAGGCACGGCTGCCGATTATAAGTCGACGCAGTCAGCCATTGATCAAATCA CTGGCAAACTTAATCGCCTCATAGAGAAGACCAATCAGCAGTTTGAACTCATCGATAACGAGTTCAACGAGGTGGAAAAACAAATTGGTA ACGTCATCAATTGGACCAGGGACAGTATTACAGAGGTTTGGTCCTACAACGCCGAATTGTTGGTGGCAATGGAAAATCAGCATACTATCG ACCTGGCTGACAGCGAAATGGACAAGCTCTATGAACGGGTAAAGCAGCCAGTTACGGGAGAACGCTGAGGAGGACGGCACTGGGTGTTT CGAGATCTTTCATAAATGCGATGATGACTGCATGGCTTCTATCCGTAATAACACGTATGATCACTCCAAGTATCGGGAAGAGGCCATGCA GAACCGGATCCAAATCGATCCCGTAAAGTTGTCATCCGGCTACAAAGACGTAATTTTGTGGTTTTCTTTCGGAGCTTCCTGTTTCATACTG CTAGCTATCGTAATGGGGTTGGTGTTCATCTGCGTGAAGAATGGAAACATGAGGTGTACCATCTGTATC

2731	ATGAACACCCAGATCCTCGTCTTCGCTCTAATCGCCATAATCCCCACTAATGCCGACAAAATTTGTCTCGGACACCACGCCGTTAGCAAC GGGACAAAAGTTAACACACTTACTGAGAGAGGGGTCGAGGTTGTAAATGCTACAGAAACCGTGGAGAGGACTAATATTCCAAGAATTTGT TCAAAGGGAAAAAAGACAGTGGATCTTGGCCAGTGCGGGCTGCTGGGGACAATCACAGGACCACCCCAGTGCGACCAATTTCTGGAATT CTCTGCAGACCTGATTATTGAACGCCGTGAGGGGAGTGACGTTTGCTATCCCGGGAAGTTTGTTAACGAAGAGGCTCTTAGACAAATCCT CCGCGAATCTGGGGGGATCGATAAAGAGGCCATGGGGTTTACCTACTCCGGCATTAGGACTAATGGAGCTACCTCCGCCTGTAGGAGGT CTGGAAGCTCATTTTACGCAGAAATGAAATGGTTGCTCAGCAACACCGACAATGCAGCGTTTCCGCAAATGACCAAGAGTTACAAGAACA CCCGAAAGTCTCCTGCGTTGATTGTGTGGGGCATTCACCATTCAGTTTCCACTGCAGAGCAGACAAAACTGTACGGTTCAGGAAACAAGC TGGTAACCGTGGGATCTTCCAACTATCAGCAGAGCTTCGTGCCCTCGCCCGGAGCCCGCCCTCAGGTGAACGGACAGAGTGGAAGGAT CGACTTCCACTGGCTGATGTTGAATCCCAACGATACCGTGACTTTCTCTTTCAACGGTGCATTTATAGCCCCTGACCGCGCTTCCTTTCTC CGAGGAAAAAGCATGGGCATCCAGTCTGGAGTGCAGGTTGACGCAAATTGCGAAGGCGACTGTTACCATAGTGGCGGCACAATAATTTC TAACCTTCCCTTTCAAAATATTGACAGCCGCGCAGTAGGTAAGTGTCCTAGATACGTGAAGCAGAGGAGCCTACTGCTCGCCACTGGCAT GAAAAATGTGCCGGAGATCCCTAAGGGCAGAGGGCTGTTCGGCGCCATCGCCGGCTTCATCGAAAATGGATGGGAGGGGTTGATAGAC GGTTGGTACGGCTTCCGGCATCAGAACGCCCAGGGTGAGGGGACCGCCGCTGATTATAAAAGTACCCAGTCTGCTATCGATCAGATCAC CGGGAAGCTCAACCGTCTTATTGAGAAGACCAATCAGCAATTCGAACTCATCGACAACGAGTTTAACGAGGTTGAAAAGCAAATTGGAAA TGTGATTAATTGGACACGGGATTCAATAACCGAAGTCTGGTCGTACAACGCAGAGCTACTTGTGGCAATGGAGAACCAGCACACCATAGA CCTTGCCGACTCCGAAATGGATAAGTTGTATGAAAGAGTGAAACGCCAGCTGCGGGAGAATGCCGAGGAAGATGGGACAGGATGTTTTG AGATCTTTCACAAGTGCGACGATGATTGCATGGCCTCCATTCGAAACAATACTTATGACCATAGCAAATACAGAGAGGAGGCCATGCAGA ACCGGATTCAAATCGATCCTGTTAAGTTATCTTCAGGTTACAAGGATGTCATCCTTTGGTTCTCGTTTGGAGCCAGCTGCTTCATATTGCT GGCTATCGTGATGGGTCTGGTGTTTATCTGCGTTAAGAACGGCAATATGAGATGTACGATCTGTATT
2732	ATGAACACTCAGATTCTGGTCTTTGCCCTGATTGCTATCATCCCAACCAACGCAGACAAGATCTGTCTGGGACACCATGCCGTATCTAAC GGAACCAAGGTTAACACGTTGACCGAGCGCGGCGTTGAAGTCGTCAATGCCACTGAGACCGTGGAGCGAACAAACATTCCCAGAATCTG TAGCAAGGGCAAACGGACTGTGGACCTGGGCCAGTGTGGGTTGTTGGGAACAATCACAGGACCTCCACAGTGTGATCAGTTCCTGGAGT TTAGTGCAGACCTTATCATCGAACGCAGAGAAGGTTCCGACGTCTGTTACCCTGGGAAATTTGTAAATGAGGAGGCCTTGAGACAGATCC TTAGGGAATCTGGAGGAATCGACAAGGAGGCCATGGGCTTTACCTATAGCGGAATAAGGACCAATGGCGCCACTTCCGCCTGTAGGAGA AGCGGTTCATCATTCTATGCAGAGATGAAGTGGTTGTTAAGTAACACGGATAATGCTGCGTTTCCACAGATGACCAAGTCCTACAAGAATA CCCGAAAATCCCCCGCACTCATTGTCTGGGGAATACACCACAGCGTCTCAACAGCTGAGCAAACAAAACTCTATGGATCCGGGAATAAGC TCGTGACCGTAGGCTCATCAAACTACCAGCAGTCGTTCGTGCCATCTCCAGGGGCCCGGCCTCAGGTCAACGGTCTGTCAGGACGCATT GATTTTCACTGGCTGATGCTTAATCCCAACGATACCGTCACATTCTCCTTCAATGGGGCGTTTATTGCGCCTGACAGAGCTAGCTTTTTGC GAGGTAAGTCTATGGGCATACAGTCTGGGGTGCAAGTGGACGCGAATTGCGAGGGGGACTGCTATCATTCCGGAGGCACCATCATCTC CAACCTCCCATTTCAAAATATCGACAGTAGGGCGGTGGGGAAGTGCCCCCGTTACGTAAAACAGCGCTCCCTCCTGCTGGCCACAGGGA TGAAGAATGTTCCAGAAATACCAAAAGGAAGGGGTCTGTTTGGTGCCATCGCAGGCTTCATCGAAAACGGATGGGAGGGGCTGATTGAT GGTTGGTACGGATTTAGACACCAGAATGCCCAGGGCGAAGGAACTGCCGCTGATTATAAGAGCACACAGTCAGCTATAGACCAGATCAC TGGGAAGCTAAACAGAATCATTGAAAAGACGAACCAGCAGTTCGAACTAATCGACAATGAGTTCAACGAGGTGGAAAAGCAGATTGGGAA CGTGATAAATTGGACACGCGACTCCATAACGGAAGTGTGGAGCTACAATGCGGAGCTGCTGGTGGCTATGGAGAATCAGCACACCATCG ACCTTGCGGATAGTGAAATGGATAAACTCTACGAGCGTGTTAAGCGGCAATTGCGGGAGAACGCCGAGGAGGATGGGACCGGGTGCTT CGAGATTTTCCACAAATGTGATGATGACTGTATGCCCTCCATACGCAATAATACCTACGACCATAGCAAATATCGGGAGGAAGCCATGCA GAACCGAATCCAAATAGACCCAGTGAAGCTGAGTTCCGGCTATAAAGATGTGATTCTCTGGTTTAGCTTTGGAGCAAGCTGTTTTATCCTG TTAGCGATTGTGATGGGTTTAGTGTTCATTTGTGTGAAAAACGGTAACATGCGATGCACCATTTGTATC
2733	ATGAATACCCAGATCCTTGTCTTTGCCCTGATAGCCATTATTCCCACCAACGCAGATAAAATTTGCCTCGGGCACCACGCTGTGAGCAAC GGCACAAAGGTGAATACTCTGACAGAACGGGGTGTAGAGGTGGTTAATGCAACAGAGACAGTGGAGCGCACAAACATCCCCAGAATCTG CTCTAAGGGCAAGCGGACTGTGGATTTAGGGCAATGTGGACTGCTAGGTACTATCACCGGGCCTCCACAGTGTGACCAGTTTTTGGAGT TCTCTGCCGATCTTATCATCGAGCGTCGAGAGGGGAGTGATGTGTGCTACCCAGGCAAATTTGTGAACGAGGAAGCCCTTCGGCAGATC CTGAGAGAGTCCGGCGGCATCGACAAAGAGGCCATGGGTTTTACCTATTCAGGAATCCGAACCAACGGAGCCACCTCCGCCTGTAGGC GGAGTGGGTCTAGCTTTTATGCAGAAATGAAATGGTTGCTGTCCAACACCGATAATGCAGCTTTCCCACAGATAACAAAGTCTTACAAAAA TACGCGCAAGTCCCCAGCGCTAATTGTGTGGGGCATTCATCACAGCGTGTCTACTGCGGAGCAGACCAAGCTGTACGGGAGCGGGAAT AAATTGGTCACAGTCGGGAGCTCGAACTACCAGCAAAGCTTCGTTCCTAGCCCCGGGGCAAGGCCTCAGGTGAACGGACTGAGCGGCA GAATTGACTTCCACTGGTTGATGCTTAATCCTAACGATACAGTGACCTTTAGCTTCAATGGAGCGTTCATTGCCCCTGATCGCGCCTCCTT TCTCCGGGGGAAAAGTATGGGGATTCAATCCGGGGTGCAGGTTGACGCTAATTGTGAGGGTGATTGCTACCACTCCGGGGGCACCATTA TAAGCAATTTACCGTTTCAGAATATTGACTCTCGAGCCGTGGGGAAATGTCCCCGGTACGTCAAGCAAAGGTCTTTGCTGCTGGCTACCG GCATGAAAAACGTACCCGAGATCCCAAAAGGACGCGGCCTCTTCGGTGCAATCGCCGGCTTTATAGAGAATGGATGGGAAGGGCTTATT GACGGGTGGTATGGGTTCCGCCATCAGAATGCGCAAGGCGAAGGAACTGCCGCCGATTACAAGAGTACACAGTCCGCCATCGATCAAAT TACTGGCAAACTAAATAGACTCATCGAGAAGACAAACCAGCAGTTTGAACTCATTGACAACGAATTCAACGAGGTAGAGAAACAGATTGG CAACGTCATCAATTGGACACGGGACAGTATCACAGAAGTCTGGTCCTACAACGCAGAGCTTCTCGTTGCTATGGAAAATCAGCACACCAT TGACCTGGCTGACTCCGAGATGGACAAGTTGTATGAGCGTGTGAAAAGGCAGTTGCGTCAAAACGCCGAGGAAGACGGCACAGGTTGTT TTGAGATCTTTCACAAATGCGACGACGATTGTATGGCAAGCATAAGAAACAATACTTATGACCACAGTAAATATCGAGAGGAAGCCATGCA GAACAGAATCCAAATCGACCCTGTGAAACTAAGCTCAGGCTACAAGGACGTGATCCTCTGGTTCTCCTTTGGTGCTTCCTGCTTTATTCTC CTTGCCATCGTCATGGGCCTCGTGTTCATCTGCGTTAAGAACGGTAATATGCGGTGTACTATTTGTATC

2734	ATGAATACCCAGATTCTCGTTTTTGCCCTAATAGCTATCATTCCTACAAATGCCGATAAAATTTGCTTGGGCCATCATGCCGTGAGCAACG GGACTAAGGTTAACACCCTGACTGAGCGGGGCGTAGAGGTGGTTAACGCCACAGAGACAGTGGAACGGACGAACATTCCAAGAATTTGT TCAAAGGGCAAGAAAACGGTGGACCTCGGGCAATGTGGTCTGCTAGGCACTATAACAGGACCTCCTCAGTGCGATCAGTTTCTGGAATT CAGTGCAGATCTGATCATCGAGCGTCGAGAGGGATCAGATGTCTGTTATCCTGGGAAGTTCGTCAATGAGGAGGCACTGAGACAGATAC TCACAGAGAGCCGAGGCATTGATAAGGAAGCTATGGGTTTTACCTATTCTCGTATTAGAACTAACGGAGCCACAAGTGCCTGTCGTCGGT CGGGCTCTTCGTTTTACGCTGAGATGAAGTGGCTGCTGAGTAATACTGACAATGCAGCGTTCCCCCAGATGACTAAGAGTTATAAAAACA CCCGTAAAAGCCCAGCACTGATCGTGTGGGGAATTCATCATTCAGTCTCTACAGCCGAGCAGACCAAGCTCTACGGGTCTGGCAACAAG CTCGTTACAGTAGGATCTAGCAACTATCAACAGAGCTTTGTACCATCCCCTGGAGCTCGTCCACAAGTGAACGGTCAGTCGGGCCGTATC GACTTTCACTGGCTGATGCTGAATCCTAATGACACGGTGACTTTCAGTTTCAATGGAGCCTTCATCGCACCAGACCGCGCAAGCTTTCTA CGCGGGAAGAGTATGGGGATTCAGTCCGGCGTTCAAGTGGACGCCAATTGCGAAGGGGACTGTTACCATAGCGGCGGCACCATCATTA GTAACCTCCCATTTCAAAACATCGATTCTCGCGCCGTGGGGAAATGTCCTAGATACGTGAAACAGCGCAGCCTGCTCCTCGCAACTGGCA TGAAGAATGTTCCAGAGATTCCGAAGGGGCGGGGTCTCTTTGGCGCCATCGCCGGGTTCATCGAAAACGGGTGGGAGGGACTGATCGA TGGCTGGTATGGCTTCCGACATCAAAACGCACAGGGTGAGGGCACAGCTGCCGATTATAAATCCACGCAGAGCGCAATCGATCAGATCA CCGGCAAACTGAACAGGCTGATCGAGAAAACTAATCAGCAGTTCGAGCTTATCGATAATGAATTCAATGAAGTGGAGAAGCAGATTGGGA ATGTGATTAATTGGACTCGGGACTCTATTACTGAAGTGTGGAGTTACAACGCCGAGCTCTTGGTTGCTATGGAGAATCAGCATACAATTGA CTTGGCAGATTCCGAGATGGATAAGCTCTACGAGCGTGTCAAGCGGCAACTGAGAGAGAACGCCGAAGAGGATGGGACGGGGTGCTTC GAGATTTTCCATAAGTGCGACGATGATTGTATGGCCAGTATCAGAAATAATACATATGACCATAGCAAATACAGAGAGGAAGCAATGCAGA ACCGGATCCAGATTGATCCGGTGAAGCTCTCTAGCGGATATAAGGATGTTATCCTGTGGTTTTCTTTTGGAGCCTCCTGTTTCATTCTGCT GGCCATTGTAATGGGACTAGTCTTCATATGTGTAAAAAACGGCAATATGAGGTGCACTATTTGTATT
2735	ATGAACACACAGATCCTGGTATTCGCGCTCATCGCAATTATACCAACAAATGCCGATAAGATATGCTTGGGACATCACGCCGTATCCAAC GGTACCAAAGTGAATACACTGACCGAACGTGGCGTAGAGGTGGTGAACGCCACTGAGACGGTTGAGCGCACCAACATCCCCCGCATCT GCTCCAAGGGAAAACGCACCGTCGACCTGGGACAGTGTGGGCTGCTCGGCACAATCACGGGTCCTCCTCAGTGTGACCAATTCTTGGA GTTCAGCGCAGATCTAATTATAGACAGGCGGGAGGGAAGCGATGTGTGCTATCCTGGCAAATTTGTCAACGAGGAAGCCCTTCGCCAGA TTCTTCGGGAGTCCGGCGGAATCGATAAAGAGGCTATGGGGTTTACATATAGTGGAATCCGCACCAATGGCGCCACGTCGGCCTGTAGG AGGAGCGGATCCAGTTTCTATGCCGAGATGAAATGGCTCCTCAGTAATACTGATAATGCTGCGTTCCCCCAGATAACCAAATCGTACAAA AATACACGGAAGTCTCCCGCGCTGATTGTATGGGGCATACACCACTCCGTATCGACTGCCGAACAGACTAAGCTCTACGGCTCTGGGAA CAAGTTAGTGACCGTCGGCTCGAGCAATTACCAGCAGAGTTTTGTCCCATCACCAGGGGCAAGACCTCAGGTGAACGGGCTTTCAGGGA GGATTGACTTCCACTGGCTGATGCTGAACCCAAATGACACTGTCACCTTTTCGTTCAACGGGGCATTTATCGCCCCAGACAGGGCCTCAT TCCTTAGAGGCAAATCTATGGGCATTCAAAGCGGGGTTCAGGTGGACGCGAACTGTGAGGGCGACTGTTATCATAGCGGTGGCACTATT ATCTCCAACCTGCCCTTTCAGAATATAGACAGTAGAGCCGTGGGTAAGTGCCCACGGTATGTGAAGCAAAGGTCCCTGCTCCTCGCTAC GGGGATGAAGAATGTGCCAGAGATTCCTAAGGGTCGCGGCCTGTTCGGAGCAATAGCGGGTTTTATAGAAAATGGTTGGGAAGGATTAA TAGATGGGTGGTACGGGTTTAGGCACCAGAATGCTCAGGGAGAGGGTACGGCAGCTGATTATAAGTCTACACAATCTGCCATCGACCAG ATCACCGGAAAGCTCAACAGACTTATCGAAAAAACTAACCAGCAGTTCGAATTGATCGACAATGAATTTAATGAAGTGGAGAAGCAGATAG GTAACGTGATTAACTGGACTCGGGACTCAATCACTGAAGTTTGGAGTTACAACGCAGAGCTCCTGGTCGCCATGGAGAACCAGCATACC ATAGATCTCGCTGACAGTGAGATGGACAAACTGTACGAAAGAGTGAAGCGTCAGCTCAGAGAGAATGCGGAGGAAGATGGGACCGGAT GCTTCGAAATTTTTCACAAGTGTGACGACGACTGTATGGCCAGCATAAGGAATAACACTTATGATCATTCTAAGTATCGTGAGGAGGCAAT GCAGAACAGGATTCAAATTGATCCGGTTAAGCTTTCTTCAGGATACAAGGACGTCATTCTGTGGTTCTCCTTCGGGGCATCTTGCTTTATT TTACTTGCAATCGTGATGGGGTTAGTCTTTATCTGCGTGAAAAATGGGAACATGAGGTGTACTATCTGCATC
2736	ATGAATACACAGATCCTGGTCTTTGCTCTGATCGCAATTATTCCTACGAATGCAGACAAGATTTGCCTTGGACACCACGCCGTATCTAACG GCACCAAAGTCAATACCCTGACCGAGCGGGGCGTTGAAGTTGTTAATGCCACCGAAACAGTAGAACGCACTAATATACCACGCATTTGCT CGAAAGGCAAAAAGACCGTTGACCTCGGCCAGTGCGGCCTGCTTGGCACAATCACAGGACCACCACAGTGTGACCAATTTCTCGAATTT TCCGCTGATCTGATAATAGAGAGGCGGGAAGGGAGCGACGTGTGTTATCCAGGGAAGTTCGTTAACGAGGAAGCACTTAGGCAAATACT GCGGGAAAGCGGTGGCATCGATAAAGAGGCCATGGGCTTTACCTACAGTGGTATTAGGACAAACGGCGCGACATCCGCGTGTAGGAGA AGCGGCTCATCCTTCTACGCTGAAATGAAGTGGCTGTTAAGTAATACCGATAATGCGGCTTTCCCGCAAATGACTAAGAGTTACAAAAACA CTAGAAAGTCTCCAGCTCTGATTGTTTGGGGGATTCATCATTCTGTGAGTACAGCCGAACAGACCAAGTTGTACGGCTCCGGAAACAAGC TGGTGACTGTAGGCTCTTCTAATTATCAACAGTCTTTTGTACCTAGTCCGGGTGCCCGCCCACAGGTAAACGGGCAGTCAGGCAGAATCG ATTTCCACTGGCTCATGCTGAATCCAAACGATACCGTCACCTTCAGTTTCAATGGAGCATTTATAGCACCGGACCGCGCGTCATTCCTAC GGGGTAAATCCATGGGCATTCAGAGCGGGGTTCAAGTAGATGCCAACTGTGAGGGCGACTGCTATCACTCTGGCGGTACGATCATAAGC AACCTTCCGTTTCAAAATATCGACAGCCGTGCTGTTGGCAAATGCCCCCGTTACGTGAAGCAGAGAAGCCTTCTGTTGGCTACTGGAATG AAGAATGTGCCCGAGATTCCTAAGGGAAGAGGCCTCTTCGGAGCGATCGCTGGATTTATAGAAAATGGCTGGGAAGGACTAATCGACGG CTGGTATGGGTTTAGACACCAGAACGCACAGGGGGAGGGCACTGCTGCGGATTATAAGTCGACACAGTCCGCGATCGACCAGATCACC GGTAAACTCAATCGGCTCATTGAGAAGACAAACCAACAGTTCGAGTTGATCGATAATGAATTTAATGAAGTGGAAAAGCAGATAGGAAATG TGATCAACTGGACACGGGACAGCATTACCGAAGTATGGAGTTACAATGCAGAGCTGCTGGTTGCTATGGAGAACCAGCACACGATTGATT TGGCCGATTCGGAAATGGACAAGCTTTACCAGCGGGTGAAAAGGCAACTCCGCGAAAATGCTGAAGAGGATGGGACTGCCTGTTTCGA GATTTTTCACAAGTGCGACGACGATTGCATGGCCAGCATAAGGAATAATACATATGACCACTCCAAATACAGGGAAGAAGCAATGCAGAA CAGAATCCAGATCGACCCTGTGAAGTTGAGTAGTGGATACAAGGATGTCATCCTGTGGTTTTCTTTTGGCGCCTCATGCTTCATTCTGCTG GCAATTGTAATGGGCCTGGTGTTCATTTGTGTGAAAAACGGAAATATGAGGTGTACAATCTGTATC

2737	ATGAACACGCAAATTTTGGTCTTCGCATTGATTGCAATTATTCCTACGAACGCAGACAAGATCTGTCTAGGACACCACGCTGTGTCAAACG GAACAAAAGTGAACACGCTGACCGAGCGGGGGGTCGAAGTGGTGAATGCAACTGAAACTGTAGAACGAACTAACATCCCAAGAATTTGT AGCAAAGGAAAGAAGACAGTTGACCTGGGTCAGTGCGGACTGCTCGGTACCATCACCGGTCCGCCGCAGTGCGACCAATTTTTAGAGTT CTCTGCAGACTTAATCATCGAGCGCAGAGAGGGCTCAGATGTCTGCTACCCCGGTAAGTTCGTGAACGAGGAGGCGCTTCGTCAGATTC TGCGAAAGAGTGGCGGGATCGACAAGGAGGCTATGGGCTTCACTTACTCAGGCATTCGTACGAATGGCGCCACATCCACCTGCAGGCG GAGCGGGTCTAGTTTTTACGCCGAGATGAAGTGGTTGTTATCGAACACCGACAATGCGGCCTTCCCCCAGATGACAAAATCATACAAAAA TACGCGCAAGTCACCCGCTATAATTGTGTGGGGCATTCACCACAGCGTCTCTACGGCCGAACAGACGAAATTATACGGGTCTGGGAACA AACTGGTGACAGTCGGCAGCAGTAACTACCAGCAATCTTTCGTGCCATCGCCAGGAGCCCGCCCGCAGGTCAATGGGTTATCTGGACGG ATAGACTTCCATTGGCTGATGCTTAACCCAAACGACACTGTTACCTTTAGCTTCAATGGGGCTTTTATCGCGCCTGATAGAGCTTGCTTCC TGAGAGGAAAGTCAATGGGAATCCAGTCAGGTGTGCAGGTGGACGCGGATTGTGAAGGTGATTGCTACCACAGTGGGGGGACAATCATT AGTAATCTGCCATTCCAGAATATTGATTCACGTGCAGTAGGAAAATGTCCACGATATGTTAAGCAAAGGTCGTTGCTACTCGCTACTGGTA TGAAAAACGTACCAGAAATACCTAAGGGAAGGGGACTATTTGGTGCAATTGCAGGGTTCATCGAGAACGGCTGGGAGGGCCTCATAGAC GGATGGTATGGTTTCCGGCACCAGAACGCACAGGGCGAAGGCACAGCAGCCGATTACAAGTCCACACAGAGCGCAATCGACCAGATCA CAGGCAAACTCAATAGCCTCATAGAAAAGACCAATCAGCAGTTTGAGCTCATTGACAACGAATTCAATGAAGTGGAGCGGCAGATAGGAA ACGTGATAAATTGGACCCGGGATAGCATTACGGAGGTCTGGTCCTATAACGCGGAGCTGCTGGTGGCCATGGAGAATCAGCACACCATT GACCTGGCCGATAGCGAAATGGATAAGCTGTATGAGCGCGTGAAGAGGCAGCTGCGGGAGAACGCTGAAGAGGACGGCACGGGCTGT TTTGAGATTTTCCACAAGTGTGATGACGACTGCATGGCCTCCATCCGAAACAACACCTATGACCATTCCAAATACCGGGAAGAGGCAATG CAAAATAGAATACAGATCGATCCTGTGAAGCTGTCCAGCGGCTATAAACACGTTATTCTGTGGTTCTCTTTCGGGGCATCCTGTTTTATAC TCTTAGCTATCGTTATGGGCCTGGTCTTTATTTGCGTGAAGAACGGCAACATGCGGTGCACGATTTGCATC
2738	ATGAACACCCAGATCCTGGTTTTCGCTTTAATCGCGATTATACCCACAAACGCTGATAAGATCTGCCTTGGGCATCATGCCGTGAGTAATG GTACAAAGGTCAATACACTCACTGAGAGAGGCGTCGAGGTCGTGAATGCTACTGAAACCGTCGAGAGAACAAATATACCCAGGATTTGCA GTAAAGGCAAGAAAACAGTAGATCTCGGTCAGTGCGGGTTGCTCGGGACCATCACTGGGCCACCTCAGTGCGACCAATTCTTGGAGTTT TCCGCCGACCTCATTATTGAGCGCCGGGAAGGCAGTGACGTCTGTTACCCCGGGAAATTCGTTAATGAGGACGCCCTGAGACAGATCCT CCGGGAGAGCGGGGGAATTGACAAGGAGGCCATGGGGTTCACCTATTCTGGAATTAGGACTAACGGCGCCACATCCGCGTGTCGCCGG TCTGGGTCATCTTTCTACGCCGAGATGAAATGGTTGCTCAGCAATACCGATAACGCCGCCTTTCCTCAGATGACCAAGTCATACAAAAAC ACACGAAAGTCACCCGCACTGATAGTGTGGGGCATACACCACAGCGTTTCAACTGCCGAACAGACAAAATTGTACGGATCTGGCAATAA GCTGGTCACCGTGGGCTCCAGTAACTACCAGCAAAGCTTCGTCCCCTCCCCCGGTGCTAGGCCACAAGTCAATGGCCAGTCCGGTAGAA TCGACTTCCATTGGCTAATGCTAAACCCCAATGACACTGTGACATTTTCTTTTAACGGAGCATTCATCGCCCCGGATAGGGCCAGCTTTCT CAGGGGCAAATCAATGGGGATTCAGAGTGGGGTGCAAGTTGATGCTAATTGTGAGGGTGATTGCTATCATTCAGGGGGAACCATTATCA GCAATTTGCCCTTCCAGAACATAGACTCACGCGCCGTCGGTAAGTGCCCTCGCTATGTCAAGCAAAGGTCCCTGCTGCTGGCCACTGGA ATGAAGAACGTGCCCGAGATCCCAAAGGGCAGGGGCTTATTCGGGGCCATTGCAGGTTTCATCGAAAACGGCTGGGAAGGACTAATAGA TGGGTGGTACGGCTTTCGTCACCAGAACGCACAAGGGGAGGGGACAGCTGCTGATTATAAAAGTACTCAGAGCGCTATCGACCAGATTA CAGGTAAGCTGAATCGGTTAATCGAGAAGACCAACCAGCAGTTCGAACTTATTGACAATGAGTTCAATGAGGTGGAGAAGCAGATTGGCA ATGTGATTAATTGGACGCGGGATAGTATTACCGAGGTCTGGTCCTATAACGCCGAGTTGCTAGTGGCAATGGAAAACCAGCACACGATAG ATCTTGCAGACAGCGAAATGGATAAGTTGTACGAAAGGGTGAAACGGCAACTAAGGGAAAACGCTGAGGAAGATGGGACAGGATGCTTC GAGATCTTCCACAAATGTGACGATGACTGTATGGCCAGTATTAGAAATAACACATATGACCATTCCAAATATCCGGAAGAGGCTATGCAAA ATCGCATCCAGATTGATCCCGTAAAACTCTCCTCGGGATACAAAGACGTGATCCTGTGGTTCTCCTTCGGAGCCTCTTGTTTCATCCTACT CGCCATCGTTATGGGGTTAGTGTTCATCTGTGTGAAGAATGGCAATATGAGGTGTACAATCTGCATT
2739	ATGAATACCCAGATCCTTGTTTTCGCCCTCATCGCCATAATTCCCACCAACGCAGATAAAATCTGTCTCGGCCACCACGCAGTGAGCAAT GGCACTAAGGTCAATACCCTCACTGAACGAGGTGTCGAGGTTGTCAACGCCACTGAAACGGTAGAGAGAACGAACATCCCCCGCATATG CAGTAAAGGCAAGAAAACTGTGGATCTCGCACAGTGCGGCCTGCTGGGCACCATTACCGGCCCGCCCCAGTGCGATCAATTCCTGGAAT TTTCTGCAGACCTCATCATCGAACGTCGGGAAGGGAGTGACGTGTGTTACCCCGGGAAATTCGTAAATGAGGAGGCTCTTCGCCAGATT CTGAGGGAAAGTGGGGGTATTGATAAGGAAGCAATGGGCTTTACCTATTCCGGAATTAGAACCAATGGCGCCACCTCCGCATGTAGACG GTCTGGTTCATCATTCTACGCTGAGATGAAGTGGCTGCTGTCCAACACTGACAACGCTGCATTTCCTCAGATGACTAAAAGTTACAAAAAC ACTCGCAAATCCCCTGCCCTTATTGTTTGGGGCATTCATCATAGTGTAAGTACAGCCGAACAAACCAAACTGTATGGCTCAGGGAATAAAT TAGTTACCGTGGGTTCATCCAACTACCAGCAGTCTTTCGTCCCTTCTCCAGGCGCTCGCCCACAAGTTAATGGGCAATCTGGGCGAATCG ATTTCCACTGGTTGATGCTGAACCCCAACGACACCGTTACCTTCAGTTTTAACGGCGCATTTATTGCCCCCGACCGCGCAAGTTTTCTGA GGGGCAAATCAATGGGGATCCAATCTGGCGTCCAGGTGGATGCCAACTGCGAAGGCGACTGTTATCATTCCGGTGGAACAATAATCTCT AATTTGCCGTTTCAGAACATTGATTCAAGAGCCGTGGGCAAGTGCCCACGGTACGTAAAACAGCGCTCCTTACTCCTTGCTACAGGGATG AAGAACGTGCCGGAGATCCCTAAAGGCCGAGGACTCTTTGGTGCTATTGCAGGGTTTATAGAAAATGGCTGGGAGGGACTGATTGATGG GTGGTACGGATTCCGACACCAGAATGCCCAAGGCGAGGGTACCGCTGCAGACTACAAAAGTACCCAGAGTGCTATCGACCAGATCACC GGGAAGTTAAATCGTTTGATAGAAAAAACAAATCAACAATTCGAGCTGATTGACAATGAGTTCAACGAGGTGGAGAAACAAATAGGAAAC GTAATCAATTGGACTCGTGACAGTATAACCGAAGTCTGGAGCTACAATGCGGAGCTACTCGTCGCTATGGAAAATCAGCATACAATCGAT CTGGCGGACAGCGAGATGGATAACCTGTACGAACGCGTGAAGAGACAACTGAGAGAAAACGCCGAGGAAGATGGCACAGGCTGCTTCG AGATTTTCCATAAGTGCGATGACGATTGCATGGCGAGTATTAGGAATAACACATATGATCACTCAAAGTATCGCGAGGAGGCGATGCAAA ACAGGATCCAGATTGACCCCGTGAAATTGTCTTCTGGCTACAAGGACGTGATACTCTGGTTTTCCTTCGGGGCCTCCTGTTTCATTCTGCT GGCGATTGTGATGGGGCTGGTGTTTATATGTGTGAAAAATGGGAATATGAGATGTACTATTTGCATT

2740	ATGAACACCCAGATCCTTGTGTTCGCACTGATAGCGATTATACCAACCAACGCCGACAAGATCTGCTTAGGGCACCACGCTGTAAGTAAT GGTACTAAGGTGAATACGCTGACTGAAAGAGGTGTGGAAGTGGTGAACGCCACAGAAACCGTAGAAAGGACCAACATCCCCAGAATCTG CAGCAAGGGGAAGCGGACCGTAGACCTGGGACAATGTGGACTTCTGGGAACAATTACTGGCCCTCCACAGTGCGACCAGTTCTTGGAG TTTAGCGCCGACCTCATTATCGAGCGACGGGAAGGCAGCGATGTATGTTATCCAGGCAAATTTGTGAACGAAGAAGCTCTCAGGCAGATT CTGCGGGAATCCGGCGGCATTGATAAAGAGGCTATGGGCTTTACCTACAGCGGGATCCGTACAAACGGGGCAACCTCTGCCTGTCGGA GATCTGGCTCAAGCTTTTATGCCGAGATGAAATGGCTCCTGTCTAATACTGACAATGCTGCTTTCCCACAGATGACGAAGTCCTATAAAAA CACCCGGAAGTCCCCCGCCTTAATTGTATGGGGCATCCATCATTCCGTTAGCACAGCCGAACAGACCAAGCTTTATGGATCTGGAAACAA GCTGGTGACCGTGGGCTCAAGTAACTATCAACAGTCATTCGTGCCTTCACCTGGAGCCCGCCCGCAGGTCAATGGCCTTAGCGGACGGA TTGACTTTCATTGGCTTATGCTGAATCCCAATGATACAGTGACGTTCAGCTTCAACGGCGCCTTCATAGCCCCCGACCGTGCCAGTTTCT TCGGGGAAAGAGCATGGGGATCCAAAGTGGCGTGCAGGTGGACGCCAATTGCGAAGGCGATTGCTATCATTCAGGCGGAACAATTATCA GTAACCTGCCGTTTCAGAATATTGACAGTCGAGCAGTTGGGAAGTGTCCAAGGTACGTGAAACAGCGCAGCCTCCTTTTAGCCACAGGC ATGAAGAATGTCCCAGAGATTCCGAAGGGCCGTGGGCTGTTTGGTGCTATTGCAGGTTTTATCGAAAATGGCTGGGAAGGACTTATTGAC GGCTGGTATGGGTTTAGGCATCAGAATGCCCAGGGTGAAGGCACTGCCGCCGACTATAAGAGCACGCAAAGCGCGATCGATCAGATTA CTGGCAAGTTAAATAGACTGATCGAAAAAACCAACCAACAGTTTGAGTTGATTGATAACGAGTTTAATGAGGTCGAAAAGCAGATTGGGAA CGTGATAAACTGGACCCGGGACAGCATAACCGAAGTATGGAGCTACAATGCAGAGCTGCTTGTGGCTATGGAGAATCAGCACACAATCG ATCTGGCAGACAGTGAGATGGACAAACTCTACGAGAGGGTCAAGCGGCAGCTCCGGGAAAATGCCGAGGAAGATGGTACCGGATGCTT TGAGATTTTCCATAAGTGTGACGATGATTGTATGGCCTCCATTCGCAATAACACCTATGATCACTCAAAGTATAGGGAGGAGGCAATGCAA AATCGCATCCAGATCGACCCCGTGAAGCTTAGCTCCGGATATAAGGACGTGATCTTGTGGTTCTCTTTTGGTGCCTCCTGCTTCATCCTTC TGGCCATTGTGATGGGATTGGTATTTATTTGCGTGAAGAATGGCAATATGCGATGCACAATTTGCATC
2741	ATGAATACACAGATACTGGTGTTTGCGCTGATTGCCATTATTCCAACAAATGCTGATAAAATCTGCCTGGGGCATCACGCAGTTTCAAATG GCACCAAAGTGAACACACTTACCGAGCGAGGCGTGGAGGTGGTGAATGCCACTGAAACAGTGGAGCGTACCAATATCCCTCGTATCTGT AGCAAGGGTAAACGGACTGTCGATCTGGGCCAGTGTGGTTTACTGGGGACAATTACAGGCCCCCCTCAGTGTGATCAGTTCTTAGAGTT TTCAGCTGATCTGATCATAGAACGGACAGAAGGCAGCGACGTGTGTTACCCTGGCAAGTTCGTGAATGAGGAGGCCCTGCGCCAAATCC TGAGAGAGTCTGGGGGGATAGATAAAGAAGCGATGGGGTTCACTTATTCTGGCATCCGAACCAATGGCGCTACGAGCGCATGCCGCAG ATCTGGCTCTAGCTTTTACGCTGAGATGAAATGGCTCCTCTCTAATACGGACAATGCTGCATTTCCCCAGATGACCAAATCTTACAAGAAT ACCCGTAAGTCCCCAGCTCTCATCGTGTGGGGAATCCACCACTCTGTGAGTACTGCCGAGCAGACAAAGCTCTACGGCTCTGGAAATAA GTTGGTGACCGTAGGTTCATCCAATTATCAACAGTCATTCGTGCCTAGTCCCGGTGCTCGGCCCCAAGTGAACGGCCTCTCCGGGAGGA TTGACTTCCATTGGCTGATGCTGAACCCTAACGATACAGTGACATTTTCATTTAACGGCGCGTTTATAGCTCCTGACCGAGCTTCCTTTCT CCGAGGTAAGAGCATGGGCATCCAGAGTGGGGTACAAGTCGATGCGAATTGCGAGGGCGATTGTTACCACAGTGGTGGCACTATCATTT CCAACCTTCCTTTTCAGAACATTGACTCACGGGCGGTCGGCAAATGTCCTCGCTACGTGAAGCAAAGGTCCCTTCTGCTGGCCACAGGG ATGAAAAACGTGCCTGAGATCCCCAAGGGGCGGGGTCTCTTTGGGGCCATCGCAGGATTTATAGAAAATGGGTGGGAGGGACTCATTAA TGGCTGGTACGGATTTAGACACCAGAACGCCCAAGGGGAGGGCACTGCCGCCGATTATAAATCCACTCAGAGCGCTATCGATCAAATAA CAGGTAAACTGAATCGGTTAATTGAGAAAACCAATCAACAGTTTGAACTGATAGATAACGAGTTTAATGAAGTGGAGAAGCAGATTGGAAA CGTGATCAACTGGACCAGAGACTCCATTACTGAGGTTTGGTCGTATAACGCGGAGCTGCTCGTCGCTATGGAGAATCAACATACCATCGA CCTGGCTGATAGTGAGATGGACAAGCTGTACGAGCGCGTTAAAAGACAGCTGCGTGAGAACGCTGAAGAGGATGGTACAGGATGTTTTG AGATCTTTCACAAGTGTGATGACGACTGCATGGCTAGTATCCGAAACAATACATACCACCACAGCAAATATAGAGAAGAGGCCATGCAAA ATCGTATACAAATTGATCCCGTGAAACTATCCTCCGGCTACAAGGATGTGATCTTGTGGTTTTCCTTTGGCGCATCTTGCTTCATCCTGCT GGCAATTGTAATGGGATTGGTCTTTATCTGCGTGAAGAACGGCAATATGCGGTGTACTATCTGCATC
2742	ATGAACACTCAGATCCTCGTGTTTGCCTTGATCGCCATCATACCAACGAATGCAGATAAGATCTGCCTGGGACACCATGCTGTATCAAAT GGCACTAAGGTGAACACCCTGACTGAGCGGGGCGTTGAAGTGGTGAACGCCACTGAAACGGTGGAAAGGACCAATATCCCCAGAATTT GCTCAAAGGGCAAAAAAACTGTTGACCTCGGCCAGTGCGGCCTCTTAGGGACGATAACCGCTCCCCCTCAGTGTGATCAGTTCCTGGAG TTCTCAGCCGATCTGATCATTGAAAGAAGAGAAGGCTCGGACGTGTGTTATCCGGGCAAGTTCGTGAACGAAGAAGCCCTGCGCCAGAT CCTCCGCGAAAGCGGGGGCATCGACAAGGAAGCCATGGGTTTCACCTACTCTGGAATTCGCACCAACGGGGCCACTTCCGCATGCCGA CGGTCCGGTTCGAGCTTTTACGCTGAGATGAAGTGGCTTTTATCAAACACAGATAACGCGGCTTTCCCGCAGATGACCAAGTCCTATAAG AACACAAGAAAGTCTCCGGCCCTGATTGTCTGGGGGATCCACCATAGTGTTTCTACTGCGGAGCAGACAAAACTTTATGGTAGCGGGAA CAAACTGGTCACCGTCGGCTCCTCCAATTACCAGCAGAGTTTTGTTCCCTCCCCCGGGGCACGACCCCAAGTCAATGGGCAAAGTGGCA GGATAGATTTCCACTGGTTGATGCTCAACCCCAATGACACCGTGACATTTAGCTTTAACGGAGCTTTTATCGCACCAGACCGGGCATCAT TCCTGCGTGGCAAGAGTATGGGTATTCAAAGTGGGGTTCAGGTAGACGCAAACTGCGAGGGCGACTGTTATCACAGTGGGGGGACCATT ATCAGCAACCTACCGTTTCAGAATATCGATAGTAGGGCAGTGGGGAAATGCCCTCGATATGTGAAGCAACGGTCCCTGCTCCTGGCTAC GGGAATGAAAAACGTCCCAGAGATTCCCAAGGGCCGGGGCCTTTTCGGCGCGATTGCTGGCTTTATTGAGAATGGGTGGGAGGGATTAA TCGACGGGTGGTATGGCTTTAGGCACCAGAACGCTCAGGGCGAAGGTACGGCCGCAGATTACAAGTCCACCCAGTCCGCCATCGACCA GATTACAGGCAAGCTCAACCGTTTAATTGAGAAAACCAATCAGCAGTTCGAGCTCATCGATAACGAGTTCAATGAGGTGGAGAAACAGAT TGGCAACGTGATAAACTGGACTAGGGACTCAATCACAGAGGTCTGGAGCTATAATGCCGAGCTGCTAGTTGCCATGGAAAACCAGCATA CAATCGATCTTGCCGACTCCGAAATGGACAAACTATATGAGAGAGTCAAACGTCAGCTAACAGAGAATGCAGAAGAGCATGGCACGGGT TGCTTCGAAATTTTCCACAAGTGTGACGACGACTGCATGGCCAGCATTAGAAACAACACATACGATCATAGCAAGTATAGGGAGGAGGCG ATGCAGAACAGGATTCAGATTGATCCCGTCAAGCTCTCTTCTGGCTATAAGGATGTCATTCTGTGGTTTTCTTTCGGCGCCTCATGTTTCA TCCTGCTTGCAATCGTGATGGGATTAGTTTTCATTTGTGTAAAGAACGGCAATATGCGGTGTACAATCTGCATC

2743	ATGAATACTCAAATCCTGGTCTTCGCCCTTATTGCGATCATACCTACCAACGCCGATAAGATCTGCTTGGGGCACCATGCCGTCTCAAAC GGCACCAAGGTCAACACTCTTACAGAACGGGGTGTAGAAGTGGTTAATGCCACTGAGACTGTCGAACGAACAAACATCCCTCGAATCTG CAGCAAAGGGAAACGGACTGTCGATCTTGGCCAGTGTGGGCTCCTGGGTACCATAACGGGGCCTCCCCAGTGCGACCAATTCTTAGAGT TCTCGGCCGATTTAATAATAGAACGCAGAGAGGGATCAGACGTATGTTACCCGGGCAAATTCGTCAATGAGGAGGCACTCCGGCAGATC CTGAGGGACAGCGGCCGGAATTGACAAAGAGGCCATGGGATTCACTTATAGTGGGATTCGTACCAATGGCGCTACAAGTGCCTGCCGTAG ATCCGGGTCGAGTTTTTATGCCGAAATGAAGTGGCTACTGTCCAATACAGATAACGCTGCGTTCCCTCAGATGACTAAGTCTTACAAGAAT ACCAGGAAAAGCCCTGCCCTTATTGTGTGGGGAATTCACCATTCGGTGAGCACCGCGGAACAAACAAAACTGTACGGCAGTGGGAACAA GCTCGTGACGGTAGGGTCCTCTAACTACCAACAAAGTTTCGTCCCATCCCCAGGAGCTAGACCTCAGGTGAACGGGTTAAGCGGGCGAA TTGATTTCCATTGGCTGATGCTGAACCCCAACGACACAGTGACATTTAGCTTCAATGGCGCCTTCATAGCTCCAGACAGAGCTTCCTTTCT GAGAGGCAAAAGTATGGGAATTCAGTCTGGAGTCCAAGTGGATGCCAACTGCGAGGGGGACTGTTATCATTCTGGCGGGACCATTATCT CTAACCTACCATTCCAGAATATAGACAGTAGGGCCGTAGGGAAGTGTCCCCGCTATGTAAAGCAGCGGAGTCTGCTGCTGGCAACAGGT ATGAAGAACGTGCCCGAAATCCCCAAGGGCCGCGGACTCTTTGGGGCTATCGCTGGGTTCATTGAGAACGGATGGGAAGGCCTCATCG ATGGGTGGTATGGTTTTCGTCACCAGAACGCTCAGGGGGAAGGCACCGCCGCTGACTATAAATCTACTCAATCGGCCATCGACCAGATT ACAGGGAAGCTGAATCGTCTAATCGAGAAAACCAATCAGCAATTCGAGCTTATTGACAATGAGTTCAATGAAGTGGAGAAACAGATTGGA AATGTTATAAACTGGACCCGGGACAGCATAACTGAGGTGTGGAGTTACAACGCGGAGCTCCTCGTTGCCATGGAGAACCAGCACACCAT AGATCTGGCCGATAGCGAAATGGACAAGCTGTACGAGCGGGTTAAGCGTCAACTGCGAGAGAATGCCGAAGAAGATGGCACCGGATGC TTTGAAATATTCCACAAATGCGATGATGACTGTATGGCTTCTATACGTAATAACACTTACGATCATAGCAAGTACCGAGAGGAGGCTATGC AAAACAGGATACAGATCGACCCCGTGAAACTTTCATCAGGCTATAAGGACGTTATACTGTGGTTCTCGTTCGGGGCAAGTTGCTTCATTCT GCTCGCCATTGTCATGGGCCTGGTTTTTATCTGTGTGAAGAACGGCAATATGCGGTGTACCATTTGCATC
2744	ATGAATACCCAGATCTTGGTTTTCGCACTGATCGCAATTATCCCAACAAACGCAGACAAAATTTGTCTTGGTCATCATGCCGTTTCCAACG GGACTAAAGTTAACACCCTCACGGAACGCGGAGTAGAGGTGGTCAACGCTACAGAAACCGTGGAGCGTACGAACATCCCTCGCATATGT TCCAAAGGAAAGAAGACTGTCGATCTGGGACAGTGCGGCCTATTAGGGACCATCACCGGCCCTCCTCAGTGCGATCAATTCCTTGAGTT TTCCGCGGATCTCATAATAGAGCGCAGGGAGGGTTCCGACGTCTGTTACCCTGGCAAGTTTGTGAATGAGGAAGCCTTACGGCAAATCT TGAGGAAGTCTGGCGGTATTGACAAGGAAGCGATGGGTTTTACTTATTCCGGCATCAGGACAAATGGGGCTACTTCTACTTGTCGGAGG AGTGGAAGCAGCTTCTATGCCGAGATGAAGTGGCTTCTGAGCAACACTGACAATGCCGCCTTTCCCCAGATGACAAAATCATACAAAAAC ACACGCAAATCGCCCGCAATCATTGTCTGGGGCATTCATCATTCTGTCTCAACCGCCGAGCAGACTAAACTCTATGGCTCAGGAAATAAA TTAGTGACCGTGGGATCCAGCAATTACCAGCAAAGTTTCGTGCCTTCCCCAGGCGCAAGACCGCAGGTTAACGGCCTCTCTGGACGCAT CGACTTTCATTGGCTCATGCTTAACCCCAATGATACAGTGACATTCTCCTTCAATGGAGCTTTTATCGCTCCGGACCGTGCTTGTTTCCTT CGAGGCAAGTCCATGGGAATTCAAAGTGGCGTTCAGGTTGACGCCGACTGCGAAGGTGACTGCTATCATAGCGGAGGCACTATCATCAG CAATCTTCCATTCCAGAACATTGACTCGCGAGCCGTGGGAAAGTGCCCAAGGTATGTCAAGCAGAGGAGCCTGCTGCTGGCTACCGCA TGAAAAACGTCCCCGAAATTCCAAAAGGCCGTGGTTTGTTTGGCGCCATTGCCGGTTTTATCGAGAACGGATGGGAGGGACTGATTGAC GGCTGGTACGGCTTCAGACACCAGAACGCGCAGGGAGAAGGGACGGCGGCCGATTATAAGAGCACTCAGAGCGCTATCGATCAGATTA CCGGTAAATTGAATAGACTGATCGAAAAGACCAACCAGCAATTTGAACTCATTGACAATGAGTTCAACGAGGTGGAACGGCAGATAGGAA ATGTCATTAACTGGACACGAGATTCTATTACTGAGGTGTGGAGTTACAACGCGGAGCTCCTAGTCGCCATGGAGAACCAGCATACAATCG ACCTGGCCGACAGCGAAATGGACAAGCTGTATGAAAGAGTAAAGAGGCAGCTGCGGGAGAACGCCGAGGAGGATGGAACCGGTTGTTT TGAGATCTTTCACAAATGCGACGATGACTGCATGGCAACCATCCGGAACAACACTTATGACCACAGCAAGTATAGGGAGGAGGCCATGC AGAATAGAATCCAGATCGACCCAGTCAAGTTAAGCAGTGGCTACAAAGATGTTATCCTCTGGTTCTCTTTTGGAGCCTCATGTTTCATTCT TCTAGCAATCGTGATGGGTCTCGTCTTTATTTGTGTCAAAAATGGAAATATGCGCTGTACCATTTGTATT
2745	ATGAACACTCAGATCCTTGTCTTTGCGCTAATTGCAATTATTCCAACGAATGCAGACAAAATCTGCCTGGGCCATCACGCTGTTTCCAACG GAACAAAGGTGAACACACTCACGGAAAGGGGCGTCGAAGTGGTGAATGCCACGGAGACTGTTGAGAGGACCAACATCCCCAGGATCTG TAGCAAAGGAAAGCGGACAGTTGACCTGGGCCAGTGTGGCCTTCTTGGTACCATCACCGGACCTCCCCAATGCGACCAGTTTCTGCAGT TTAGCGCAGACCTTATCATCGAGCGGCGCGAGGGGAGCGACGTCTGTTACCCGGGCAAGTTCGTAAATGAAGAGGCGCTAAGGCAGAT CCTTAGGGAATCCGGGGGTATAGACAAAGAGGCAATGGGCTTTACCTACAGCGGAATCCGGACTAATGGCGCCACCTCCGCTTGTAGGC GTTCAGGATCTAGTTTCTACGCTGAGATGAAGTGGTTGTTGAGTAATACCGATAATGCGGCCTTCCCTCAGATGACCAAATCCTATAAAAA TACAAGAAAAAGCCCAGCCCTGATCGTATGGGGGATCCACCATTCTGTATCTACCGCAGAACAGACTAAGCTCTACGGGTCTGGGTCCA AATTGGTAACTGTAGGGAGCTCTAATTATCAACAGAGCTTTGTTCCGAGTCCTGGCGCCAGGCCCCAGGTGAACGGTCTGTCCGGGAGA ATTGATTTTCACTGGCTCATGCTCAATCCTAATGATACGGTTACCTTCAGCTTTAATGGAGCGTTCATCGCACCCGATAGGGCTTCATTTTT ACGTGGCAAGTCCATGGGAATCCAGTCAGGAGTCCAGGTCGACGCTAATTGCGAGGGTGATTGCTACCACAGCGGAGGAACTATCATTT CGAACCTCCCCTTTCAGAATATTGATTCTAGGGCCGTCGGTAAATGTCCAAGATATGTGAAGCAAAGGTCTTTGCTACTCGCCACTGGAA TGAAAAACGTGCCTGAAATCCCTAAAGGCCGGGGCCTGTTTGGCGCAATCGCTGGATTTATCGAGAATGGTTGGGAGGGACTGATTGAC GGCTGGTATGGATTCCGGCACCAAAACGCTCAGGGGGAAGGCACTGCTGCCGATTACAAGTCCACTCAATCCGCGATTGACCAAATAAC CGGTAAGCTGAATAGGCTGATCGAGAAGACAAATCAGCAGTTTGAGCTCATCGATAACGAGTTTAATGAAGTAGAAAAGCAGATTGGGAA CGTTATTAATTGGACAAGAGATAGTATAACCGAGGTGTGGAGCTATAATGCGGAGCTCCTGGTGGCCATGGAAAATCAGCATACAATAGA CCTCGCCGATAGCGAAATGGATAAATTATACGAGAGAGTGAAGCGGCAGCTGAGGGAAAACGCTGAGGAGGACGGAACTGGCTGTTTT GAGATCTTTCATAAGTGCGATGATGACTGTATGGCCAGCATTCGGAACAACACCTACGATCATTCTAAGTATAGGGAGGAGGCAATGCAG AACCGAATCCAAATAGATCCTGTTAAGCTGTCTAGCGGATATAAAGACGTGATACTTTGGTTCAGCTTCGGGGCCAGCTGTTTCATTCTGC TTGCTATCGTCATGGGACTTGTATTTATCTGCGTGAAGTCCAGAAACATGAGGTGCACTATATGTATC

2746	ATGAACACTCAAATCCTGGTGTTCGCGCTGATCGCCATCATACCTACCAATGCTGACAAAATCTGCTTGGGCCACCACGCCGTTAGCAAT GGAACCAAAGTGAACACGTTGACGGAAAGAGGAGTGGAGGTGGTGAATGCCACGGAAACAGTGGAGCGGACAAACATCCCCCGGATCT GTAGTAAAGGGAAAAAGACGGTTGATCTCGGCCAGTGCGGCCTCCTGGGGACAATAACCGGGCCTCCTCAATGTGACCAGTTCCTGGAA TTCTCAGCCGATTTAATCATAGAGCGGCGGGAGGGGAGTGATGTGTGTTATCCAGGGAAATTCGTGAATGAAGAAGCCTTGAGACAGAT CCTTCGCGAAAGTGGAGGAATCGACAAGGAGGCCATGGCGTTTACTTATAGTGGAATCAGAACCAACGGCGCCACGTCCGCATGCCGA CGCAGCGGTTCTAGCTTCTATGCCGAAATGAAGTGGCTGCTCTCCAATACCGATAATGCAGCCTTCCCCCAAATGACGAAATCTTACAAG AATACCCGCAAAAGTCCTGCTTTGATTGTCTGGGGCATCCACCACAGCGTCTCTACTGCCGAACAAACAAAGTTATACGGATCAGGCAAC AAACTGGTCACCGTAGGTTCCAGCAACTACCAACAGTCATTTGTCCCGAGCCCTGGAGCCCGCCCACAGGTGAACGGGCAGTCCGGCC GAATTGACTTTCACTGGCTGATGCTCAACCCGAACGATACTGTGACCTTCAGCTTCAATGGCGCGTTCATTGCACCTGACCGTGCATCAT TCTTGAGGGGCAAAAGCATGGGGATCCAGTCTGGAGTCCAGGTCGACGCAAATTGCGAAGGAGATTGTTATCACAGCGGGGGCACGAT TATCTCAAATCTCCCCTTTCAAAATATAGATAGCCGAGCAGTTGGCAAATGCCCTCGCTACGTGAAACAAAGGAGCCTGCTCCTGGCTAC CGGGATGAAGAACGTACCTGAAATCCCCAAGGGACGCGGACTTTTCGGCGCTATTGCTGGGTTCATAGAAAACGGCTGGGAAGGGTTGA TCGACGGCTGGTATGGCTTCAGACACCAGAATGCTCAGGGAGAAGGCACAGCAGCCGACTACAAATCAACCCAGTCTGCCATCGATCAG ATTACAGGAAAGCTGAATCGGCTCATCGAGAAAACTAACCAGCAGTTCGAACTCATCGACAATGAGTTTAACGAGGTCGAAAAACAGATT GGTAATGTGATCAACTGGACTAGGGACTCCATTACAGAGGTTTGGTCCTATAATGCCGAGCTTTTGGTTGCCATGGAAAACCAACATACC ATCGACCTGGCCGATAGCGAGATGGACAAGTTGTACGAGAGGGTAAAAAGGCAGCTCAGAGAAAACGCCGAGGAAGATGGAACCGGTT GTTTCGAAATTTTTCATAAGTGCGATGACGACTGTATGGCTTCAATTCGAAACAATACTTACGATCACAGCAAGTACCGCGAAGAGGCCAT GCAGAATAGGATTCAGATTGATCCTGTTAAGTTGTCATCCGGCTATAAGGATGTAATCTTATGGTTTAGTTTCGGCGCTAGCTGCTTCATA CTGCTGGCAATCGTGATGGGGCTGGTATTTATATGCGTCAAAAATGGGAATATGCGCTGTACCATCTGCATC
2747	ATGAATACTCAAATCTTGGTCTTTGCACTGATCGCCATTATTCCTACTAACGCCGATAAAATTTGCTTGGGTCATCACGCCGTGTCAAATG GCACTAAAGTTAATACCTTAACAGAGCGCGGAGTAGAGGTGGTGAATGCAACGGAGACTGTTGAGCGGACCAATATACCCCGCATTTGT AGTAAAGGCAAGAAAACAGTGGACCTGGGCCAATGCGGGTTGCTTGGGACGATCACCGGCCCCCCGCAGTGCGATCAGTTCTTAGAATT CTCCGCTGACCTAATCATCGAAAGACGGGAGGCAGCGATGTGTGTTATCCTGGCAAATTCGTCAACGAGGAAGCCCTCAGGCAGATTC TTCGCAAGTCAGGGGGTATTGATAAGGAGGCGATGGGATTCACATACTCCGGAATTCGAACAAACGGCGCTACAAGTACTTGTAGGAGG AGCGGGAGTTCATTCTACGCAGAGATGAAGTGGCTGCTCAGCAACACCGACAACGCAGCGTTCCCACAGATGACCAAAAGCTATAAAAA TACCCGAAAGAGTCCCGCAATCATCGTATGGGGAATTCATCACAGTGTCTCTACCGCTGAGCAGACCAAGTTGTATGGCTCCGGCAACAA GCTCGTTACCGTTGGCAGCTCAAATTACCAACAGTCCTTCGTCCCCTCCCCAGGGGCAAGGCCCCAAGTCAACGGCCTGTCCGGCAGAA TTGACTTCCACTGGCTCATGCTAAACCCCAATGATACGGTCACCTTCAGTTTTAATGGGGCATTCATCGCACCCGACAGAGCCTGTTTTCT GCGAGGTAAATCAATGGGTATACAATCCGGGGTCCAGGTCGACGCTGACTGCGAGGGCGACTGCTATCATTCGGGAGGCACCATTATTT CCAACCTGCCTTTCCAGAACATTGATTCCAGAGCCGTGGGCAAGTGTCCTCGGTACGTGAAACAACGATCTCTGCTTCTGGCTACCGGAA TGAAGAACGTGCCCGAGATCCCCAAGGGACGAGGCCTGTTCGGGGCCATTGCCGGATTTATTGAGAATGGCTGGGAGGGTCTCATCGA TGGATGGTACGGCTTCAGGCACCAGAACGCACAAGGTGAAGGCACAGCTGCTGATTACAAATCCACCCAGAGTGCAATTGACCAGATCA CCGGAAAACTCAATAGACTGATTGAAAAAACCAACCAACAGTTTGAATTGATTGATAATGAGTTTAACGAAGTCGAACGGCAGATTGGTAA TGTGATCAATTGGACCCGGGATTCCATTACAGAGGTTTGGTCCTACAACGCCGAGCTGCTCGTCGCAATGGAAAATCAGCACACCATAGA TCTGGCCGACTCCGAGATGGACAAGCTTTACGAAAGGGTGAAGAGACAGCTCAGGGAGAACGCCGAAGAGGACGGAACAGGGTGTTTC GAAATCTTCCATAAGTGCGACGATGATTGTATGGCTTCTATCCGGAACAACACCTATGATCACTCCAAATATAGACAGGAAGCAATGCAGA ACAGGATTCAAATCGACCCTGTCAAACTCTCAAGTGGGTACAAGGACGTAATCCTATGGTTCAGCTTTGGCGCCAGCTGTTTTATTCTGCT GGCAATCGTGATGGGTTTGGTGTTTATCTGCGTCAAGAATGGCAACATGAGGTGTACCATTTGCATC
2748	ATGAATACACAGATCCTGGTCTTCGCCCTTATCGCTATCATCCCCACAAACGCCGACAAGATATGCCTGGGCCACCACGCCGTTAGCAAC GGCACCAAAGTAAACACTCTCACTGAAAGGGGTGTGGAAGTTGTGAACGCAACAGAAACCGTAGAACGCACGAACATTCCAAGAATTTG CTCAAAGGGGAAACGTACAGTCGATCTAGGACAGTGCGGCCTGCTCGGCACCATAACCGGGCCCCCCCAATGCGATCAATTTCTGGAGT TCTCTGCCGATCTTATCATCGAGCGTCGGGAAGGGAGCGATGTGTGCTACCCCGGCAAATTTGTGAACGAGGAGGCCCTCAGACAGATC TTAAGAGAGTCTGGAGGGATTGACAAAGAGGCCATGGGGTTCACATACTCCGGCATACGTACGAATGGGGCCACGAGTGCTTGTCGGA GGAGCGGTTCAAGTTTCTACGCGGAGATGAAATGGCTGCTGAGCAATACCGATAACGCCGCCTTCCCCCAGATGACCAAAAGTTATAAAA ATACAAGGAAATCACCCGCCCTTATCGTTTGGGGAATTCATCACTCCGTCAGTACCGCCGAACAGACTAAGCTCTATGGATCGGGTTCAA AGCTAGTGACCGTGGGCTCAAGCAATTACCAGCAATCCTTTGTACCAAGCCCTGGGGCCCGTCCCCAAGTGAACGGATTAAGCGGTAGG ATTGACTTCCATTGGTTGATGCTCAATCCGAATGATACGGTGACCTTCAGTTTCAACGGGGCGTTCATAGCTCCTGATCGCGCGAGCTTC CTCCGGGGAAAATCGATGGGAATTCAGTCAGGCGTTCAGGTGGACGCCAATTGTGAAGGAGACTGCTACCATAGCGGAGGGACTATAAT TTCCAATCTGCCTTTTCAGAACATTGACTCCCGCGCTGTGGGGAAGTGTCCCCGGTACGTGAAGCAACGCTCCCTGCTTCTCGCCACAG GCATGAAGAACGTACCTGAGATTCCTAAGGGCCGGGGGCTTTTCGGCGCGATTGCTGGATTCATCGAAAACGGCTGGGAAGGTTTAATA GATGGCTGGTATGGCTTCAGGCACCAGAATGCACAGGGCGAGGGGACCGCGGCCGACTATAAGTCTACCCAAAGTGCTATTGATCAGAT AACCGGAAAGCTTAACCGCCTGATCGAGAAAACCAATCAACAGTTTGAACTTATCGACAACGAATTCAACGAGGTTGAGAAGCAGATTGG GAACGTCATTAATTGGACCCGAGATTCGATCACAGAGGTGTGGAGCTACAACGCCGAGCTGCTGGTCGCGATGGAAAATCAGCATACAA TCGACTTGGCCGATAGTGAGATGGACAAACTATACGAGAGAGTGAAACGGCAGCTGAGGGAGAACGCCGAAGAGGACGGCACCGGGTG CTTTGAGATTTTTCACAAGTGCGATGACGATTGTATGGCTTCCATCAGGAATAATACTTACGACCACAGTAAGTATCGCGAAGAGGCTATG CAAAACCGCATTCAGATCGACCCCGTCAAACTCTCATCGGGCTATAAAGATGTGATACTCTGGTTTTCTTTTGGTGCATCATGCTTCATTC TTTTGGCAATAGTGATGGGGCTGGTGTTCATCTGCGTAAAGAACGGTAACATGAGATGCACTATCTGTATT

2749	ATGAATACACAGATCTTAGTGTTTGCTTTAATAGCTATTATTCCAACCAATGCCGATAAAATCTGTCTGGGACACCACGCCGTGAGCAATG GGACGAAAGTGAACACTCTTACCGAACGAGGAGTCGAGGTTGTCAATGCTACGGAAACTGTGGAAAGGACCAACATCCCAAGGATCTGC AGCAAAGGCAAAAAAACTGTAGATCTGGGGCAGTGTGGACTGCTGGGGACCATTACCGGACCCCCCCAGTGCGACCAGTTCTTGGAGTT TTCCGCCGATCTGATTATTGAGCGGCGCGAGGGGTCTGATGTGTGCTATCCTGGTAAATTTGTCAACGAAGAGGCGCTGCGACAAATTCT CCGCAAAAGTGGGGGAATCGACAAGGAAGCCATGGGCTTCACATACTCTGGCATTCGAACAAATGGGGCCACTTCCACATGCAGGCGGT CTGGGTCTTCATTTTATGCCGAGATGAAGTGGCTTCTTTCTAATACTGACAACGCCGCCTTCCCCCAAATGACCAAGTCCTACAAAAACAC CCGGAAGTCCCCAGCCATTATCGTCTGGGGAATCCATCACAGCGTTTCTACAGCCGAACAGACCAAGCTCTACGGATCAGGAAACAAAC TGGTTACAGTGGGTAGCAGCAACTATCAGCAGTCGTTCGTTCCTTCTCCTGGGGCGCGGCCGCAAGTAAACGGGCTCAGTGGGCGTATC GATTTCCACTGGCTGATGCTGAACCCAAACGACACAGTCACCTTCAGCTTTAATGGAGCCTTCATCGCCCCTGATAGGGCCTGCTTTCTG CGGGGAAAATCCATGGGCATCCAAAGCGGAGTGCAGGTCGACGCCGATTGCGAAGGGGACTGCTACCATTCTGGTGGGACCATCATCT CCAACCTCCCATTTCAGAACATTGACTCCCGGGCGGTCGGAAAATGCCCACGTTACGTGAAGCAGCGCAGTCTCTTACTCGCCACTGGG ATGAAGAACGTGCCCGAAATTCCAAAGGGGCGTGGTCTGTTTGGGGCGATTGCCGGCTTCATTGAGAACGGCTGGGAGGGGCTTATCG ACGGCTGGTACGGATTCCGTCACCAGAATGCCCAGGGTGAGGGCACCGCAGCAGACTATAAAAGCACACAGAGCGCCATTGACCAAAT CACTGGGAAGCTGAATCGCCTGATTGAGAAAACTAACCAGCAGTTTGAACTCATAGATAATGAATTTAACGAAGTCGAAAGGCAGATTGG CAACGTAATTAATTGGACTCGCGACTCAATCACTGAGGTCTGGAGTTACAATGCAGAACTGCTTGTCGCTATGGAGAACCAGCATACCAT TGACCTGGCTGATAGCGAGATGGATAAACTGTACGAAAGGGTGAAGCGACAGCTCCGCGAGAACGCAGAGGAGGACGGGACCGGCTGT TTCGAGATCTTTCACAAGTGTGACGACGACTGTATGGCTTCGATTCGGAACAACACCTATGACCACTCCAAATATAGGGAAGAGGCTATG CAGAATAGGATCCAAATCGATCCTGTCAAGCTGAGCTCCGGCTACAAAGACGTAATCCTGTGGTTCTCTTTTGGCGCCTCGTGTTTTATC CTACTCGCCATCGTCATGGGTTTAGTTTTTATCTGCGTAAAAAATGGAAATATGAGATGTACTATCTGCATC
2750	ATGAACACGCAGATCCTTGTCTTCGCCCTGATTGCCATTATACCCACTAACGCCGATAAAATTTGCTTGGGCCACCACGCCGTGTCCAAT GGAACAAAAGTGAATACTTTGACCGAGAGGGGTGTCGAAGTCGTCAACGCCACCGAGACTGTCGAACGGACAAATATACCGCGTATCTG TTCTAAAGGGAAGCGAACCGTGGACCTGGGTCAATGCGGACTGCTGGGGACTATTACAGGGCCCCCTCAGTGCGATCAGTTTCTGGAAT TTAGCGCTGACCTCATTATTGAAAGGAGGGAGGGATCAGATGTGTGTTACCCAGGTAAGTTTGTCAATGAGGAAGCCCTCCGACAGATTC TCAGGGAATCTGGAGGCATTGATAAAGAGGCTATGGGGTTCACCTATTCAGGCATCAGGACCAATGGGGCTACCAGCGCTTGCCGCCG GAGCGGATCATCTTTTTATGCCGAGATGAAATGGCTACTGTCAAACACCGACAACGCCGCATTCCCACAGATGACCAAGAGTTACAAGAA TACACGCAAGTCACCCGCCCTCATCGTCTGGGGCATCCATCATTCCGTGAGCACAGCCGAGCAGACGAAGCTCTACGGATCGGGCAGC AAGCTCGTAACCGTCGGGAGCTCTAACTATCAGCAATCTTTTGTGCCCTCCCCCGGAGCCCGGCCACAGGTCAATGGACTATCCGGGCG CATCGATTTCCACTGGCTGATGTTGAACCCGAACGACACCGTGACATTTTCATTCAATGGGGCATTCATCGCGCCAGACCGCGCGTCCTT TCTGAGGGGAAAGTCCATGGGTATCCAGTCTGGTGTACAGGTGGATGCCAACTGTGAGGGTGACTGCTACCATAGTGGAGGAACAATTA TCTCCAACTTGCCATTCCAAAACATCGACAGTAGAGCTGTTGGTAAATGTCCCAGGTACGTGAAGCAAAGGTCATTGCTGCTGGCCACTG GGATGAAGAATGTGCCCGAAATTCCAAAAGGAAGGGGGCTTTTCGGGGCAATCGCCGGCTTCATTGAGAATGGCTGGGAAGGCCTCATC GACGGGTGGTACGGGTTTAGGCATCAAAACGCGCAGGGGGAGGGGACAGCTGCTGATTACAAGAGTACACAGAGCGCAATCGACCAGA TCACCGGGAAACTAAATCGCCTCATCGAGAAGACTAATCAGCAGTTCGAACTGATCGACAACGAGTTCAACGAGGTCGAAAAACAGATTG GTAACGTCATTAACTGGACCAGAGATTCCATAACCGAAGTCTGGTCCTATAACGCAGAACTGCTGGTAGCCATGGAGAACCAGCACACTA TCGACCTGGCAGACTCGGAAATGGATAAACTGTACGAGAGGGTGAAGCGGCAGCTCCGGGAGAACGCTGAAGAAGATGGTACCGGCTG TTTCGAAATTTTTCACAAATGTGACGATCATTGTATGGCCTCTATACGCAACAATACCTACGATCACTCCAAATACAGGGAGGACGCCTATG CAGAATCGTATCCAGATCGATCCGGTAAAACTGAGTAGCGGATATAAGGATGTGATCCTGTGGTTCAGCTTCGGGGCTAGTTGCTTTATT CTGTTGGCAATTGTGATGGGCCTGGTGTTCATTTGCGTCAAAAATGGCAACATGCGCTGCACCATTTGTATC
2751	ATGAACACCCAAATATTAGTGTTCGCATTGATTGCGATTATCCCAACCAACGCAGATAAGATCTGCTTGGGACATCACGCTGTCAGCAAC GGGACCAAGGTGAACACGTTGACAGAGAGGGGCGTAGAAGTGGTCAACGCCACTGAGACAGTGGAACGCACCAACATACCGCGGATCT GCTCCAAGGGAAAGCGCACGGTGGATCTTGGCCAGTGCGGCCTGCTGGGTACGATAACTGGACCTCCCCAGTGCGACCAGTTCCTGGA GTTCTCCGCCGACCTCATCATCGAAAGGAGAGAAGGTAGCGATGTCTGCTACCCAGGGAAATTCGTTAACGAAGAGGCCCTTAGACAAA TCCTGAGAGAATCTGGCGGGATCGATAAGGAGGCAATGGGATTTACTTACTCAGGAATTAGGACTAACGGGGCTACCTCTGCGTGCCGA CGGTCCGGTTCGAGCTTCTATGCCGAGATGAAATGGCTTTTGAGCAACACCGATAATGCTGCTTTTCCCCAGATGACTAAGTCCTATAAG AATACCAGAAAAAGCCCCGCCTTAATTGTGTGGGGCATCCACCACTCAGTGAGTACAGCCGAACAAACGAAACTGTACGGCTCTGGTTC CAAGTTGGTCACAGTCCGATCCTCAAATTATCAGCAGAGCTTCGTACCCTCTCCGGGTGCGAGACCACAGGTGAACGGCCTGAGTGGTC GGATTGACTTCCACTGGCTCATGCTGAACCCCAACGATACCGTAACTTTCTCTTTCAACGGAGCCTTTATAGCTCCTGATAGAGCTTCTTT TCTGCGTGGGAAGTCCATGGGGATCCAGAGCGGCGTGCAGGTGGACGCGAACTGTGAGGGTGACTGCTATCACTCAGGAGGAACAATT ATCTCGAATCTTCCTTTTCAAAACATCGACTCCCGAGCCGTCGGAAAATGCCCTCGCTATGTGAAACAGCGGAGCTTGTTGCTCGCCACG GGGATGAAGAACGTCCCTGAAATACCGAAGGGGCGGGGCCTGTTTGGAGCAATCGCAGGATTTATCGAGAACGGGTGGGAAGGGCTTA TTGATGGGTGGTATGGCTTCCGCCACCAAAACGCTCAAGGCGAGGGGACTGCTGCTGATTATAAATCGACACAGTCTGCTATTGACCAG ATTACAGGAAAGCTCAACCGATTGATAGAAAAGACTAACCAGCAGTTTGAGCTGATCGATAATGAATTTAACGAGGTGGAGAAGCAGATC GGGAATGTGATCAACTGGACCAGAGATTCCATCACTGAAGTTTGGAGTTACAACGCCGAGCTCCTGGTGGCTATGGAAAATCAGCATACC ATCGACCTCGCTGACAGTGAGATGGACAAGCTCTATGAAAGAGTTAAGCGACAGCTGCGAGAGAATGCCGAGGAGGATGGTACAGGCT GCTTTGAGATTTTCCACAAGTGTGATGATGACTGTATGGCCTCAATACGGAATAATACATATGATCACTCCAAGTATCGGGAGGAAGCCAT GCAGAACCGTATCCAGATCGACCCTGTGAAACTGAGCTCTGGATACAAGGATGTCATACTCTGGTTCTCATTCGGTGCTTCTTGTTTTATT CTGCTGGCCATTGTGATGGGCCTCGTGTTTATATGCGTAAAGTCCCGCAACATGAGATGCACCATCTGTATC

2752	ATGAATACTCAGATACTGGTATTCGCTCTCATTGCTATTATTCCTACGAACGCAGATAAGATTTGCCTAGGCCATCATGCTGTCTCTAACG GCACTAAGGTCAACACCCTGACCGAGCGCGGAGTGGAGGTAGTTAACGCAACTGAAACAGTCGAACGCACCAATATTCCCAGAATCTGC TCGAAGGGAAAGAAGACCGTCGATCTGGGGCAATGTGGGCTTCTCGGCACCATCACGGGCCCTCCACAGTGTGACCAGTTTCTGGAATT TTCCGCCGACCTGATCATAGAGCGCCGGGAAGGATCCGATGTGTGTTATCCCGGGAAGTTCGTGAACGAGGAAGCACTGCGACAGATC CTGAGGGAGTCCGGCGGGATTGACAAGGAGGCCATGGGCTTCACTTATAGCGCTATACGTACCAACGGTGCAACCAGTGCTTGCAGGC GATCCGGGTCCTCCTTCTACGCTGAGATGAAGTGGCTCCTCTCCAATACAGACAACGCCGCCTTTCCACAGATGACTAAAAGTTATAAGA ACACACGCAAGTCCCCCGCGCTGATCGTGTGGGGGATTCATCATAGCGTGAGCACCGCGGAGCAGACAAAACTCTACGGAAGCGGTAA CAAGCTGGTAACAGTGGGATCTTCTAACTACCAACAGAGTTTCGTGCCTTCCCCAGGTGCGCGCCCCCAGGTAAACGGACAGAGTGGGA GAATCGACTTCCACTGGCTGATGCTGAATCCCAACGATACTGTCACCTTTAGTTTTAATGGTGCATTTATCGCCCCTGATAGAGCATCCTT CTTACGGGGGAAGAGTATGGGGATTCAGTCAGGCGTGCAGGTTGACGCTAACTGCGAGGGCGATTGTTACCACTCGGGCGGGACTATA ATCTCGAACCTTCCCTTTCAAAACATAGATAGCCGGGCCGTGGGCAAATGTCCGAGATACGTGAAACAGAGGAGTCTGTTGCTGGCTAC GGGCATGAAGAACGTGCCAGAAATTCCTAAGGGCCGCGGTCTGTTCGGAGCAATCGCAGGGTTCATCGAGAATGGCTGGGAAGGGCTG ATCGATGGTTGGTATGGATTCAGACACCAGAACGCGCAGGGGGAGGGAACAGCCGCTGACTATAAATCCACACAGAGCGCTATAGATCA AATCACAGGGAAACTTAACAGGCTGATCGAGAAAACAAACCAGCAGTTTGAGTTGATTGATAATGAGTTCAACGAGGTGGAGAAGCAAAT CGGCAACGTCATCAATTGGACTCGCGATTCTATCACTGAGGTATGGAGCTACAATGCAGAGCTGCTCGTAGCGATGGAGAATCAGCATA CAATTGATCTGGCCGACTCCGAAATGGACAAGCTGTATGAGCGCGTAAAGAGACAACTTAGAGAAAATGCCGAGGAAGATGGCACTGGC TGCTTCGAAATCTTTCATAAATGCGATGATGACTGTATGGCATCTATCAGAAACAACACTTATGACCACTCCAAGTATAGGGAAGAAGCTA TGCAGAATAGAATCCAGATCGATCCCGTAAAACTTAGTTCCGGGTACAAGGACGTTATTCTTTGGTTTTCTTTCGGCGCAAGCTGTTTCAT CCTTCTTGCAATCGTAATGGGTCTCGTCTTCATCTGCGTTAAAAATGGCAACATGCGTTGCACAATTTGCATT
2753	ATGAATACCCAGATCCTGGTCTTTGCGCTGATCGCTATTATCCCAACTAACGCCGATAAAATCTGCCTGGGCCACCACGCTGTCAGCAAT GGCACAAAGGTAAATACCCTAACAGAGCGGGGGGTGGAAGTCGTTAATGCTACTGAGACCGTGGAACGAACAAATATTCCTAGGATCTG TTCCAAGGGCAAGCGCACTGTGGACTTGGGTCAGTGCGGCTTACTTGGTACAATCACTGGGCCGCCACAGTGCGACCAGTTTCTCGAGT TCTCGGCCGATCTGATCATAGAAAGAACGGAGGCAGCGACGTGTGCTACCCCGGCAAGTTTGTTAACGAGGAGGCACTGCGGCAGAT CCTGAGAGAAAGCGGAGGCATCGACAAAGAGGCAATGGGGTTCACCTACTCTGGCATTAGGACTAACGGTGCAACTTCTGCCTGCCGGA GAAGCGGTTCTTCCTTCTACGCCGAAATGAAGTGGCTGCTGTCCAATACTGACAACGCTGCCTTTCCTCAGATGACAAAGAGTTACAAAA ACACACGGAAAAGCCCAGCTCTGATCGTGTGGGGCATCCACCATAGTGTAAGCACAGCTGAACAGACGAAGCTGTACGGGTCTGGCAAT AAACTTGTCACAGTTGGATCGTCCAATTACCAACAGAGCTTCGTCCCCAGTCCAGGTGAGAGACCTCAGGTTAACGGCCTCTCGGGGAG AATCGACTTCCACTGGCTGATGCTTAACCCCAACGACACAGTAACCTTCTCCTTTAACGGCGCCTTTATTGCTCCTGATAGGGCCTCCTTT TTGAGAGGAAAAAGTATGGGCATCCAAAGCGGGGTGCAGGTGGACGCCAACTGTGAAGGCGACTGCTACCACTCGGGGGGCACCATAA TATCCAACCTACCTTTTCAGAATATTGACTCTCGGGCCGTGGGGAAGTGTCCTCGTTACGTGAAGCAGCGAAGCCTCCTACTCGCAACCG GCATGAAGAACGTCCCAGAGATCCCAAAGGGTCGGGGCCTGTTCGGTGCCATTGCCGGGTTTATAGAAAACGGGTGGGAAGGCCTGAT CGATGGCTGGTATGGCTTCCGCCATCAGAACGCACAGGGTGAGGGCACAGCAGCAGACTATAAAAGCACTCAATCGGCCATCGACCAG ATCACGGGGAAGTTAAATAGGCTGATTGAAAAGACAAACCAACAATTCGAACTTATTGATAACGAGTTTAACGAAGTGGAGAAGCAGATA GGAAACGTTATCAATTGGACGCGAGATTCCATAACAGAAGTTTGGTCATATAATGCCGAGCTCCTAGTCGCCATGGAAAATCAGCACACT ATTGATCTGGCTGATTCCGAGATGGATAAATTATATGAGAGAGTGAAGCGCCAGCTTAGGGAAAACGCCGAAGAGGACGGAACTGGGTG TTTCGAGATCTTTCATAAATGCGATGATGACTGCATGGCCTCTATTCGGAATAACACATACGATCACTCCAAGTATCGGGAGGAGGCGAT GCAGAACAGAATTCAGATTGATCCAGTGAAGCTGTCCAGCGGATACAAAGACGTGATACTATGGTTCAGCTTTGGTGCTAGTTGCTTTAT CCTGTTGGCGATCGTCATGGGCCTGGTATTCATTTGCGTGAAGAATGGTAACATGCGCTGCACCATATGTATT
2754	ATGAACACACAGATACTGGTGTTCGCACTGATAGCCATAATACCTACCAATGCTGATAAAATTTGTCTTGGTCACCATGCCGTCAGCAATG GAACAAAAGTCAATACTCTGACAGAGAGGGGAGTCGAAGTGGTGAACGCAACAGAGACGGTAGAACGGACAAACATTCCAAGAATCTGT TCCAAAGGTAAGAGGACGGTCGATCTCGGACAGTGCGGCCTATTACGAACTATTACTGGTCCTCCGCAGTGCGACCAGTTTCTGGAATTT AGCGCAGACTTGATTATAGAGCGGCGAGAAGGGTCCGATGTATGCTATCCCGGCAAATTCGTGAATGAGGAGGCCCTGCGACAGATTTT GAGAGAAAGCGGGGGGATTGATAAAGAAGCGATGGGGTTTACATATTCAGGCATCAGGACCAATGGCGCTACCTCAGCTTGCCGAAGAA GTGGTAGCTCTTTTTATGCCGAAATGAAGTGGCTTCTTTCAAACACTGATAACGCAGCCTTCCCACAGATGACCAAGAGCTACAAGAATAC TCGTAAATCACCAGCCCTAATTGTGTGGGGTATCCACCATTCAGTCTCTACCGCAGAACAAACAAAACTGTACGGCAGCGGGAACAAGCT AGTGACCGTTGGATCCAGCAATTACCAGCAAAGCTTTGTCCCTTCCCCCGGCGCACGTCCACAGGTAAACGGCTTGAGCGGGAGAATTG ATTTCCACTGGCTAATGCTAAATCCCAACGATACCGTTACTTTTAGCTTCAATGGGGCTTTCATTGCCCCAGATAGAGCCTCTTTCCTGAG AGGTAAAAGTATGGGCATCCAATCCGGGGTGCAGGTGGATGCGAATTGTGAAGGCGATTGTTACCACAGCGGGGGGACCATTATTTCCA ACTTACCCTTCCAAAACATCGATTCGCGAGCAGTGGGCAAATGTCCCCGGTATGTCAAGCAGAGGTCGCTGTTGCTGGCCACCGGCATG AAAAATGTCCCTGAGATTCCAAAAGGTCGCGGTTTGTTTGGCGCTATTGCGGGATTTATCGAGAACGGCTGGGAGGGGCTGATTGACGG GTGGTACGGGTTCAGACACCAGAACGCTCAGGGCGAGGGAACCGCTGCGGATTATAAAAGCACGCAGTCGGCCATAGACCAGATCACA GGCAAGCTCAATAGGCTCATCGAGAAGACGAATCAGCAATTCGAATTGATAGATAACGAATTCAACGAGGTGGAGAAACAGATTGGGAAT GTGATCAATTGGACAAGGGATTCAATTACGGAAGTGTGGTCCTATAACGCCGAGCTGCTCGTGGCCATGGAGAATCAGCATACAATCGAT CTGGCCGACAGTGAGATGGATAAACTCTACGAACGTGTGAAGCGGCAGCTACGCGAGAACGCCGAGGAAGATGGGACAGGCTGTTTTG AGATCTTCCACAAGTGCGACGACGACTGTATGGCGTCAATCCGCAATAACACATACGACCACAGCAAGTACCGAGAGGAAGCCATGCAA AACAGGATCCAAATCGATCCTGTAAAGCTTTCCTCTGGGTATAAAGACGTCATTCTTTGGTTCTCTTTTGGAGCCAGTTGCTTCATCCTTCT AGCCATTGTGATGGGGCTGGTATTCATCTGCGTGAAAAATGGGAATATGCGATGTACCATATGCATT

2755	ATGAATACCCAGATCCTCGTTTTTGCTCTGATCGCAATTATTCCCACTAATGCAGACAAAATCTGTCTCGGTCATCATGCTGTGTCTAATG GAACGAAGGTCAATACCCTGACCGAACGCGGTGTCGAAGTGGTCAATGCGACAGAAACTGTAGAACGAACCAATATCCCTAGGATTTGTT CCAAGGGGAAGAAGACTGTTGACTTGGGGCAATGCGGGCTCCTAGGCACCATTACGGGCCCCCCCCAATGTGACCAGTTCCTAGAGTTT AGCGCAGATCTGATCATCGAAAGAAGGGAAGGCTCCGATGTGTGCTATCCCGGGAAGTTCGTGAATGAAGAGGCGCTGAGGCAGATTCT CAGAGACAGCGGCGGTATTGAGAAAGAAGCTATGGGATTTACATATAGCGGGATTCCGCCAAACGGCGCAACTAGTGCGTGTCGACGC AGCGGTTCTTCTTTCTATGCCGAGATGAAGTGGTTACTAAGCAACACCGATAACGCCGCGTTCCCCCAGATGACCAAGTCTTATAAGAAC ACACGCAAATCCCCCGCGCTCATTGTGTGGGGAATTCACCACAGCGTCTCTACTGCAGAGCAGACAAAACTGTACGGTTCAGGGAATAA GCTCGTAACTGTAGGGTCAAGTAACTACCAGCAGTCCTTTGTCCCTTCTCCAGGGGCTCGGCCTCAGGTCAATGGGCTGAGCGGCCGGA TCGATTTCCACTGGCTGATGCTGAATCCCAACGACACCGTGACTTTCTCTTTCAACGGGGCCTTTATCGCCCCAGATCGGGCCAGCTTTC TGAGAGGGAAGTCAATGGGAATCCAGTCTGGAGTCCAGGTGGATGCGAACTGTGAAGGGGATTGCTACCACAGCGGCGGGACTATCAT CTCCAACCTTCCATTCCAAAACATTGACTCTCGCGCCGTTGGCAAGTGTCCACGATACGTCAAGCAGAGGTCTCTGCTCCTGGCTACCGG AATGAAGAATGTGCCCGAGATACCCAAAGGCCGTGGCCTGTTCGGCGCGATAGCTGGCTTTATTGAAAACGGATGGGAGGGATTGATAG ACGGCTGGTACGGCTTTCGCCATCAGAACGCGCAGGGCGAAGGAACCGCCGCCGATTATAAATCAACACAGTCAGCTATCGACCAGATA ACCGGCAAGCTGAACCGATTAATTGAAAAAACAAACCAGCAGTTTGAACTGATCGATAATGAATTCAACGAAGTGGAAAAACAGATTGGG AATGTTATAAATTGGACTCGGGACTCCATCACTGAAGTTTGGTCTTACAATGCCGAATTACTGGTTGCCATGGAAAATCAACACACTATCG ATCTCGCGGACAGCGAAATGGACAAGCTTTACGAACGCGTTAAACGGCAGCTTCGTGAGAATGCGGAAGAGGATGGAACGGGCTGCTT CGAGATCTTCCATAAATGCGACGACGATTGTATGGCCTCAATTAGGAATAACACCTACGATCACAGCAAGTACCGGGAGGAGGCCATGC AGAACCGCATTCAAATTGATCCCGTCAAACTCTCCAGCCGGCTATAAAGATGTGATCTTGTGGTTCTCTTTTGGTGCTTCCTGCTTTATCCT ACTGGCCATTGTCATGGGACTGGTGTTCATCTGCGTCAAAAACGGGAACATGCGCTGTACCATCTGTATT
2756	ATGAACACCCAGATCCTTGTCTTTGCTCTGATTGCTATCATCCCTACAAATGCGGACAAGATTTGTCTGGGTCATCATGCAGTATCCAACG GAACCAAAGTCAACACCCTAACCGAACGCGGCGTAGAAGTGGTCAACGCCACCGAGACCGTGGAACGCACAAATATTCCTAGGATCTGC TCAAAAGGTAAAAAGACAGTGGATTTGGGCCAGTGTGGTCTGCTGGGGACTATAACTGGCCCCCCCCAATGCGATCAGTTCCTTGAATTT AGCGCCGATCTCATTATAGAACGGCGGGAGGGCAGCGATGTTTGCTATCCAGGAAAGTTTGTCAATGAAGAAGCACTAAGGCAGATTCT GAGAGAATCTGGAGGAATCGACAAGGAGGCCATGGGATTCACGTACTCCGGCATTCGCACAAATGGGGCCACCTCAGCTTGTAGGCGC AGTGGCAGCAGCTTTTACGCGGAGATGAAGTGGTTGCTTTCCAATACAGATAACGCTGCATTCCCCCAGATGACAAAAAGCTACAAGAAT ACACGAAAATCCCCCGCTCTGATAGTGTGGGGGATCCACCATAGTGTGTCAACCGCCGAGCAAACGAAGTTGTATGGCTCCGGAAACAA ATTGGTGACTGTCGGTTCATCCAACTACCAGCAGTCTTTTGTGCCTTCTCCCGGGGCCCGACCTCAGGTAAATGGTCAGAGCGGTCGGA TCGACTTCCACTGGCTCATGCTGAATCCCAACGACACAGTTACTTTTTCATTCAATGGGGCATTCATTGCGCCAGACAGGGCCTCCTTTCT AAGGGGTAAGTCGATGGGAATACAGTCTGGCGTGCAAGTAGATGCAAATTGCGAAGGGGACTGCTATCACTCTGGGGGCACGATCATCA GTAACCTGCCTTTTCAGAACATCGACAGCCGAGCAGTAGGAAAATGCCCACGCTACGTGAAGCAGCGCTCACTGCTCTTAGCCACCGGG ATGAAGAACGTCCCAGAAATTCCTAAAGGAAGGGGGCTGTTCGGCGCAATTGCTGGGTTTATTGAGAATGGATGGGAGGGTCTGATCGA TGGCTGGTACGGTTTCAGACACCAGAACGCCCAAGGCGAAGGCACAGCCGCTGACTATAAGAGTACTCAAAGTGCAATTGACCAGATTA CCGGCAAGTTGAATCGTCTGATTGAAAAGACAAATCAGCAGTTCGAACTGATAGATAATGAGTTTAATGAGGTTGAGAAACAGATAGGCA ACGTGATTAACTGGACCAGGGACTCAATCACAGAAGTGTGGAGCTACAACGCGGAATTACTAGTGGCAATGGAAAATCAGCATACTATCG ACCTTGCAGATTCTGAAATGGACAAACTGTATGAAAGAGTGAAACGCCAGCTCCGTGAAAATGCCGAAGAGGACGGGACCGGTTGTTTT GAGATTTTCCATAAGTGTGACGACGATTGTATGGCAAGCATCCGTAATAATACATACGATCATTCGAAGTATCGCGAGGACGCAATGCAG AACAGGATACAAATCGATCCTGTAAAACTGAGTAGTGGGTATAAAGATGTGATTTTGTGGTTCTCATTTGGCGCCTCTTGTTTCATCCTCC TCGCTATCGTGATGGGACTGGTGTTTATCTGCGTGAAGAATGGTAACATGCGGTGCACTATATGCATC
2757	ATGAATACCCAGATCCTAGTGTTTGCCCTTATTGCCATCATACCGACAAACGCCGACAAGATATGCCTAGGACATCACGCTGTCTCAAAC GGTACCAAGGTTAACACGTTGACGGAGCGAGGAGTTGAGGTGGTGAACGCTACTGAGACCGTGGAACGCACTAACATTCCCCGCATTTG TAGTAAAGGCAAGAGAACTGTCGACCTGGGACAGTGCGGCCTCTTAGGGACCATCACAGGACCCCCGCAGTGTGACCAGTTCCTCGAAT TTTCTGCTGACCTGATCATTGAGCGCGAGAGAGGGGAGCGATGTGTGCTACCCCGGAAAATTCGTAAATGAAGAAGCACTGCGACAGATT CTGCGGGAGAGTGGCGGAATTGATAAGGAAGCTATGGGGTTTACATACAGCGGGATCAGGACTAACGGAGCGACCAGCGCTTGCAGGC GGTCAGGAAGTTCTTTCTATGCAGAAATGAAGTGGCTGCTGTCCAACACGGACAATGCTGCTTTCCCCCAGATGACTAAATCCTACAAAA ATACTCGAAAATCTCCCGCCCTTATTGTTTGGGGGATCCATCACAGCGTGTCGACCGCCGAACAGACAAAGCTCTATGGAAGCGGTAACA AGCTTGTGACCGTTGGATCTTCCAATTATCAGCAGTCCTTTGTCCCCTCTCCTGGTGCCAGACCCCAGGTTAATGGCCTAAGCGGGCGCA TCGACTTTCATTGGCTCATGCTGAACCCTAATGATACCGTTACGTTTAGTTTTAATGGCGCCTTCATTGCCCCTGACAGAGCATCTTTCCT GCGAGGGAAGTCAATGGGCATCCAGAGCGGTGTACAGGTGGACGCCAATTGTGAGGGCGATTGTTATCACTCCGGGGGAACAATTATTA GTAATCTGCCCTTCCAGAATATAGATTCGAGGGCGGTAGGTAAGTGTCCTCGCTACGTTAAACAACGCTCTCTTCTTTTGGCCACTGGGA TGAAGAACGTACCTGAAATACCTAAGGGTCGCGGGCTCTTTGGCGCGATCGCAGGCTTCATTGAAAACGGCTGGGAGGGTTTAATCAAC GGGTGGTACGGTTTCCGACACCAGAACGCTCAGGGGGAAGGTACCGCTGCAGATTACAAGTCCACTCAATCAGCTATTGATCAGATCAC GGGCAAGCTGAACAGGCTCATCGAAAAGACCAATCAGCAATTCGAACTCATCGATAACGAGTTCAACGAGGTGGAGAAGCAAATTGGAA ATGTAATAAATTGGACGCGCGATTCTATTACAGAAGTGTGGTCCTACAACGCGGAACTCTTAGTGGCAATGGAGAACCAGCATACTATTG ACCTGGCTGACTCAGAAATGGATAAGCTGTACGAAAGGGTGAAAAGACAGTTGAGGGAAAATGCTGAGGAACATGGGACGGGATGCTTT GAGATTTTCCATAAGTGCGATGATGACTGCATGGCAAGCATCCGTAACAACACATATGACCATTCTAAGTACCGGGAGGAAGCCATGCAA AATCGCATACAGATCGACCCTGTGAAGCTCAGTTCTGGCTACAAAGATGTGATCCTGTGGTTTTCGTTCGGAGCCAGCTGCTTTATCCTA CTTGCAATAGTCATGGGTCTTGTGTTCATATGCGTAAAGAATGGCAACATGAGGTGTACCATCTGTATC

2758	ATGAACACCCAGATTCTGGTGTTTGCCCTGATTGCTATAATTCCAACCAATGCGGACAAAATCTGCCTGGGCCACCACGCCGTGTCGAAC GGCACGAAAGTGAATACCTTAACGGAAAGAGGCGTTGAAGTTGTGAATGCAACAGAAACTGTAGAAAGAACTAATATTCCTAGAATCTGTT CTAAGGGGAAAAAAACGGTGGACTTAGGCCAGTGTGGTCTGCTAGGGACTATCACCGGCCCCCCTCAATGTGACCAGTTCCTGGAGTTC AGCGCCGACCTCATTATAGAAAGAAGAGAAGGATCAGACGTGTGCTACCCTGGCAAATTCGTAAACGAGGAGGCCCTCCGGCAGATCCT CCGCGAATCCGGTGGAATCGACAAGGAAGCCATGGGATTTACCTATTCTGGTATTCGGACCAACGGCGCCACCAGTGCTTGTCGGCGGT CAGGGTCTTCCTTTTACGCAGAGATGAAGTGGCTCCTGTCCAATACTGACAATGCCGCATTCCCTCAAATGACTAAGTCCTATAAAAATAC CAGAAAATCTCCTGCACTTATCGTGTGGGGCATTCATCACTCTGTCAGTACTGCCGAGCAGACCAAGCTATACGGGAGCGGTAACAAGCT TGTGACAGTAGGGTCATCCAACTATCAGCAAAGCTTTGTACCGTCCCCTGGGGCACGTCCCCAGGTGAACGGCCAGTCCGGGCGGATC GACTTCCACTGGTTGATGCTGAATCCTAACGATACCGTGACCTTCTCTTTCAACGGGGCATTTATCGCACCAGACAGGGCTTCCTTTCTG CGCGGAAAGTCCATGGGCATCCAGTCCGGCGTGCAGGTCGATGCAAACTGCGAGGGTGATTGCTACCATTCTGGGGGGACAATAATCT CTAATCTGCCTTTTCAGAACATTGACAGTCGAGCGGTGGGCAAGTGTCCCAGGTACGTGAAACAACGCTCCCTTCTGTTGGCAACAGGG ATGAAGAACGTGCCTGAGATCCCGAAAGGCCGTGGATTATTTGGCGCTATTGCTGGCTTCATTGAGAATGGATGGGAGGGCTTAATCGA CGGTTGGTACGGCTTCCGGCATCAGAATGCACAGGGCGAAGGAACAGCTGCCGACTATAAATCCACCCAGTCCGCAATCGATCAGATAA CCGGCAAGCTGAACCGGCTCATTGAGAAAACAAATCAGCAGTTCGAGCTCATAGACAATGAGTTTAACGAGGTAGAGAAACAGATTGGTA ATGTCATTAACTGGACACGGGATAGCATCACGGAAGTTTGGAGCTATAATGCTGAGCTCTTGGTCGCTATGGAGAACCAGCATACGATTG ACTTGGCAGACTCTGAGATGGATAAGCTTTATGAGCGCGTGAAGAGGCAACTTAGAGAGAACGCTGAGGAGGACGGTACTGGGTGCTTC GAGATATTCCATAAGTGCGACGATGACTGCATGGCCAGCATTAGAAACAACACATATGATCACTCCAAGTATCGTGAAGAAGCCATGCAG AATCGGATTCAGATTGATCCAGTGAAGCTGTCCTCCGGCTACAAGGATGTGATCCTGTGGTTCTCGTTCGGTGCTAGCTGCTTCATATTG TTAGCTATCGTTATGGGCCTCGTGTTCATCTGCGTGAAGAACGGTAACATGCGTTGCACTATCTGCATA
2759	ATGAACACTCAAATTCTGGTGTTCGCACTCATTGCCATCATACCAACTAACGCTGACAAGATATGCTTGGGCCATCACGCCGTGAGCAAT GGGACAAAGGTAAACACCCTCACAGAAAGAGGCGTCGAAGTGGTTAATGCTACAGAGACAGTCGAGAGGACAAACATCCCCCGTATTTG TTCCAAGGGTAAGAAAACTGTTGATCTGGGCCAGTGTGGGTTGTTAGGAACCATCACCGGACCCCCGCAATGTGATCAATTCCTGGAATT TTCTGCGGATCTGATTATTGAGAGGCGCCAAGGCAGCGATGTGTGTTATCCCGGAAAGTTTGTTAACGAAGAAGCACTGCGGCAGATTCT GCGAGAATCCGGTGGCATCGACAAAGAGGCTATGGGCTTTACTTATTCCGGCATCAGGACGAATGGAGCCACCTCTGCCTGCCGCCGG TCAGGGAGTTCCTTCTATGCAGAGATGAAGTGGCTGCTCTCCAATACAGATAATGCCGCATTCCCTCAGATGACGAAATCTTATAAAAACA CACGGAAGAGTCCCGCACTGATCGTCTGGGGAATTCACCACTCCGTGAGCACTGCAGAGCAAACCAAGCTGTACGGGTCAGGGAACAA GTTAGTGACAGTCGGAAGCTCTAATTATCAACAGTCGTTCGTGCCAAGTCCTGGGGCCCGACCGCAGGTGAATGGTCAGTCAGGGCGTA TCGATTTCCACTGGTTAATGCTGAATCCAAACGACACCGTGACCTTCTCCTTTAATGGCGCCTTTATTGCACCCGATAGAGCTTCCTTTTTA AGAGGAAAATCCATGGGAATTCAAAGTGGTGTGCAGGTAGATGCGAATTGCGAGGGAGATTGTTATCACTCCGGTGGGACCATTATCTC CAACCTCCCTTTCCAGAACATCGACTCGCGAGCTGTGGGCAAGTGCCCAAGATACGTTAAGCAGAGATCGCTTCTGTTGGCCACCGGAA TGAAGAACGTTCCTGAAATTCCTAAGGGCCGCGGATTATTCGGCGCCATTGCAGGCTTTATAGAAAACGGTTGGGAGGGCCTTATCGAT GGATGGTATGGCTTCAGACACCAGAATGCACAAGGGGAGGGGACTGCCGCCGACTATAAGTCGACCCAGAGCGCAATTGATCAGATTAC TGGGAAGCTTAATAGACTCATCGAGAAAACAAACCAACAGTTCGAACTCATTGACAACGAGTTTAATGAGGTTGAGAAGCAGATCGGTAA TGTCATCAACTGGACAAGAGACAGCATCACTGAGGTGTGGTCCTATAATGCTGAGCTGCTGGTAGCTATGGAGAACCAACACACAATTGA TCTCGCCGACTCAGAGATGGACAAGTTGTACGAGCGCGTCAAGAGGCAATTAAGAGAAAACGCCGAGGAGGATGGGACCGGCTGTTTT GAGATCTTCCATAAATGCGATGATCACTGTATGGCTAGCATCCGAAACAACACTTATGACCACTCCAAATATCGGGAAGAAGCAATGCAG AATCGAATTCAAATTGACCCAGTTAAACTCAGTTCCGGATATAAGGACGTTATCCTCTGGTTCAGCTTCGGAGCATCTTGCTTTATACTGC TGGCAATCGTTATGGGACTCGTGTTCATCTGTGTCAAGAACGGTAACATGAGATGTACTATCTGCATT
2760	ATGAACACACAAATCTTGGTCTTTGCTCTCATTGCAATAATACCGACAAATGCCGATAAAATCTGTCTAGGACACCACGCTGTATCCAATG GAACAAAGGTCAATACCTTGACAGAGCGCGGCGTAGAGGTGGTAAATGCCACAGAGACAGTCGAGCGAACAAATATTCCCCGCATATGT TCCAAAGGTAAAAACACCGTTGACCTCGGCAGTGCGGATTACTGGGAACAATCACGGCCCCCCACAGTGTGACCAATTTCTGGAGTT CAGTGCCGATTTAATCATAGAGCGCAGGGAAGGTTCCGACGTCTGTTATCCTGGCAAGTTTGTAAACGAGGAGGCCTTACGGCAAATACT GAGGGAGTCAGGCGGCATCGAGAAGGAAGCCATGGGCTTCACCTATTCCGGAATTCGGGCGAACGGCGCAACCTCGGCCTGCCGACG GAGCGGAAGCTCATTTTACGCCGAGATGAAGTGGCTGCTTTCCAACACTGACAACGCAGCTTTCCCTCAAATGACAAAGTCGTATAAGAA CACAAGAAAATCTCCTGCCCTCATTGTTTGGGGCATACACCATTCCGTGTCAACAGCAGAACAGACAAAACTCTACGGCTCGGGTAACAA GCTCGTGACAGTCGGTTCCAGTAATTACCAGCAGTCCTTTGTACCTTCCCCCGGTGCACGGCCCCAGGTGAACGGCCTCTCTGGCAGGA TTGATTTCCACTGGCTGATGCTGAATCCTAACGATACAGTCACTTTTAGCTTTAATGGAGCTTTCATTGCACCCGACCGCGCCAGCTTTTT GAGGGGAAAATCCATGGGGATCCAATCCGGAGTGCAGGTGGATGCTAACTGCGAAGGCGATTGTTATCATAGTGGGGGGACCATAATCA GTAACCTCCCTTTTCAGAATATCGACTCCCGCGCAGTAGGAAAATGCCCTAGGTACGTGAAACAGCGCTCCTTACTGCTCGCCACGGGC ATGAAGAATGTTCCTGAGATACCAAAAGGGCGGGGCCTGTTTGGGGCTATTGCGGGATTCATTGAAAACGGGTGGGAGGGGTTGATCGA TGGCTGGTACGGGTTCAGGCACCAGAATGCCCAGGGTGAAGGAACAGCAGCCGACTACAAGTCTACGCAGTCTGCTATCGATCAGATTA CCGGCAAGCTAAACCGCTTAATTGAAAAAACCAACCAACAGTTCGAACTGATCGACAATGAGTTTAACGAAGTGGAAAAGCAGATAGGGA ACGTGATTAATTGGACCAGAGACAGTATTACCGAGGTGTGGTCGTATAATGCCGAACTGTTGGTAGCTATGGAGAACCAGCACACGATAG ATCTGGCAGACAGTGAAATGGATAAGCTGTATGAGAGAGTAAAGCGGCAACTCCGGGAAAATGCAGAAGAGGATGGCACAGGATGCTTC GAGATATTCCATAAGTGTGATGACGATTGTATGGCCAGCATCAGAAACAATACCTACGACCACTCAAAATACCGCGAAGAGGCCATGCAG AATAGGATTCAAATTGACCCAGTGAAGCTCTCCTCGGGATACAAAGATGTGATCTTGTGGTTCAGCTTCGGCGCCTCCTGTTTTATCCTCC TGGCGATAGTGATGGGACTCGTCTTTATTTGCGTTAAGAACGGGAACATGAGGTGCACCATATGCATC

2761	ATGAACACCCAGATCCTAGTGTTTGCACTTATCGCCATCATCCCGACTAATGCGGACAAGATCTGCCTGGGCCATCACGCTGTGTCTAAT GGAACCAAGGTGAATACGCTGACGGAACGAGGAGTGGAGGTGGTAAATGCTACGGAAACAGTCGAGAGAACCAACATCCCACGGATCT GCTCTAAGGGAAAAAAGACCGTTGATCTGGGCCAATGCGGTCTACTCGGTACCATTACCGGACCACCACAATGCGACCAGTTTCTCGAAT TCAGCGCCGACTTGATCATAGAAAGGCGCGAGGGATCGGATGTGTGCTACCCTGGCAAATTCGTGAATGAGGAGGCCCTCCGCCAGAT CCTTAGAGAATCAGGTGGCATCGATAAGGAAGCTATGGGCTTCACATACTCCGCCATAAGAACAAATGGCGCTACCAGCGCCTGTCGAC GCTCGGGCAGCTCCTTTTATGCCGAGATGAAGTGGTTGCTCAGTAACACAGATAACGCAGCCTTCCCACAGATGACAAAGTCTTACAAGA ATACCCGTAAAAGCCCAGCCCTTATTGTGTGGGGAATACACCATAGCGTGAGCACGGCAGAGCAGACGAAACTGTATGGTAGCGGGAAC AAACTCGTTACGGTCGGGAGCTCAAACTACCAGCAGTCTTTTGTACCTTCACCCGGCGCTCGCCCACAGGTTAACGGTCAGTCGGGGAG AATCGATTTCCACTGGCTGATGCTGAACCCAAATGACACCGTCACCTTCTCCTTCAACGGAGCCTTCATTGCCCCAGATCGCGCCTCTTTT CTTAGAGGCAAGTCCATGGGCATCCAATCAGGCGTCCAGGTCGATGCCAATTGTGAAGGCGACTGTTATCATTCCGGAGGCACAATTATA AGCAACCTGCCCTTCCAGAACATCGACAGTCGAGCTGTCGGTAAGTGCCCTCGTTACGTGAAGCAGCGGAGTCTCCTCTTAGCTACAGG AATGAAAAATGTCCCAGAGATCCCTAAAGGACGCGGTCTCTTTGGGGCCATAGCTGGATTCATCGAAAACGGCTGGGAAGGCCTGATCG ACGGCTGGTACGGCTTCAGACACCAGAATGCACAAGGGGAAGGAACCGCTGCTGATTATAAGAGTACCCAGTCAGCTATCGACCAAATA ACCGGTAAGCTTAACCGACTGATTGAGAAAACAAACCAGCAATTTGAGCTCATTGATAATGAGTTTAATGAGGTGGAAAAGCAGATTGGC AATGTCATTAACTGGACCCGCGACTCGATTACAGAAGTGTGGTCTTATAACGCGGAGTTACTTGTCGCCATGGAAAATCAGCACACCATC GACTTGGCCGACTCTGAAATGGACAAACTGTACGAGCGAGTCAAGCGACAACTGAGGGAGAACGCCGAGGAAGACGGGACCGGTTGTT TTGAGATATTTCACAAATGCGACGATGACTGCATGGCCTCGATCAGGAACAATACTTACGACCACTCCAAGTACAGGGAAGAAGCGATGC AAAACAGGATACAGATAGATCCTGTTAAGCTCTCCAGTGGGTATAAAGACGTGATCCTGTGGTTCAGCTTCGGAGCATCCTGTTTTATCCT TTTGGCCATCGTAATGGGATTAGTCTTCATCTGCGTGAAGAATGGGAACATGCGGTGCACCATCTGTATA
2762	ATGAATACCCAGATATTGGTATTTGCCCTGATAGCGATCATTCCAACTAACGCAGATAAGATATGCCTCGGCCACCACGCTGTGTCGAAT GGGACTAAGGTGAATACCCTGACTGAAAGAGGTGTCGAGGTGGTCAACGCTACCGAAACAGTGGAAAGGACCAACATACCGAGAATCTG CAGCAAAGGTAAAAAGACTGTAGACCTCGGCCAGTGCGGCCTTCTGGGGACCATTACTGGACCTCCCCAATGCGATCAGTTCCTGGAGT TCTCTGCCGACCTCATCATTGAGCGGAGAGAGGCAAGCGATGTGTGCTACCCCGGCAAATTTGTGAATGACGAGGCTCTGAGACAGATT TTGAGGGAGAGTGGCGGGATCGAAAAGGAGGCAATGGGGTTTACCTACTCAGGAATCAGGGCCAACGGAGCAACCTCTGCATGCAGAA GGTCCGGGTCCTCCTTCTACGCAGAGATGAAATGGCTGTTAAGCAATACGGATAATGCCGCCTTCCCTCAGATGACTAAGTCCTATAAAA ATACCCGTAAGAGTCCAGCACTCATAGTCTGGGGCATTCACCATTCCGTCTCTACAGCAGAGCAGACTAAACTCTATGGAAGCGGGAACA AACTCGTGACCGTCGGCTCGAGCAACTATCAACAATCTTTCGTTCCATCTCCCGGAGCCAGACCCCAGGTGAACGGGCTATCGGGACGG ATCGACTTCCACTGGCTGATGTTAAACCCCAATGATACGGTCACCTTTAGCTTCAATGGTGCATTCATCGCTCCAGACCGGGCCTCGTTT CTCAGAGGAAAATCTATGGGAATCCAGAGTGGTGTGCAGGTGGACGCCAACTGCGAGGGTGATTGTTATCATTCTGGGGGCACTATCAT CTCAAACCTCCCATTCCAGAACATAGATTCAAGAGCAGTTGGAAAGTGTCCAAGATATGTGAAGCAGCGCTCTCTGCTTTTGGCCACCGG CATGAAAAATGTTCCTGAGATACCGAAAGGACGGGGCCTATTTGGCGCGATCGCAGGATTCATCGAAAATGGATGGGAGGGACTGATCG ACGGTTGGTATGGCTTCCGCCACCAGAATGCCCAGGGGGAAGGTACCGCAGCTGACTATAAATCAACGCAGTCCGCCATCGATCAGATA ACTGGCAAACTGAATAGGCTAATAGAAAAGACTAACCAGCAGTTTGAGCTGATTGACAATGAATTCAACGAAGTGGAGAAGCAGATTGGC AACGTTATTAACTGGACCCGGGACTCCATCACGGAGGTGTGGTCTTACAACGCAGAGTTACTAGTAGCAATGGAGAATCAGCACACTATA GACCTAGCCGATAGTGAAATGGACAAACTATATGAGAGAGTCAAACGCCAACTGCGGGAGAACGCCGAAGAAGATGGAACAGGATGTTT CGAGATTTTCCACAAATGTGATGATGATTGCATGGCGTCTATCCGCAATAATACTTACGATCATTCAAAGTACCGCGAAGAGGCCATGCA GAACCGAATCCAGATTGATCCCGTGAAGCTTTCCTCCGGATACAAAGATGTAATCCTGTGGTTCTCATTCGGCGCTTCCTGTTTCATCTTA CTGGCAATCGTCATGGGGCTGGTTTTCATTTGCGTCAAAAACGGCAATATGAGATGTACTATTTGCATC
2763	ATGAACACCCAGATCCTGGTTTTTGCCTTGATCGCAATTATTCCTACCAATGCGGACAAAATTTGCCTTGGGCACCACGCCGTTTCCAATG GCACAAAGGTCAACACTTTGACCGAACGGGGAGTGGAAGTGGTGAACGCTACAGAAACCGTTGAGCGGACTAACATTCCCAGGATCTGT AGTAAGGGCAAGCGCACCGTGGACCTGGGCCAGTGCGGATTGTTAGGCACCATCACCGGCCCACCCCAGTGCGACCAGTTCCTGGAAT TTAGTGCAGATTTAATCATAGAACGAAGAGAGGGTTCTGATGTGTGTTACCCGGGGAAATTCGTGAACGAAGAGGCCCTCCGCCAGATTT TGAGAGAGAGCGGAGGCATAGATAAAGAAGCAATGGGCTTTACCTATAGCGGGATCCGGACTAATGGAGCTACCTCTGCATGTCGGCGT AGTGGAAGTTCTTTCTATGCTGAGATGAAATGGCTCCTTTCTAACACAGACAATGCTGCATTTCCACAAATGACAAAATCCTATAAAAATAC TCGGAAATCCCCAGCCCTGATCGTTTGGGGAATCCACCATTCAGTTTCGACTGCAGAACAGACTAAATTGTATGGGTCCGGGTCTAAGCT GGTAACAGTTGGAAGTTCCAATTACCAGCAAAGCTTCGTCCCCTCGCCAGGTGCACGCCCACAGGTGAACGGGCTAAGCGGGCGTATTG ATTTCCACTGGTTAATGCTGAATCCAAACGACACTGTCACGTTCAGCTTTAACGGAGCGTTCATCGCTCCTGACCGCGCCAGTTTCCTGC GAGGTAAATCAATGGGCATCCAGTCAGGTGTTCAGGTAGATGCCAACTGTGAGGGGGATTGCTATCATTCTGGGGGGACTATTATTAGTA ACCTGCCCTTCCAGAATATAGATTCTCGGGCCGTCGGCAAGTGTCCAAGGTACGTTAAACAGCGCAGTCTGTTACTTGCCACTGGAATGA AGAACGTTCCCGAGATCCCAAAGGGCCGAGGTCTGTTCGGGGCCATCGCCGGCTTTATCGAGAATGGATGGGAAGGGTTGATTGATGG ATGGTATGGTTTCCGTCACCAAAATGCCCAGGGAGAGGGGACAGCTGCTGACTATAAGTCCACCCAGAGTGCAATAGACCAAATCACAG GGAAACTGAATAGACTTATCGAAAAGACCAACCAGCAGTTTGAGTTGATTGATAACGAATTCAATGAAGTCGAGAAACAGATTGGGAATGT CATTAATTGGACACGAGACTCCATCACAGAGGTGTGGTCCTACAATGCCGAATTGTTGGTGGCCATGGAGAATCAACATACTATCGATCT GGCAGACTCAGAAATGGACAAGCTGTATGAGCGAGTGAAGAGGCAGCTGACAGAGAACGCCGAAGAAGATGGAACCGGGTGCTTTGAG ATTTTTCATAAGTGTGACGACGATTGTATGGCAAGTATCCGCAATAATACCTACGACCACTCCAAGTATAGAGAGGAAGCCATGCAGAAC CGGATTCAGATTGATCCAGTGAAACTCAGTTCCGGCTATAAAGACGTGATCCTGTGGTTTAGCTTCGGCGCCTCCTGTTTCATCCTGTTG GCCATTGTCATGGGGCTGGTCTTTATCTGTGTCAAGTCCCGAAACATGCGATGCACAATCTGCATC

2764	ATGAACACTCAAATTCTAGTGTTCGCACTGATTGCCATTATCCCAACTAACGCCGACAAAATCTGTTTGGGCCACCATGCTGTTAGCAACG GTACCAAAGTGAATACCCTGACCGAAAGAGGCGTTGAAGTAGTAAACGCCACTGAAACCGTTGAGAGGACAAACATTCCACGGATTTGTA GCAAAGGAAAGAAAACTGTCGATCTGGGGCAATGTGGACTGCTGGGCACCATCACTGGCCCTCCCCAGTGTGACCAGTTTCTGGAGTTT TCCGCAGATTTAATCATAGAGCGCAGAGAAGGAAGTGACGTGTGCTACCCTGGGAAATTCGTTAATGAAGAAGCCCTCCGGCAGATCCT AAGACAATCTGGGGGTATCGAAAAGGAGGCAATGCGATTCACGTACTCGGGGATCCGTGCTAACGGAGCAACTTCCGCATGCCGGAGG TCAGGGTCATCTTTTTATGCTGAAATGAAATGGTTGCTGTCAAATACCGATAACGCCGCGTTCCCGCAAATGACAAAGAGCTACAAAAATA CCCGCAAGAGTCCCGCGCTCATCGTGTGGGGCATCCATCATAGCGTGTCCACTGCTGAGCAGACCAAACTTTATGGTAGTGGGAACAAG CTGGTCACGGTCGGATCAAGCAATTACCAGCAGTCCTTCGTACCAAGCCCGGGGGCCCGGCCCCAGGTGAACGGCCTGTCAGGACGCA TCGACTTTCACTGGTTGATGTTGAACCCGAATGATACAGTGACTTTTTCATTCAACGGCGCTTTTATCGCACCAGACAGGGCCTCCTTTCT CCGCGGGAAGAGCATGGGCATCCAGAGTGGTGTGCAGGTTGACGCGAATTGCGAGGGAGACTGTTATCACTCAGGCGGAACAATTATC AGCAATCTTCCTTTCCAAAACATTGATTCGAGAGCCGTCGGGAAGTGCCCCAGGTACGTCAAGCAGCGAAGTCTGCTTTTAGCTACAGGA ATGAAGAACGTGCCCGAGATACCCAAAGGCAGGGGACTGTTCGGCGCAATCGCTGGCTTTATTGAGAACGGATGGGAAGGTCTGATCG ATGGCTGGTATGGATTTAGGCACCAGAACGCTCAGGGCGAGGGCACTGCAGCCGATTATAAGAGCACGCAGTCTGCCATAGACCAAATC ACCGGGAAACTCAACAGGCTGATCGAGAAGACCAACCAGCAATTTGAGCTCATTGACAATGAATTTAATGAAGTGGAGAAGCAGATCGGC AATGTTATTAATTGGACACGAGACTCTATTACTGAAGTGTGGTCTTACAATGCCGAGCTGCTTGTTGCAATGGAGAATCAGCATACAATCG ATCTGGCCGATTCAGAGATGGACAAGCTGTACGAAAGGGTGAAACGGCAGCTACGGGAGAACGCTGAGGAGGATGGCACAGGGTGCTT TGAGATTTTCCATAAGTGCGACGACGATTGTATGGCGAGTATACGCAATAACACCTACGATCATTCAAAGTATCGCGAGGAAGCCATGCA GAACCGCATTCAGATCCATCCCGTGAAGTTGTCCTCTGGCTACAAAGACGTCATTCTGTGGTTCAGCTTTGGAGCATCCTGTTTCATCCTC CTGGCAATCGTGATGGGACTCGTGTTTATCTGCGTCAAAAACGGCAATATGAGGTGCACCATCTGTATT
2765	ATGAATACACAGATCTTGGTTTTCGCTCTGATCGCCATCATACCGACCAATGCTGACAAAATCTGTCTGGGTCACCATGCAGTGAGCAAC GGTACCAAAGTCAATACATTGACCGAGCGTGGCGTCGAAGTTGTTAATGCCACAGAAACAGTCGAGCGAACCAATATTCCGAGAATTTGC TCCAAGGGGAAGAAGACCGTGGACCTAGGCCAGTGTGGACTGCTGGGTACAATTACTGGGCCACCACAGTGTGATCAGTTCCTGGAGTT CAGCGCAGATCTGATCATCGAACGGAGAGAGGGATCAGATGTATGCTATCCTGGTAAATTCGTAAACGAAGAGGCCCTGCGACAGATTC TAAGAAAAAGCGGAGGCATCGATAAGGAAGCTATGGGGTTCACATATTCAGGCATCAGAACTAATGGTGCCACTTCCACTTGCCGACGAA GCGGGAGTAGCTTCTACGCAGAAATGAAATGGCTTCTATCAAACACGGACAACGCGGCCTTTCCTCAGATGACCAAATCCTATAAGAACA CCAGAAAAAGCCCCGCAATCATCGTATGGGGAATTCATCACTCTGTGAGTACCGCCGAGCAGACAAAGCTTTACGGCTCCGGAAACAAG CTGGTGACCGTTGGGAGTAGCAATTACCAGCAGTCTTTCGTACCATCCCCCGGTGCGAGACCCCAAGTTAACGGCCTTTCAGGACGCAT TGATTTCCACTGGCTGATGTTGAATCCGAACGACACCGTGACGTTCTCCTTCAACGGGGCATTCATTGCACCCGATAGAGCCTGTTTCCT CAGGGGAAAATCCATGGGCATACAGAGCGGGGTGCAGGTGGACGCCGACTGTGAGGGAGACTGCTACCACTCTGGTGGTACGATCATA AGCAACCTCCCATTCCAGAATATCGATTCTAGGGCGGTGGGTAAATGTCCAAGATATGTGAAGCAGCGGTCTCTTCTGCTGGCCACCGG CATGAAAAACGTTCCTGAGATTCCCAAAGGCAGAGGGCTGTTCGGCGCAATCGCCGGCTTTATAGAAAACGGATGGGAGGGCCTGATCG ATGGGTGGTACGGATTCAGACACCAGAATGCCCAGGGCGAAGGTACAGCTGCAGATTACAAAAGCACACAGTCAGCCATTGACCAAATC ACAGGCAAACTGAATCGCCTCATTGAGAAAACGAATCAGCAGTTCGAGCTCATTGATAACGAGTTCAACGAGGTCGAAAGACAAATTGGT AATGTGATCAACTGGACGAGGGACTCAATCACTGAGGTGTGGTCTTATAATGCAGAGCTGTTGGTGGCTATGGAGAATCAACACACAATT GACCTGGCGGATTCTGAAATGGATAAACTGTACGAACGAGTCAAGCGTCAGCTGAGGGAGAACGCTGAGGAAGACGGGACTGGATGCT TCGAAATTTTCCATAAATGCGATGACGATTGCATGGCCTCTATCCGAAACAATACATACGATCATAGTAAGTACAGAGAAGAGGCCATCCA GAACAGAATCCAGATTGACCCGGTGAAACTGTCATCGGGATACAAGGACGTCATCCTCTGGTTTTCTTTTGGTGCCTCTTGCTTCATATTA CTGGCAATTGTTATGGGACTGGTCTTCATATGTGTTAAGAATGGTAACATGCGGTGTACAATATGCATC
2766	ATGAACACTCAAATCCTCGTTTTCGCTCTGATTGCTATTATTCCTACTAATGCCGATAAGATTTGTCTGGGCCACCATGCAGTTTCCAATGG AACCAAGGTTAACACCCTGACGGAGCGTGGTGTTGAGGTGGTTAATGCAACTGAAACTGTTGAGCGTACCAATATCCCTAGAATATGCAG TAAGCGCGAAAAAAACTGTGGACCTCGGCCAATGTGGCCTGCTGGGCACCATCACAGGGCCGCCCCAATGTGATCAATTTTTAGAATTTA GCGCAGACCTGATTATCGAGCGCCGAGAAGGTAGCGACGTGTGCTACCCTGGCAAGTTTGTGAATGAGGAAGCCCTGAGACAAATTCTT AGGGAGTCAGGAGGCATCGACAAGGAAGCCATGGGCTTTACCTATAGCGGGATCAGAACAAATGGCGCAACTTCAGCCTGTAGGCGCT CGGGGTCCTCTTTCTACGCAGAGATGAAATGGCTGCTCAGCAATACAGACAATGCTGCTTTCCCTCAGATGACCAAGTCCTACAAAAATA CCCGGAAGTCGCCCGCTTTGATCGTTTGGGGAATTCACCATTCTGTCAGCACGGCCGAGCAGACTAAGCTATACGGGTCCGGTAACAAA CTGGTGACAGTCGGCAGCAGCAATTACCAGCAGTCTTTCGTGCCCTCCCCAGGCGCAAGGCCCCAGGTAAATGGTCAGTCTGGACGCAT AGATTTTCACTGGCTCATGCTCAACCCCAATGACACAGTTACCTTTTCCTTCAATGGAGCCTTTATCGCCCCGGATCGCGCCAGTTTTCTG CGGGGTAAATCCATGGGGATCCAGTCAGGGGTGCAGGTCGACGCTAACTGCGAGGGAGATTGTTACCACTCAGGAGGAACCATCATCA GCAATTTGCCCTTCCAGAATATCGATTCTAGGGCAGTGGGCAAGTGTCCAAGGTACGTAAAGCAGCGATCCCTCCTGCTGGCCACCGGT ATGAAAAATGTGCCAGAGATCCCTAAAGGTAGGGGGCTCTTCGGGGCTATCGCGGGCTTCATCGAGAATGGGTGGGAGGGACTCATCG ATGGATGGTACGGCTTCAGACACCAGAACGCACAAGGCGAGGGGACAGCAGCTGATTATAAATCCACCCAGTCCGCTATCGACCAGATT ACAGGGAAACTGAATCGGCTCATCGAGAAGACCAACCAGCAGTTCGAGCTGATTGACAATGAGTTCAATGAGGTTGAAAAGCAGATTGG CAACGTCATCAATTGGACTCGTGACAGCATCACCGAGGTTTGGTCCTACAACGCCGAGCTGCTCGTCGCCATGGAGAATCAACACACTAT AGACTTGGCCGACTCAGAAATGGATAAGCTATATGAGAGAGTTAAGCGACAGTTACGCGAGAACGCAGAAGAAGACGGTACCGGATGCT TTGAAATCTTTCACAAATGCGACGACGACTGTATGGCATCAATAAGAAACAACACCTACGATCACTCAAAGTACCGCGAGGAGGCCATGC AAAACCGGATCCAAATAGACCCCGTTAAGTTGTCCAGCGGGTATAAGGACGTGATTTTGTGGTTCTCTTTTGGGGCCTCCTGTTTCATTCT TCTGGCTATCGTGATGGGCCTGGTGTTCATATGTGTTAAGAATGGTAACATGAGATGTACTATCTGTATT

2767	ATGAATACACAAATTCTGGTCTTCGCTCTGATCGCCATAATTCCCACCAACGCTGATAAGATCTGTCTGGGCCACCACGCCGTCTCCAAT GGTACTAAAGTAAACACTCTCACTGAGCGCGGCGTTGAAGTGGTCAATGCTACCGAGACTGTGGAGCGGACAAACATCCCAAGGATTTG CTCTAAGGGAAAGAGAACTGTGGACTTGGGCCAGTGTGGATTACTGGGCACAATTACAGGCCCTCCCCAGTGCGATCAGTTCTTGGAGT TTAGTGCCGACCTAATAATTGAACGCAGAGAGGGATCTGACGTGTGCTATCCAGGGAAATTTGTTAATGAAGAGGCACTCCGCCAGATTC TGCGAGACTCCGGCGGGATCGACAAAGAGGCCATGGGATTCACGTATTCGGGCATCAGGACCAATGGCCGCACCTCCGCGTGTCGACG GAGCGGCAGTAGCTTTTATGCTGAGATGAAGTGGCTCCTCAGTAACACCGACAACGCTGCTTTCCCTCAGATGACCAAGTCATACAAGAA TACCCGGAAATCCCCTGCACTTATCGTGTGGGGAATCCATCATTCCGTTAGTACCGCCGAGCAAACTAAACTGTACGGGAGTGGCAACAA GCTTGTGACTGTGGGCTCATCGAATTATCAACAGTCGTTCGTCCCATCACCAGGCGCTAGGCCACAGGTTAATGGACTGTCGGGGCGCA TCGATTTTCACTGGTTAATGTTAAACCCTAATGACACCGTTACGTTTAGCTTCAATGGTGCCTTTATCGCCCCCGACCGGGCCAGCTTCCT CCGTGGAAAGTCCATGGGAATTCAGTCAGGCGTACAAGTGGATGCCAACTGTGAAGGCGATTGCTACCATTCTGGAGGAACCATTATTA GCAACCTGCCCTTCCAGAACATTGATAGCAGGGCAGTAGGTAAATGTCCCCGGTATGTGAAACAAAGAAGCCTGCTGCTCGCCACTGGC ATGAAAAACGTCCCTGAGATCCCGAAAGGTAGAGGCTTGTTCGGGGCAATTGCCGGGTTCATTGAAAACGGATGGGAAGGACTGATTGA CGGATGGTATGGGTTCCGGCACCAAAATGCTCAGGGAGAAGGGACCGCAGCCGATTATAAGTCAACCCAAAGCGCCATTGACCAGATTA CCGGAAAGCTTAACCGGCTCATCGAAAAGACAAATCAGCAGTTTGAGCTGATCGACAACGAATTTAATGAGGTGGAGAAGCAGATCGGA AACGTGATTAATTGGACTAGGGACAGCATCACCGAGGTCTGGTCTTATAACGCCGAACTGCTGGTAGCGATGGAAAATCAACACACCATC GACCTGGCTGACTCAGAGATGGATAAGCTGTATGAAAGGGTCAAACGTCAGCTTCGTGAAAATGCTGAAGAGGATGGCACAGGTTGCTT CGAAATCTTCCATAAGTGTGATGATGATTGTATGGCTAGTATTAGGAACAACACATACGACCATTCCAAATATCGGGAGGAAGCTATGCAA AACCGGATCCAAATCGACCCAGTTAAACTGTCTTCCGGCTACAAAGATGTTATTCTCTGGTTCAGTTTTGGTGCATCCTGCTTTATACTCCT TGCCATCGTAATGGGCCTGGTGTTTATCTGCGTCAAGAACGGAAACATGCGATGCACGATCTGTATC
2768	ATGAACACTCAGATCCTCGTATTCGCCTTGATCGCTATTATTCCAACTAACGCCGATAAAATCTGCCTTGGCCACCATGCCGTTAGCAACG GTACTAAAGTGAATACTCTTACAGAGAGGGGCGTCGAGGTCGTAAATGCCACTGAAACAGTTGAGCGAACAAACATTCCTCGAATTTGTT CTAAGGGAAAAAGAACAGTCGACCTCGGCCAGTGCGGATTGCTGGGAACCATTACGGGGCCTCCCCAGTGCGATCAGTTTCTGGAGTTT TCCGCCGACCTGATCATAGAGCGGCGCGAGGGTTCCGACGTGTGCTACCCAGGAAAATTTGTCAACGAAGAAGCACTACGACAGATCTT GAGAGAGAGCGGAGGGATTGACAAGGAAGCAATGGGATTCACTTATAGCGGCATCAGGACTAATGGTGCGACCTCTGCTTGCAGACGTT CCGGGTCATCCTTCTACGCAGAGATGAAGTGGTTACTAAGCAATACCGATAACGCAGCTTTTCCCCAGATGACCAAGTCCTACAAGAATA CACGGAAAAGCCCCGCTTTGATCGTTTGGGGTATCCATCACTCTGTGAGCACCGCAGAACAGACAAAGTTGTACGGATCGGGAAGTAAG CTTGTCACCGTGGGCTCGTCCAACTACCAACAGTCTTTTGTACCTTCTCCAGGGGCTCGTCCACAAGTCAACGGCCTGTCTGGGAGAATC GACTTTCACTGGCTCATGTTAAATCCCAATGACACAGTCACCTTCTCCTTTAATGGCGCCTTCATTGCACCCGACCGTGCCTCATTCCTGA GAGGTAAGTCCATGGGAATTCAGAGCGGTGTCCAGGTCGACGCTAACTGCGAGGGGGACTGCTATCACTCCGGGGGGACAATTATCAG TAATTTGCCCTTTCAGAATATTGATTCAAGAGCTGTTGGAAAGTGCCCACGCTACGTAAAGCAGAGAAGCCTCCTATTGGCCACAGGAAT GAAGAACGTGCCGGAGATCCCCAAGGGTCGCGGTTTGTTTGGCGCCATCGCTGGCTTTATTGAAAACGGATGGGAGGGTTTGATTGATG GCTGGTATGGGTTCAGGCATCAGAACGCCCAGGGAGAAGGTACCGCAGCAGACTACAAGAGCACACAGTCCGCCATTGACCAGATTACT GGTAAGCTGAACAGATTGATCGAAAAGACTAACCAGCAGTTTGAGCTGATTGACAATGAATTCAACGAAGTGGAAAAGCAGATCGGGAAT GTGATCAATTGGACCCGGGATTCAATCACCGAGGTGTGGAGCTATAATGCCGAGCTGCTCGTAGCCATGGAGAATCAACATACAATAGAT CTCGCTGACTCCGAAATGGATAAGCTCTACGAGCGCGTAAAACGGCAGCTGCGCGAAAACGCTGAAGAAGATGGAACTGGATGCTTCGA GATCTTTCATAAGTGCGACGATGATTGTATGGCCTCCATTAGAAATAATACCTATGACCATACCAAGTACAGGGAAGAGCGCCATGCAAAAT AGAATCCAGATCGATCCCGTCAAACTAAGTTCGGGGTATAAGGATGTGATCCTATGGTTTTCCTTTGGTGCTAGCTGTTTCATTCTGTTAG CCATCGTCATGGGGCTTGTGTTCATATGTGTGAAAAATGGTAACATGCGGTGCACAATTTGTATA
2769	ATGAATACACAGATCTTAGTATTTGCGCTAATCGCTATTATCCCTACCAATGCTGATAAGATCTGTCTGGGCCACCACGCCGTTTCCAATG GCACTAAAGTGAACACTCTCACTGAGCGCGGCGTCGAGGTGGTCAACGCCACCGAGACAGTTGAGAGAACAAACATACCTCGCATCTGT AGTAAAGGCAAGAAGACTGTCGATCTGGGCCAGTGTGGGCTGCTGGGAACGATAACCGGTCCCCCTCAGTGTGATCAGTTTCTGGAATT CTCTGCAGACCTGATTATCGAAAGACGAGAAGGGAGCGACGTGTGTTACCCCGGAAAGTTTGTTAATGAGGAAGCACTGCGCCAGATCC TTCGGGAATCTGGGGGCATCGATAAGGAAGCGATGGGCTTCACTTATAGTGGGATTCGCACCAACGGGGCTACGTCCGCTTGCCGGCG GAGCGGTTCGTCTTTTTACGCGGAAATGAAGTGGCTGCTCAGCAATACCGATAACGCCGCCTTCCCGCAGATGACAAAAAGCTATAAGAA TACCCGGAAAAGTCCGGCACTCATAGTGTGGGGGATCCACCATAGCGTTAGCACGGCGGAGCAGACAAAGCTCTATGGGAGTGGAAAT AAGCTGGTGACCGTCGGCTCTTCAAATTACCAGCAATCATTTGTGCCCTCTCCCGGCGCCCGTCCTCAAGTGAACGGACAGTCAGGAAG GATTGACTTCCACTGGCTCATGTTGAACCCTAATGACACAGTGACATTTTCGTTTAACGGAGCATTCATCGCCCCTGACAGAGCTAGTTTC CTCAGAGGGAAGAGTATGGGGATCCAGAGCGGGGTACAGGTGGATGCAAACTGTGAGGGGGACTGCTATCATTCTGGTGGGACCATCA TTTCGAACCTGCCATTCCAGAACATTGACTCTCGCGCTGTTGGCAAATGTCCTAGGTACGTTAAGCAAAGATCATTGCTGCTCGCAACGG GCATGAAAAACGTGCCGGAAATCCCTAAAGGTCGGGGTCTCTTCGGCGCCATCGCCGGATTCATTGAGAATGGCTGGGAGGGACTCATC GACGGGTGGTATGGGTTCCGGCATCAGAACGCACAGGGCGAGGGGACTGCCGCTGACTACAAATCGACCCAGTCCGCTATTGACCAGA TCACCGGCAAACTCAACCGGCTGATCGAGAAGACTAACCAACAGTTTGAACTCATCGATAATGAGTTCAACGAGGTAGAAAAACAAATTG GAAACGTGATCAATTGGACACGCGATTCTATTACTGAGGTCTGGAGTTATAACGCAGAGCTGCTGGTGGCTATGGAAAACCAGCACACCA TTCATCTAGCTGACTCGGAAATGGATAAGCTGTACGAAAGGGTGAAACGCCAGCTACGCGAAAATGCCCAGGAAGACGGTACAGGATGC TTCGAGATATTCCACAAGTGCGACGACGACTGCATGGCCAGCATCCGCAATAACACATATGACCATTCTAAATACCGGGAGGAAGCCATG CAGAATCGAATCCAGATCGACCCTGTCAAATTAAGCAGTGGGTACAAGGACGTGATCCTGTGGTTTAGCTTCGGAGCTAGTTGTTTTATAT TACTCGCCATCGTTATGGGATTGGTATTTATCTGCGTCAAGAATGGAAACATGCGCTGCACTATCTGCATC

2770	ATGAACACCCAGATACTCGTCTTTGCCCTCATTGCCATAATCCCGACTAATGCGGATAAGATTTGTCTAGGGCATCACGCTGTATCAAACG GCACGAAAGTCAACACCTTGACCGAAAGGGGCGTAGAGGTGGTAAACGCTACAGAAACAGTCGAGCGGACAAATATCCCAAGGATTTGC AGCAAGGGTAAAAAGACTGTAGACTTAGGACAGTGCGGCTTACTCGGCACAATCACAGGCCCCCCCCAATGTGATCAGTTCCTTGAATTC AGTGCAGATCTGATAATCGAAAGGAGGGAGGGATCTGACGTGTGTTACCCAGGCAAATTCGTCAACGAGGAAGCACTCAGACAGATCTT AAGAAAGTCCGGCGGGATAGACAAAGAAGCTATGGGTTTTACGTACAGTGGGATAAGAACCAATGGCGCCACTAGCACCTGTAGCAGGA GTGGGTCGTCCTTCTATGCTGAAATGAAGTGGCTGCTGTCCAATACTGACAATGCAGCCTTTCCTCAGATGACAAAGAGCTACAAGAATA CAAGGAAAAGCCCGGCAATTATAGTATGGGGAATCCACCATTCAGTCAGCACCGCTGAACAAACTAAGCTGTATGGCTCCGGCAATAAG CTCGTCACAGTGGGGAGCTCAAATTATCAGCAGTCATTCGTGCCAAGCCCAGGGGCGCGGCCGCAAGTTAACGGCCTCTCCGGAAGAA TTGATTTTCATTGGTTAATGCTTAATCCTAACGACACGGTTACCTTCAGCTTTAACGGCGCTTTCATCGCTCCCGACAGGGCTTGTTTTCTC CGCGGCAAGTCTATGGGAATCCAGTCCGGGGTGCAGGTCGACGCGGATTGTGAAGGAGACTGCTACCACAGTGGGGGCACTATAATCT CAAATCTGCCATTCCAGAATATCGACAGTCGCGCCGTGGGGAAGTGTCCGAGATACGTTAAGCAACGATCACTGCTACTGGCCACCGGT ATGAAAAATGTGCCAGAGATTCCCAAAGGGAGAGGCCTGTTTGGTGCCATCGCCGGGTTTATCGAGAACGGGTGGGAAGGCTTGATTGA CGGCTGGTACGGATTTCGGCACCAAAACGCCCAAGGGGAAGGTACCGCTGCCGATTACAAGAGCACCCAAAGTGCCATTGACCAGATTA CGGGGAAACTGAACAGACTGATCGAGAAAACGAATCAACAATTCGAACTGATAGATAACGAATTTAACGAAGTCGAGCGACAGATCGGGA ATGTGATCAACTGGACTAGGGATTCTATTACCGAGGTATGGAGCTATAATGCTGAGCTTCTTGTGGCAATGGAAAATCAACACACAATAGA TTTAGCGGATAGTGAGATGGACAAGCTCTACGAACGGGTAAAAAGACAGTTGAGAGAGAATGCCGAGGAGGACGGGACAGGCTGTTTTG AAATCTTCCATAAGTGTGACGATGATTGTATGGCGAGCATCCGAAACAATACTTACGATCACAGCAAATACCGTGAGGAGGCGATGCAGA ATAGAATCCAGATAGATCCGGTTAAACTGTCCAGCGGGTACAAGGATGTGATCCTGTGGTTCTCATTCGGCGCTTCATGTTTCATTTTGCT GGCAATTGTGATGGGACTCGTGTTCATATGTGTGAAAAACGGCAATATGCGCTGCACTATTTGCATC
2771	ATGAATACACAGATCCTAGTGTTCGCACTCATAGCTATAATCCCTACAAATGCCGACAAGATATGTCTTGGCCACCACGCTGTGTCAAACG GGACAAAAGTTAACACCTTGACAGAGCGGGGCGTCGAGGTCGTTAACGCCACCGAGACTGTGGAAAGAACAAATATACCCCGGATTTGT TCAAAAGGGAAGAGAACTGTCGACCTGGGGCAGTGCGGACTCCTGGGGACTATCACTGGACCTCCACAATGCGACCAGTTCCTCGAGTT CTCTGCCGATCTCATCATTGAGAGGCGAGACGGGTCCGATGTGTGTTACCCAGGGAAATTTGTGAATGAGGAAGCCCTTCGGCAGATAC TTCGTGAAAGCGGGGGGCATCGATAAAGAGGCCATGGGTTTTACGTATTCTGGAATTCGCACCAACGGGGCCACAAGTGCATGTCGGAGG AGCGGATCATCATTCTATGCAGAGATGAAATGGCTGCTTTCTAACACCGATAATGCCGCTTTCCCGCAGATGACAAAAAGTTATAAAAATA CCAGGAAATCCCCCGCTCTAATTGTGTGGGGGATTCATCACAGCGTTTCCACTGCCGAACAGACTAAGCTGTACGGTTCGGGCAGCAAG CTGGTGACAGTGGGTTCATCAAACTACCAACAGTCCTTCGTGCCTTCTCCAGGCGCGCGGCCGCAGGTTAACGGACTCTCTGGCAGAAT CGACTTTCATTGGTTGATGCTGAATCCCAACGACACCGTCACCTTCTCCTTTAACGGTGCATTTATCGCCCCAGATCGTGCTTCATTTCTA CGCGGCAAAAGCATGGGGATTCAGTCCGGCGTCCAGGTAGATGCAAACTGTGAGGGCGACTGTTACCACTCAGGCGGAACAATTATTAG CAACCTGCCATTTCAGAATATTGACTCTCGCGCCGTGGGGAAATGCCCAAGATATGTCAAGCAGCGGAGCCTGCTCCTGGCCACTGGGA TGAAGAATGTGCCAGAAATACCAAAGGGTAGAGGTCTGTTCGGCGCCATCGCTGGCTTTATAGAGAATGGATGGGAGGGACTGATCGAT GGCTGGTACGGATTCAGACATCAAAACGCCCAAGGTGAAGGAACCGCTGCCGATTACAAGTCTACCCAAAGTGCTATAGATCAGATAAC CGGCAAGCTTAATAGGCTCATTGAGAAAACAAATCAGCAGTTCGAGCTGATCGACAACGAGTTCAACGAGGTGGAGAAGCAGATTGGAA ACGTGATTAATTGGACGCGCGATAGCATCACAGAGGTGTGGTCCTATAACGCAGAGCTGCTGGTGGCTATGGAGAACCAGCACACTATC GATCTGGCCGACTCCGAAATGGACAAGCTGTACGAGAGGGTGAAGCGGCAGTTGCGAGAGAATGCCGAGGAGGACGGGACAGGCTGC TTTGAGATCTTTCACAAGTGCGACGATGATTGTATGGCTTCAATCCGCAATAACACATACGATCATTCCAAATACAGGGAGGAAGCTATGC AGAACCGAATCCAGATAGATCCAGTGAAGCTCTCTAGTGGCTACAAGGACGTGATTCTGTGGTTCAGTTTCGGTGCTAGCTGTTTTATTCT GCTTGCAATAGTAATGGGGTTGGTGTTTATCTGTGTTAAGTCACGGAACATGCGCTGCACCATATGCATC
2772	ATGAACACACAGATCCTAGTTTTCGCGCTGATCGCAATCATACCTACAAATGCTGACAAGATTTGCCTAGGACATCACGCCGTGTCTAATG GAACCAAAGTAAACACTCTGACTGAGCGCGGGGGTAGAGGTCGTGAATGCAACAGAAACGGTAGAGCGGACAAATATCCCTAGGATATGC AGTAAAGGCAAGCGTACAGTAGATCTCGGCCAGTGTGGCCTACTCGCAACAATCACCGGTCCTCCCCAGTGCGACCAGTTCCTTGAATT CAGTGCTGATCTGATCATTGAAAGGCGGGAGGGGTCCGATGTATGTTACCCTGGGAAGTTTGTCAATGAAGAGGCACTGCGCCAAATCC TGAGAGAGTCAGGAGGCATAGACAAAGAGGCTATGGGATTTACATACAGTGGGATAAGAACTAACGGCGCGACCAGCGCCTGCAGGAG GTCCGGGTCGTCTTTCTATGCCGAAATGAAATGGCTCCTTTCTAATACAGATAACGCAGCCTTTCCCCAGATCACTAAATCATACAAGAAT ACTAGGAAGTCACCCGCACTCATCGTTTGGGGCATACATCACTCGGTCAGCACCGCCGAACAGACTAAATTATACGGTTCAGGGAACAA GCTCGTAACCGTGGGCTCGTCTAACTATCAACAGAGTTTCGTTCCGTCCCCCGGGGCGAGACCGCAGGTGAATGGCCTCTCCGGGCGA ATCGATTTTCATTGGCTGATGCTGAACCCCAACGATACCGTAACCTTTTCGTTCAACGGGGCTTTCATCGCTCCGGATAGAGCATCATTTC TGCGCGGGAAGTCCATGGGCATTCAGAGCGGTGTTCAAGTCGACGCCAACTGTGAGGGAGACTGTTATCACAGCGGGGGGACCATCAT CAGCAACCTCCCGTTTCAGAATATTGACTCAAGGGCGGTCGGAAAGTGTCCACGTTACGTCAAGCAGCGGAGTCTCCTGTTAGCGACAG GCATGAAGAATGTCCCTGAAATTCCTAAAGGCCGGGGCCTATTTGGTGCCATCGCCGGTTTCATCGAGAACGGGTGGGAAGGTCTGATC GATGGTTGGTATGGCTTCCGCCACCAGAATGCGCAAGGGGAAGGAACCGCGGCGGACTACAAAAGTACTCAGTCCGCAATCGATCAGA TCACAGGTAAGTTGAATAGGCTTATCGAGAAGACAAACCAGCAGTTTGAGCTTATAGACAACGAATTCAATGAGGTGGAAAAGCAAATTG GAAATGTAATCAATTGGACCAGGGATAGCATCACCGAAGTGTGGAGCTACAATGCCGAGCTGTTGGTGGCAATGGAGAACCAACATACA ATCCACCTAGCCGATAGTGAGATGGACAAGCTGTATGAGCGGGTGAAGCGGCAGCTGAGGGAAAATGCGGAGGAAGACGGAACTGGGT GTTTCGAGATTTTCCACAAGTGTGATGACGATTGTATGGCCTCAATTCGGAACAACACATACGATCATTCCAAGTATCGAGAAGAGGCCAT GCAAAACCGGATTCAGATTGACCCCGTCAAGTTATCCTCCGGGTACAAGGACGTTATTCTCTGGTTTAGCTTTGGAGCCAGTTGCTTCATT CTGTTGGCAATTGTGATGGGTCTTGTATTCATCTGTGTGAAAAATGGGAATATGCGCTGCACCATCTGTATC

2773	ATGAACACGCAAATCTTGGTGTTCGCACTGATCGCCATTATTCCAACCAATGCTGACAAGATTTGCCTTGGTCACCACGCGGTTAGCAAC GGGACCAAAGTGAACACCCTGACCGAGCGAGGGGTGGAGGTAGTGAACGCAACCGAGACAGTGGAGAGAACTAATATACCCCGGATCT GTAGCAAGGGTAAGAGAACGGTGGATCTTGGGCAGTGTGGCTTGTTGGGGACTATCACAGGCCCTCCTCAGTGCGACCAGTTCTTGGAA TTTTCTGCCGACCTTATCATTGAGCGCAGAGAGGGGTCAGATGTCTGTTATCCTGGAAAATTTGTGAACGAAGAGGCTCTTCGGCAGATA CTGCGCGAATCAGGCGGCATTCACAAAGAAGCCATGGGGTTTACCTATAGCGGAATCACGACTAACGGCGCCACCTCCGCCTGCAGGA GGAGTGGGAGCAGCTTCTACGCTGAAATGAAGTGGTTGCTCTCTAACACAGACAATGCTGCATTCCCTCAGATTACCAAGAGTTATAAGA ACACACGCAAATCTCCCGCGCTGATCGTGTGGGGCATTCATCACAGCGTTAGCACGGCCGAGCAGACAAAACTGTACGGCTCGGGAAAT AAACTTGTGACGGTTGGCAGCAGCAACTATCAGCAGTCCTTCGTTCCTTCTCCAGGCGCGCGCCCCCAGGTGAATGGGTTGTCTGGGAG AATCGACTTTCACTGGCTAATGCTTAATCCTAACGACACGGTGACTTTCTCATTTAACGGAGCTTTCATCGCACCAGACCGCGCATCCTTT CTGAGAGGGAAATCGATGGGCATCCAATCCGGGGTGCAGGTCGATGCGAACTGCGAGGGTGATTGTTATCACTCCGGGGGCACCATAA TCAGCAACCTGCCTTTCCAGAATATCGACTCCCGGGCAGTAGGAAAGTGCCCACGATATGTGAAGCAGAGGAGCTTGCTGCTGGCTACA GGCATGAAGAATGTCCCAGAAATCCCCAAGGGGAGAGGTCTATTTGGGGCCATCGCTGGCTTTATAGAGAACGGGTGGGAAGGTCTGAT TGACGGGTGGTACGGCTTTAGGCACCAGAACGCCCAGGGCGAAGGAACAGCTGCCGACTATAAATCTACACAGAGCGCCATTGATCAG ATAACTGGCAAGTTAAACCGGCTCATCGAGAAGACTAATCAGCAATTTGAGCTTATCGATAACGAATTCAATGAAGTAGAGAAACAAATCG GGAATGTGATCAACTGGACTCGAGATTCAATAACGGAGGTATGGTCCTACAACGCCGAACTGCTCGTTGCTATGGAGAACCAGCACACC ATCGATCTGGCAGACAGCGAGATGGATAAGCTCTACGAGAGAGTCAAGCGACAGTTACGTGAAAATGCCGAAGAAGATGGAACCGGTTG CTTTGAGATATTTCATAAATGCGATGATGATTGCATGGCTTCGATTCGGAATAACACATATGACCACAGCAAGTACAGGGAAGAAGCGATG CAAAACCGCATTCAGATCGACCCAGTCAAACTAAGCTCTGGATACAAAGACGTTATACTGTGGTTCTCCTTTGGAGCCAGCTGCTTCATCC TGCTCGCCATAGTGATGGGACTGGTGTTTATCTGCGTGAAAAACGGAAACATGAGGTGTACCATTTGTATC
2774	ATGAACACACAGATTCTCGTCTTCGCACTCATTGCCATCATCCCTACAAACGCGGACAAAATTTGCCTTGGGCACCACGCCGTGAGCAAC GGCACAAAAGTCAACACATTGACAGAGCGAGGCGTAGAGGTCGTTAATGCCACCGAAACGGTAGAAAGGACGAACATTCCTAGGATATG TAGTAAAGGAAAGCGTACTGTAGATTTAGGACAGTGTGGCTTGCTGGGAACTATCACAGGACCCCCACAATGCGACCAGTTCCTAGAGTT TAGCGCCGATCTGATAATTGAACGTACGCAGGGAAGCGACGTATGCTACCCAGGGAAATTTGTGAAGGAGGAGGCCCTCCGACAGATTC TGAGAGAGTCCGGGGGAATCGACAAGGAAGCCATGGGATTCACCTATTCGGGCATAAGAACCAACGGCGCAACATCCGCTTGCCGGAG GTCCGGGAGTAGTTTTTATGCTGAGATGAAGTGGCTGCTTTCCAACACCGATAACGCCGCCTTTCCTCAGATGACGAAATCCTATAAGAA TACAAGAAAAAGCCCGGCACTCATCGTGTGGGGCATCCATCACAGCGTCTCCACGGCAGAGCAGACTAAACTCTACGGATCTGGAAATA AACTGGTAACAGTGGGGTCAAGTAACTACCAACAGTCGTTCGTGCCATCCCCTGGCGCTCGGCCTCAGGTTAACGGGCTGTCCGGTAGG ATTGATTTCCATTGGCTGATGTTAAACCCAAACGACACCGTGACCTTCAGTTTTAACGGCGCTTTTATTGCTCCCGACAGGGCAAGTTTCT TGAGGGGCAAGAGCATGGGAATCCAGTCTGGTGTGCAGGTCGATGCCAACTGTGAGGGAGATTGTTATCATAGCGGAGGTACGATTATC TCTAATCTACCTTTTCAAAATATCGACTCCAGGGCGGTCGGGAAATGTCCTCGCTACGTCAAGCAGAGGTCACTGCTTCTGGCAACCGGA ATGAAGAACGTGCCCGAAATCCCAAAGGGCAGAGGTCTGTTTGGCGCTATAGCCGGGTTCATTGAGAACGGATGGGAGGGTTTAATAGA CGGGTGGTATGGATTCCGCCACCAAAATGCCCAGGGGGAAGGGACAGCTGCCGATTACAAGTCAACACAGAGCGCGATCGATCAGATC ACAGGAAAGTTGAACAGGTTGATAGAGAAAACAAATCAACAGTTTGAGCTGATCGATAATGAGTTCAATGAGGTCGAGAAGCAAATTGGG AACGTAATAAATTGGACACGAGATTCCATAACTGAAGTGTGGTCTTACAATGCTGAACTGCTCGTGGCGATGGAAAACCAACACACGATT GATTTGGCCGATTCCGAGATGGATAAACTGTATGAACGTGTGAAGCGTCAACTTCGTGAAAATGCCGAAGAGGACGGTACCGGATGCTT CGAGATCTTCCACAACGTGTGATGATCATTGCATGGCCAGTATCAGGAATAACACATATGACCATTCCAAATACCGCGAAGAGGCCATGCA GAATCGTATTCAGATTGACCCCGTTAAGCTGAGTTCCGGTTACAAAGATGTCATTCTCTGGTTTTCTTTTGGCGCCTCATGTTTCATTCTGC TAGCCATTGTGATGGGGCTTGTCTTTATCTGTGTCAAGAACGGTAACATGAGGTGTACCATCTGTATC
2775	ATGAATACCCAGATTCTGGTGTTCGCGTTAATAGCCATTATCCCAACTAACGCTGATAAGATATGTCTGGGCCATCACGCTGTGAGCAATG GAACTAAGGTGAACACTCTGACCGAGAGGGGTGTGGAGGTAGTGAATGCTACCGAGACAGTAGAGCGGACTAACATTCCCAGGATTTGC TCTAAGGGCAAGCGCACCGTCGACCTGGGACAGTGCGGGTTGCTAGGTACGATAACTGGTCCACCACAATGTGATCAATTCTTGGAATT TTCTGCTGACCTCATTATAGAGAGGAGGGAGGGCTCCGATGTGTGCTACCCCGGGAAATTCGTCAACGAAGAAGCACTGCGTCAGATTT TGAGAGAATCGGGGGGTATTGACAAGGAGGCCATGGGTTTTACCTACTCCGGCATTCGTACTAACGGAGCCACCAGTGCCTGCAGGCG GAGCGGGAGTAGCTTCTACGCTGAGATGAAATGGCTTCTAAGCAACACTGATAACGCCGCTTTCCCCCAGATGACTAAATCTTACAAAAA CACACGGAAGAGTCCCGCTCTGATAGTATGGGGAATACACCATTCAGTCTCTACAGCCGAGCAGACTAAGCTTTATGGAAGCGGGAACA AACTGGTGACCGTCGGTTCATCTAATTACCAGCAGTCTTTCGTGCCCTCTCCTGGGGCCAGGCCCCAGGTGAATGGGCTGTCTGGCCGC ATTGATTTCCACTGGTTAATGCTGAATCCCAATGATACCGTGACCTTTTCTTTCAATGGCGCTTTCATTGCTCCTGATAGAGCCTCATTTCT GAGAGGGAAGTCGATGGGGATACAGTCTGGGGTCCAGGTCGACGCTAATTGCGAAGGCGATTGCTACCACTCTGGGGGGACAATTATA AGTAATTTGCCGTTCCAGAATATCGATAGTAGAGCTGTGGGGAAATGCCCCCGCTACGTAAAGCAGCGTTCCCTTTTGCTTGCCACAGGT ATGAAAAACGTGCCCGAAATTCCAAAAGGAAGAGGACTATTTGGTGCCATTGCGGGGTTCATCGAAAACGGATGGGAGGGCCTGATCGA TGGGTGGTACGGCTTCAGACATCAAAATGCTCAGGGGGAAGGAACTGCAGCGGACTATAAATCTACCCAGTCTGCCATCGATCAGATCA CTGGCAAACTAAATCGACTTATCGAGAAAACAAACCAGCAGTTTGAGCTTATCGATAACGAGTTCAACGAGGTAGAAAAGCAGATTGGAA ACGTCATTAACTGGACTCGGGATTCAATCACAGAAGTATGGAGTTACAACGCTGAGCTTTTAGTGGCCATGGAAAATCAGCATACGATTG ATCTCGCCGACTCCGAGATGGATAAGCTGTACGAACGTGTGAAGCGCCAGCTCAGAGAGAATGCAGAGGAGGACGGGACTGGCTGCTT CGAGATCTTTCATAAATGTGACGATGACTGCATGGCAAGCATTCGAAATAACACATACGATCATAGTAAGTATCGGGAGGAGGCTATGCA AAATCGGATTCAAATTGATCCGGTGAAGCTGTCGAGTGGCTACAAGGATGTGATTTTATGGTTTTCATTCGGGGCCTCCTGCTTCATCCTA CTGGCCATAGTGATGGGCCTGGTGTTTATCTGCGTGAAAAATGGTAATATGCGGTGCACCATCTGCATC

2776	ATGAATACTCAAATTTTAGTTTTCGCTCTTATTGCCATCATTCCAACTAACGCTGATAAAATCTGCCTGGGCCACCACGCCGTCAGCAATG GGACCAAGGTAAACACACTCACCGAACGCGGCGTGGAAGTAGTGAATGCGACCGAAACTGTTGAGCGCACCAACATTCCACGAATATGT TCCAAGGGTAAGAAAACCGTGGATCTGGGGCAATGTGGCCTTCTGGGGACGATAACAGGACCACCCCAGTGTGACCAGTTTCTGGAGTT TTCCGCAGACCTGATCATCGAACGTAGGGAGGGAAGCGACGTCTGTTATCCCGGCAAGTTCGTTAATGAGGAGGCCCTTCGACAAATCC TGCGCGAAAGCGGTGGGATAGACAAAGAAGCCATGGGCTTCACATACACGCGCATCAGAACCAACGCCCCCACAAGTGCCTGCAGGCG ATCTGGCTCGTCCTTCTACGCGGAGATGAAGTGGCTGCTGTCAAACACCGACAATGCCGCTTTTCCCCAGATGACAAAAAGCTACAAAAA CACCCGAAAAAGCCCTGCATTAATAGTCTGGGGCATCCACCACAGCGTGTCTACCGCCGAACAAACGAAGCTGTATGGTTCCGGAAACA AACTAGTGACGGTGGGCTCATCAAATTATCAGCAGTCCTTCGTGCCTTCCCCAGGGGCCAGGCCTCAGGTGAATGGACAATCCGGGCGA ATTGATTTTCATTGGCTCATGTTGAACCCCAACGACACTGTAACGTTTTCTTTCAACGGTGCGTTTATCGCTCCAGATAGGGCCTCCTTCC TCCGAGGAAAAAGTATGGGTATCCAGAGTGGCGTGCAGGTGGATGCCAATTGCGAGGGTGACTGCTACCACTCTGGCGGTACAATTATC TCTAATCTGCCCTTCCAGAACATAGACAGCCGTGCGGTCGGGAAGTGCCCCCGCTACGTTAAGCAACGGTCTTTATTGCTTGCCACTGGT ATGAAGAACGTGCCAGAGATACCAAAAGGGAGAGGGCTCTTTGGCGCCATCGCTGGATTTATAGAGAACGGGTGGGAGGGCCTGATTG ACGGGTGGTACGGATTCCGCCATCAGAACGCCCAAGGTGAAGGCACCGCCGCCGACTACAAGAGCACCCAGAGCGCCATTGATCAGAT CACGGGAAAACTGAATAGGCTGATTGAAAAGACAAATCAGCAGTTCGAGCTGATAGACAACGAATTTAACGAAGTAGAAAAACAAATCGG CAACGTGATCAACTGGACAAGAGACTCCATCACTGAAGTGTGGAGCTATAATGCCGAGTTACTCGTGGCAATGGAGAATCAGCATACTAT TGATCTGGCCGATTCGGAAATGGACAAGCTGTACGAACGCGTCAAGCGGCAACTGCGGGAGAATGCCGAGGAGGACGGTACCGGTTGT TTCGAGATTTTTCATAAATGTGATGACGACTGCATGGCCTCAATTAGAAACAACACATATGATCACAGCAAGTATCGGGAAGAAGCTATGC AAAACCGGATTCAGATCGACCCAGTTAAGCTGAGTAGCGGGTATAAGGACGTAATCCTCTGGTTTTCTTTCGGCGCCTCCTGTTTTATTTT ATTAGCGATCGTGATGGGTCTGGTGTTTATCTGCGTCAAAAATGGTAATATGAGGTGTACTATATGTATC
2777	ATGAACACTCAGATTCTAGTATTTGCACTTATTGCCATCATACCAACCAACGCTGACAAGATCTGTCTCGGCCATCATGCGGTCAGTAACG GTACAAAGGTCAATACACTGACAGAACGTGGGGTAGAGGTCGTAAACGCAACCGAGACCGTGGAGAGAACCAATATTCCTCGAATATGT TCCAAAGGCAAGAGGACTGTTGATTTAGGACAGTGCGGGCTGCTGGGAACTATAACAGGGCCCCCACAATGTGATCAATTCCTCGAGTT CTCACCCGACCTCATTATCGAGAGACGCGAGGGATCTGATGTGTGCTACCCTGGCAAATTCGTGAACGAGGAGGCCCTGCGCCAGATCT TGCGGGAAAGCGGTGGAATCGATAAGGAGGCCATGGGGTTTACTTATAGCGGCATAAGGACCAACGGCGCCACCTCTGCATGCAGGAG GAGCGGCAGTAGCTTCTACGCAGAGATGAAATGGCTGCTGTCTAATACCGACAATGCCGCGTTCCCTCAAATGACTAAAAGCTACAAGAA TACGCGCAAGTCTCCAGCCCTGATCGTGTGGGGCATCCATCACAGTGTCTCAACTGCAGAACAGACCAAACTTTACGGTTCAGGCAACAA ACTCGTCACCGTCGGCTCTTCTAATTATCAACAGAGTTTCGTGCCTAGCCCAGGCGAGCGACCCCAGGTGAACGGTCTTTCCGGTCGGA TCGATTTTCACTGGCTCATGCTCAACCCGAACGATACTGTGACTTTCAGTTTCAACGGGGCCTTCATTGCTCCGGATCGGGCATCATTTCT GAGGGGCAAGTCTATGGGCATCCAGTCCGGCGTGCAGGTGGACGCTAACTGCGAAGGTGATTGTTACCACTCGGGTGGCACAATCATTT CTAATCTACCCTTCCAGAACATTGACTCAAGAGCTGTCGGGAAGTGTCCACGATACGTGAAGCAGAGATCCCTGCTCCTCGCCACAGGG ATGAAGAATGTCCCAGAGATTCCTAAAGGGCGCGGCCTGTTCGGAGCTATAGCGGGGTTCATTGAGAATGGGTGGGAAGGCCTAATTGA CGGTTGGTATGGTTTTAGGCATCAGAATGCCCAGGGCGAGGGTACCGCCGCAGATTATAAAAGTACTCAGTCTGCCATCGATCAAATCAC CGGGAAGCTAAACCGGTTGATCGAAAAAACTAACCAGCAGTTTGAGCTGATTGACAACGAATTTAACGAGGTGGAGAAACAGATCGGCAA CGTAATTAATTGGACGCGGGATAGCATCACAGAAGTCTGGAGCTATAATGCAGAGCTCCTGGTGGCGATGGAGAATCAGCACACTATCG ACTTGGCTGATTCGGAAATGGATAAGCTGTATGAGAGGGTGAAACGTCAGCTGCGGGAGAATGCCGAAGAAGATGGTACCGGGTGTTTC GAAATATTCCATAAATGTGACCATGATTGTATGGCCAGCATCCGAAATAACACATATGACCATTCTAAATACAGAGAAGAGGCCATCCAGA ATCGAATACAAATCGACCCTGTGAAACTGTCCTCCGGATATAAAGATGTTATCCTTTGGTTTAGCTTCGGCGCCTCTTGCTTTATACTGTT GGCAATCGTGATGGGCTTGGTATTTATTTGTGTCAAGAATGGCAATATGCGCTGTACGATCTGCATT
2778	ATGAACACCCAGATCTTGGTGTTTGCTCTTATTGCAATCATCCCTACTAACGCCGATAAAATCTGTCTGGGTCACCATGCCGTCAGTAATG GAACCAAGGTAAACACACTGACGGAGAGGGGCGTGGAGGTGGTTAACGCAACCGAAACAGTTGAGCGGACGAATATCCCCCGGATATG TAGTAAGGGAAAGAAAACTGTCGATCTGGCACAGTGCGGCCTACTAGGAACCATCACTGGGCCTCCCCAGTGTGACCAGTTCCTGCAAT TTTCTGCCGACCTGATTATCGAACGCCGTGAGGGATCCGACGTGTGCTATCCAGGTAAATTCGTGAATGAAGAAGCACTTAGGCAGATCC TTCGTGAATCTGGGGGCATCGATAAGGAAGCTATGGGGTTTACTTATTCCGGTATTAGGACCAACGGGGCAACAAGCGCCTGCAGACGA AGTGGTTCGTCATTTTACGCCGAGATGAAATGGCTGTTGTCCAACACTGACAATGCTGCTTTTCCCCAAATGACCAAAAGCTATAAAAACA CTCGGAAATCACCCGCCCTCATCGTATGGGGGATCCACCATTCTGTTTCTACTGCAGAACAGACAAAACTGTACGGGAGCGGCAATAAG CTGGTGACCGTGGGGTCCTCCAATTATCAACAGTCTTTTGTCCCGTCCCCAGGAGCGAGACCACAGGTCAATGGACAGTCAGGCCGAAT CGACTTCCACTGGCTTATGCTTAATCCAAACGACACCGTCACCTTTAGTTTCAATGGCGCCTTCATCGCCCCAGACAGAGCCTCATTCCT GCGCGGGAAATCGATGGGCATTCAGTCGGGAGTCCAAGTGGACGCAAACTGCGAAGGTGACTGTTACCATAGTGGGGGCACAATCATC AGTAACCTGCCCTTTCAGAACATAGATTCCCGCGCCGTAGGAAAGTGTCCCCGGTATGTGAAACAGAGATCATTGTTGCTGGCAACAGG GATGAAAAATGTACCCGAAATTCCCAAAGGGCGAGGGCTGTTTGGAGCCATTGCCGGTTTTATCGAGAACGGGTGGGAGGGCCTGATCG ATGGGTGGTACGGTTTCAGACACCAGAACGCCCAGGGCGAAGGGACCGCTGCAGATTACAAGTCGACCCAATCCGCGATAGATCAGAT CACTGGTAAACTCAATCGACTCATTGAAAAGACAAATCAACAGTTCGAGCTGATTGATAACGAGTTCAACGAAGTCGAAAAGCAGATAGG GAACGTGATCAATTGGACACGCGACTCCATCACCGAAGTGTGGAGTTACAATGCAGAGCTGTTGGTAGCGATGGAGAACCAGCATACTA TTCACCTGGCCGATTCAGAAATGGACAAGCTGTACGAGCGCGTCAAGCGGCAACTGCGAGAGAATGCTGAAGAGGATCGAACCGGATG CTTCGAAATCTTCCACAAGTGCGATGACGACTGCATGGCCTCCATCAGGAACAATACATACGATCACTCAAAGTACCGGGAGGAGGCCAT GCAGAACCGTATTCAGATCGACCCCGTAAAACTGAGCTCCGGCTACAAGGATGTAATCTTGTGGTTCAGCTTCGGCGCTTCATGCTTTAT TTTACTGGCCATCGTCATGGGACTTGTCTTTATCTGTGTTAAGAACGGGAATATGCGGTGCACCATCTGCATC

2779	ATGAATACACAGATACTGGTGTTTGCCCTCATTGCAATCATTCCAACCAATGCCGACAAGATTTGCCTGGGCCATCACGCTGTGTCCAAC GGCACCAAGGTGAACACCCTTACAGAACGCGGGGTGGAGGTGGTCAACGCAACCGAGACCGTGGAGAGAACCAATATTCCAAGGATCT GCTCGAAGGGAAAGAAGACCGTAGATCTGGGACAGTGTGGGCTGCTCGGCACAATAACCGGCCCCCCGCAGTGTGATCAGTTCTTGGA GTTCAGCGCTGATCTGATTATCGAGCGGAGGGAGGGATCCGACGTCTGTTACCCCGGGAAGTTCGTGAATGAAGAAGCTCTTAGACAGA TCCTTCGCGAATCTCGCCGCAATTGACAACGAAGCTATGGGCTTCACTTATTCTGGAATAACAACCAACGGCGCAACCTCCGCCTGTAGG CGCAGCGGATCCAGCTTTTATGCCGAGATGAAATGGTTGTTGTCCAATACTGACAATGCCGCTTTCCCCCAGATGACCAAGTCCTACAAG AATACTCGAAAAAGCCCAGCTTTGATCGTCTGGGGCATCCACCATTCTGTCTCCACCGCGGAGCAGACCAAGCTTTATGGCTCTGGTAAC AAGCTGGTGACTGTGGGAAGCAGCAATTACCAGCAAAGTTTCGTGCCTAGTCCCGGGGCCCGCCCACAGGTAAACGGCCAGTCAGGGA GAATAGACTTCCACTGGCTGATGCTGAACCCTAATGACACGGTGACATTTAGCTTTAATGGTGCTTTCATCGCTCCCGATCGCGCCAGTTT TCTCAGAGGGAAGTCGATGGGAATCCAATCCGGTGTCCAAGTGGACGCCAATTGTGAGGGAGACTGCTATCATAGCGGGGGAACAATCA TATCTAATCTTCCATTTCAGAACATTGATTCTCGAGCAGTTGGAAAGTGCCCCCGGTACGTGAAGCAACGAAGCCTGCTCCTTGCAACCG GCATGAAGAATGTGCCCGAGATCCCAAAGGGACGGGGCCTATTCGGGGCCATCGCCGGGTTCATCGAGAATGGTTGGGAGGGACTAAT CGATGGATGGTATGGCTTTCGGCACCAGAACGCTCAGGGCGAAGGAACCGCAGCAGACTACAAATCAACACAGTCCGCCATAGACCAG ATCACAGGTAAACTCAACCGCCTGATAGAGAAAACAAATCAGCAATTTGAACTTATCGACAATGAGTTCAACGAAGTGGAGAAACAAATAG GAAACGTAATTAACTGGACCAGGGACTCAATCACCGAAGTGTGGAGCTATAATGCCGAGCTCCTGGTGGCTATGGAGAATCAGCATACTA TCGACTTAGCTGACAGCGAGATGGACAAGCTGTATGAACGGGTCAAACGGCAGCTTAGGGAGAATGCTGAAGAGGATGGCACTGGCTG CTTCGAGATCTTCCACAAATGCGATGATGACTGTATGGCCTCTATTAGGAATAATACGTACGACCATTCTAAATACAGGGAGGAGGCTATG CAGAACCGGATTCAGATTGATCCAGTTAAGCTGTCTAGCGGTTATAAAGATGTGATTTTGTGGTTCTCCTTTGGCGCCAGCTGTTTCATCT TGCTAGCAATAGTTATGGGCCTAGTGTTTATCTGCGTGAAGAACGGCAATATGAGATGTACAATCTGCATC
2780	ATGAACACACAGATCCTGGTATTTGCTCTTATCGCTATCATCCCGACCAACGCCGATAAGATCTGTCTTGGCCACCACGCCGTTTCGAAT GGCACCAAGGTGAATACTCTTACCGAACGAGGAGTGGAAGTTGTGAACGCCACGGAAACAGTTGAACGCACCAACATCCCGAGGATCTG TTCCAAAGGCAAGAAAACGGTTGACCTGGGCCAGTGTGGACTGTTGGGGACTATCACCGGGCCCCCACAGTGTGATCAGTTCCTGGAGT TTTCTGCTGACCTGATTATTGAAAGACGGGAGGGGTCAGACGTGTGTTACCCCGGCAAGTTCGTCAATGAGGAAGCTCTTCGACAGATTT TAAGGGAATCGGGCGGAATTGACAAGGAAGCCATGGGGTTCACCTACTCTGGGATTCGTACCAACGGTGCTACCTCCGCTTGTCGAAGG TCTGGCAGTAGCTTCTACGCGGAGATGAAATGGCTCCTGTCAAACACAGACAATGCCGCGTTCCCTCAAATGACTAAGTCATACAAAAAC ACTCGTAAGTCCCCGGCTCTCATAGTGTGGGGCATCCACCATAGTGTGAGCACAGCGGAGCAGACCAAATTGTACGGAAGCGGCAACAA GCTTGTGACTGTGGGTTCCTCCAACTACCAACAGAGTTTCGTGCCAAGCCCGGGCGCTCGCCCTCAGGTCAACGGCCAGTCCGGCAGA ATTGACTTCCACTGGCTCATGCTCAACCCAAACGACACAGTTACATTTAGTTTCAATGGAGCGTTCATAGCACCTGATCGCGCTTCTTTTC TGAGAGGAAAATCCATGGGGATTCAGAGTGGTGTGCAAGTAGATGCTAACTGCGAGGGCGACTGTTACCACTCTGGCGGTACTATCATC TCCAACCTCCCATTTCAAAACATAGATTCTCGTGCGGTGGGCAAATGTCCGAGGTACGTGAAACAGCGATCACTCCTGCTAGCCACCGGA ATGAAAAACGTGCCCGAGATCCCGAAGGGCCGGGGCCTGTTCGGTGCCATCGCAGGGTTTATCGAAAACGGGTGGGAAGGCCTGATCG ATGGGTGGTATGGGTTTCGGCATCAGAACGCACAGGGAGAAGGAACAGCTGCTGACTACAAAAGCACACAGTCTGCCATCGACCAAATT ACAGGTAAACTCAATAGGCTCATTGAGAAGACCAATCAGCAGTTCGAGCTCATTGACAATGAGTTTAACGAGGTCGAAAAGCAAATCGGG AATGTAATAAATTGGACCCGCGACAGTATCACAGAAGTCTGGAGTTACAACGCAGAGCTGCTCGTGGCTATGGAGAACCAGCACACAATT GACTTGGCTGACTCCGAAATGGATAAGTTGTACGAACGAGTAAAGCGCCAGCTGCGCGAGAACGCTGAAGAGGACGGGACGGGGTGTT TCGAAATTTTCCACAAATGTGACGATGATTGCATGGCCTCTATCCGCAACAATACCTACGACCACAGCAAGTACCGCGAGGAGGCCATGC AGAACCGCATCCAAATCGACCCTGTGAAGTTGTCTTCCGGATACAAGGATGTGATCCTGTGGTTCTCCTTTGGGGCCAGTTGCTTTATCC TTCTCGCAATTGTGATGGGACTCGTCTTTATCTGCGTGAAAAACGGTAACATGCGCTGTACAATTTGTATA
2781	ATGAACACCCAGATCCTAGTCTTTGCTCTGATTGCGATCATCCCCACCAATGCGGATAAGATCTGCCTCGGACACCACGCCGTCTCTAAT GGCACGAAAGTCAACACCTTAACAGAAAGGGGGGTGGAGGTGGTTAATGCCACCGAGACAGTCGAACGAACGAATATTCCCAGAATTTG TTCTAAAGGAAAACGCACCGTGGACTTAGGGCAGTGCGGCCTCCTCGGCACTATTACCGGCCCTCCACAGTGTGACCAGTTCTTAGAGT TCAGCGCCGACCTGATCATAGAGCGAAGGGAAGGGTCTGACGTGTGCTATCCTGGCAAATTTGTGAATGAGGAGGCCTTGAGACAGATT CTTAGGGAGTCGGGCGGTATAGACAAGGAAGCTATGGGCTTTACATACAGTGGGATCCGGACAAATGGCGCGACAAGCGCATGTAGGA GAAGCGGAAGCAGCTTTTACGCTGAGATGAAGTGGCTACTCTCCAATACCGATAATGCCGCCTTCCCACAAATGACAAAATCCTATAAGA ACACCAGGAAGTCCCCAGCCCTCATCGTCTGGGGAATTCACCACTCCGTCTCTACAGCAGAACAAACTAAACTCTACGGTTCCGGAAATA AGCTCGTTACTGTCGGCAGCTCCAATTACCAGCAGTCATTCGTCCCAAGTCCCGGAGCAAGACCTCAGGTTAATGGCCTCAGCGGAAGA ATCGACTTCCACTGGTTGATGTTAAACCCGAACGACACAGTGACATTCTCCTTTAACGGAGCCTTCATCGCACCGGATCGGGCCTCTTTTT TAAGGGGAAAGAGCATGGGCATTCAGTCCGGGGTGCAGGTCGATGCTAACTGTGAAGGCGATTGTTATCATAGCGGAGGCACCATCATC TCAAATCTGCCTTTTCAGAACATTGACTCCAGAGCTGTAGGAAAGTGTCCTCGCTATGTGAAACAAAGGTCTCTTCTGCTGGCAACTGGAA TGAAGAATGTGCCAGAGATTCCTAAGGGTCGGGGACTATTTGGAGCCATAGCCGGCTTTATAGAAAATGGATGGGAGGGCCTGATAGAC GGTTGGTACGGGTTCCGCCATCAGAATGCCCAAGGTGAGGGTACGGCCGCAGACTATAAGTCTACACAGAGCGCCATTGATCAGATAAC CGGAAAGCTGAACCGTCTGATAGAGAAAACCAACCAGCAATTTGAACTTATTGATAACGAGTTCAACGAGGTTGAGAAGCAGATCGGCAA TGTTATTAACTGGACAAGAGATTCGATCACCGAAGTTTGGTCATACAACGCCGAACTGCTGGTAGCCATGGAGAACCAGCACACAATCGA CCTCGCAGATAGTGAGATGGACAAACTGTACGAGCGTGTGAAGAGACAGCTGAGAGAGAACGCGGAGCAAGACGGCACTGGCTGTTTC GAGATTTTCCATAAGTGCGATGACGATTGCATGGCCTCCATTAGGAATAATACCTATGACCATTCCAAATACCGTGAAGAAGCCATGCAAA ACCGGATCCAGATTGACCCCGTCAAGCTTTCGTCAGGGTATAAAGATGTGATCCTGTGGTTCTCTTTCGGCGCCAGTTGCTTCATCCTGC TCGCAATCGTGATGGGACTCGTGTTCATATGTGTCAAAAACGGCAACATGCGCTGTACGATCTGCATA

2782	ATGAACACTCAGATCCTTGTCTTTGCCCTAATAGCAATAATTCCAACAAATGCCGATAAAATCTGTTTGGGTCATCACGCCGTCTCAAACG GAACAAAGGTGAACACACTGACAGAACGGGGTGTCGAGGTGGTCAATGCCACCGAAACTGTCGAACGCACTAATATTCCCAGAATATGC TCGAAGGGTAAGAGGACGGTAGACTTGGGCCAGTGTGGCCTGCTTGGTACCATCACTGGCCCTCCCCAGTGTGACCAATTCCTGGAGTT TAGCGCGGACCTGATAATTGAGAGACGTGAGGGTAGCGATGTTTGCTATCCTGGAAAGTTCGTCAATGAAGAGGCCCTGAGGCAGATCC TACCAGAATCCGGCGGGATCGATAAAGAAGCTATGGGATTTACCTACTCCGGGATCAGGACAAACGGGGCAACTTCGGCATGTCGGCG GTCTGGCAGCTCATTTTATGCTGAGATGAAATGGTTGCTCTCAAACACCGATAACGCCGCTTTCCCACAAATCACCAAGTCCTACAAGAAC ACAAGGAAGTCCCCCGCTTTGATTGTCTGGGGCATCCACCACAGCGTGTCAACTGCTGAGCAGACCAAGTTATACGGCTCAGGCAACAA ACTCGTTACCGTGGGATCTTCGAATTATCAACAGAGTTTTGTGCCATCCCCTGGTGCTAGGCCGCAAGTTAATGGTCTTAGTGGGCGTAT AGACTTCCACTGGCTTATGCTGAATCCGAACGATACCGTCACCTTTTCATTCAACGGTGCTTTCATAGCCCCCGATCGCGCCTCTTTCCTC AGGGGGAAGTCTATGGGTATTCAGTCAGGTGTGCAGGTGGATGCGAATTGTGAGGGAGATTGTTACCACTCCGGGGGCACCATAATAAG CAATCTTCCCTTTCAAAATATCGACTCTCGGGCGGTCGGAAAATGTCCGCGCTATGTGAAGCAGCGGAGTCTCCTGCTGGCAACAGGGA TGAAAAATGTGCCCGAGATTCCTAAAGGAAGAGGTTTGTTTGGTGCGATCGCTGGTTTTATCGAGAATGGGTGGGAGGGCTTAATCGAC GGTTGGTATGGGTTTAGACACCAGAACGCCCAGGGAGAGGGTACCGCAGCCGACTATAAGAGTACACAGAGCGCAATCGACCAGATCA CAGGCAAACTCAACCGCCTAATTGAGAAAACTAACCAGCAATTCGAGTTAATCGACAATGAGTTCAACGAGGTAGAAAAGCAAATAGGCA ACGTCATTAATTGGACCCGGGACAGTATAACAGAGGTCTGGAGCTATAATGCCGAGCTGTTGGTGGCCATGGAAAACCAGCACACGATC GACCTGGCAGACAGCGAAATGGACAAGCTGTACGAGAGAGTTAAGCGCCAGTTGCGCGAGAACGCCGAGGAAGATGGCACCGGGTGC TTCGAAATCTTTCACAAGTGCGATGACGACTGCATGGCTTCTATTCGAAATAATACTTATGATCATAGCAAGTATAGAGAAGAAGCCATGC AGAATAGGATCCAGATCGACCCAGTAAAGTTGTCTTCTGGCTACAAGGATGTGATACTGTGGTTCTCCTTTGGGGCTAGTTGCTTTATTTT GCTGGCTATTGTTATGGGTCTGGTGTTCATTTGTGTGAAGAATGGGAACATGCGCTGCACGATATGTATC
2783	ATGAATACCCAGATCCTGGTGTTCGCACTGATCGCCATTATACCCACCAACGCCGACAAGATCTGCCTTGGCCACCACGCTGTTTCTAAT GGAACGAAGGTCAATACTCTGACCGAGAGGGGGGTGGAAGTCGTGAACGCTACAGAGACTGTTGAAAGGACAAATATTCCGCGCATTTG CAGCAAAGGGAAGCGAACCGTGGATTTGGGCCAATGTGGCCTCCTTGGCACTATTACAGGCCCACCTCAGTGTGATCAGTTTCTGGAAT TCTCCGCTGACCTCATTATAGAACGGAGAGAAGGATCAGACGTTTGTTACCCTGGCAAATTCGTGAATGAGGAGGCCTTGAGACAGATAC TCAGGGAGTCCGGCGGCATCGACAAGGAAGCAATGGGTTTCACCTACAGCGGCATTCGCACCAACGGCGCCACCTCCGCCTGCCGCCG CAGCGGGTCTAGCTTTTATGCCGAAATGAAATGGCTCTTAAGTAATACAGACAACGCTGCCTTTCCCCAGATGACAAAATCATACAAGAAC ACAAGAAAATCACCCGCACTTATAGTTTGGGGGATCCATCACTCCGTGTCCACCGCTGAGCAGACTAAGTTGTACGGGAGTGGAAACAAA CTGGTTACGGTGGGTAGTTCTAACTATCAGCAGTCCTTTGTGCCTTCGCCCGGAGCTCGCCCACAGGTTAACGGTCTGAGCGGTCGCAT CGACTTCCACTGGCTAATGCTGAACCCTAACGATACCGTGACATTCTCTTTTAATGGAGCTTTCATCGCCCCCGACCGCGCTTCCTTTTTG AGGGGGAAGTCCATGGGTATCCAGAGCGGAGTGCAGGTGGACGCCAATTGCGAGGGAGATTGTTATCATTCTGGCGGCACCATTATCTC TAATTTACCGTTTCAGAATATCGACTCAAGAGCAGTGGGCAAGTGCCCTCGATACGTGAAGCAGCGCAGCCTGCTTCTTGCTACAGGAAT GAAAAACGTCCCGGAGATTCCCAAGGGGCGGGGCCTGTTCGGCGCCATCGCAGGCTTCATAGAGAACGGTTGGGAGGGACTGATTGAT GGATGGTACGGGTTCCGCCATCAGAACGCCCAGGGCGAAGGAACCGCTGCGGACTACAAGTCCACCCAGTCCGCAATCGACCAGATAA CGGGCAAGTTGAATAGGCTCATCGAAAAAACGAACCAACAATTCGAGCTTATTGATAACGAGTTTAACGAAGTAGAAAAGCAGATTGGGA ATGTGATAAACTGGACCAGAGATAGCATCACCGAGGTGTGGAGCTATAATGCAGAACTGCTAGTAGCTATGGAAAATCAGCACACAATCG ACTTAGCCGATAGTGAGATGGACAAATTGTATGAGCGAGTAAAAAGACAATTAAGAGAAAACGCAGAGGAGGATGGCACCGGTTGTTTCG AAATCTTTCACAAATGTGATGACGACTGTATGCCCTCAATTCGGAACAACACCTACGACCACAGCAAATACAGAGAAGAGGCCATGCAGA ATCGCATACAGATAGATCCAGTCAAACTGTCATCTGGGTACAAGGACGTTATATTATGGTTCTCTTTCGGGGCAAGCTGCTTCATCCTTCT GGCTATCGTGATGGGCTTGGTCTTCATCTGCGTCAAGAATGGAAACATGCGCTGCACCATCTGCATC
2784	ATGAATACCCAGATCCTGGTTTTCGCCCTGATCGCTATCATTCCTACTAACGCAGATAAGATCTGCCTGGGGCACCATGCAGTCAGCAAC GGGACCAAGGTGAACACACTGACTGAGCGTGGAGTCGAGGTCGTCAATGCGACTGAGACAGTAGAGCGGACGAACATACCCAGAATCT GCAGCAAAGGGAAACGGACGGTAGACCTCGGCCAGTGCGGTCTCCTGGGTACCATCACAGGTCCACCCCAATGCGATCAGTTTCTGGA GTTCTCCGCCGACCTCATCATCGAACGGCGCGAGGGGTCCGATGTCTGCTACCCTGGTAAATTCGTCAATGAAGAGGCTCTCCGTCAGA TTCTTCGGGAATCAGGGGGGATTGACAAGGAAGCTATGGGATTTACATACTCCGGTATACGAACCAACGGAGCTACCAGTGCTTGCCGA CGGTCAGGGTCATCATTCTACGCCGAGATGAAATGGCTTCTCTCGAACACTGATAATGCTGCGTTTCCACAAATTACCAAAAGCTACAAAA ACACCCGGAAGTCCCCTGCACTGATCGTGTGGGGAATTCACCACAGCGTCTCTACGGCTGAACAGACCAAACTGTACGGATCAGGCAAC AAGTTGGTGACTGTCGGGTCATCCAATTATCAGCAGAGTTTCGTTCCCAGCCCTGGCGCAAGACCTCAGGTGAACGGCTTGTCCGGGAG GATTGATTTTCACTGGCTCATGCTGAATCCTAATGACACTGTTACCTTTTCTTTTAATGGGGCTTTTATTGCACCGGACCGAGCAAGTTTTC TACGCGGAAAGTCCATGGGGATACAGAGTGGGGTGCAGGTGGATGCTAATTGCGAGGGCGATTGTTATCACAGCGGCGGCACCATAAT ATCAAACCTGCCCTTCCAGAACATCGACTCCAGGGCCGTGGGCAAGTGTCCAAGATACGTCAAACAAAGATCTCTGCTGCTGGCCACTG GGATGAAAAACGTCCCAGAAATCCCAAAAGGGAGGGGGCTCTTCGGAGCCATAGCCGGGTTTATCGAAAATGGATGGGAAGGTCTGATA GACGGGTGGTACGGTTTCAGACATCAGAACGCACAGGGCGAGGGAACAGCCGCCGACTACAAGAGCACCCAGAGTGCAATCGATCAGA TCACTGGTAAGCTTAATAGGCTGATCGAAAAGACTAATCAACAATTTGAGCTGATAGACAATGAATTCAACGAAGTTGAAAAGCAGATAGG TAATGTAATCAATTGGACCCGAGATTCAATAACTGAAGTCTGGAGCTACAATGCCGAACTGTTGGTGGCTATGGAGAACCAACATACGATT GACCTGGCCGACTCCGAGATGGACAAGCTGTATGAGCGGGTAAAGAGACAGCTGAGGGAGAATGCCGAGGAAGATGGCACCGGATGTT TCGAGATTTTTCACAAGTGTGATGACGACTGCATGGCGTCTATCCGGAATAACACCTACGACCACTCAAAATATCGGGAAGAAGCGATGC AGAATCGAATTCAGATCGACCCAGTCAAACTGTCGAGTGGCTATAAAGATGTGATACTGTGGTTTAGCTTCGGGGCATCTTGTTTCATTCT GCTGGCTATTGTGATGGGCCTCGTGTTCATTTGCGTTAAGAATGGGAATATGCGATGCACCATTTGTATC

2785	ATGAATACTCAAATACTTGTCTTTGCACTAATCGCTATCATACCCACTAATGCCGATAAAATCTGCTTAGGACACCATGCTGTGTCCAACG GAACTAAAGTGAACACTCTGACGGAGCGAGGCGTCGAAGTTGTCAACGCAACCGAGACTGTGGAGCGCACGAATATCCCCCGAATCTGC TCGAAAGGAAAGAAGACTGTTGACCTTGGTCAATGCGGATTACTGGGAACCATCACAGGACCCCCCCAATGCGACCAGTTTCTGGAATTT AGTGCTGATCTTATAATCGAGCGCCGGGAAGGTAGCGACGTTTGTTATCCGGGAAAGTTTGTAAACGAGGAGGCTCTGAGACAGATCCT GCGCGAGTCAGGGGCGATCGACAAGGAGGCCATGGGCTTTACCTACTCTGGAATCAGAACAAATGCCGCTACTTCCGCTTGTCGGCGC TCTGGAAGCAGCTTCTATGCTGAAATGAAATGGCTGCTGTCCAACACCGACAATGCAGCCTTTCCTCAGATGACTAAATCTTACAAAAATA CCCGGAAATCCCCAGCTCTGATAGTGTGGGGTATCCATCACTCAGTCAGTACTGCTGAACAAACCAAGTTATATGGCTCCGGCAATAAGC TAGTAACCGTGGGAAGCTCCAACTACCAACAGTCTTTCGTGCCCTCACCCGGCGCAAGACCCCAGGTGAACGGCCAGTCAGGTCGCATT GATTTCCACTGGTTGATGCTTAACCCAAATGACACCGTTACCTTTTCCTTCAATGGCGCCTTTATAGCGCCCGACAGAGCTTCATTCCTTA GAGGCAAGTCTATGGGTATCCAATCCGGAGTCCAGGTAGACGCAAACTGTGAGGGAGACTGCTATCATTCCGGTGGAACCATCATATCG AATTTGCCTTTCCAGAACATCGATAGCCGGGCGGTCGGCAAGTGCCCTAGATACGTTAAACAAAGGAGCCTGCTCTTAGCAACAGGTATG AAGAACGTCCCAGAGATTCCAAAGGGCCGAGGACTGTTCGGTGCAATAGCTGGCTTTATCGAGAATGGGTGGGAGGGGCTCATCGATG GTTGGTATGGGTTCAGACACCAGAATGCCCAGGGCGAAGGAACCGCTGCTGACTACAAAAGCACACAGAGCGCTATAGATCAGATAACA GGCAAGCTTAACCGGCTGATCGAGAAAACGAATCAGCAGTTCGAACTCATTGATAATGAGTTCAACGAAGTGGAGAAGCAGATTGGTAAT GTCATCAACTGGACTAGAGATAGTATAACAGAAGTGTGGTCTTATAACGCCGAACTCCTGGTCGCAATGGAAAATCAACATACCATTGAC CTGGCTGATTCGGAGATGGACAAACTTTATGAGCGGGTGAAGAGGCAGCTCCGAGAGAATGCAGAAGAGGACGGCACAGGGTGCTTTG AGATCTTCCATAAGTGCGACGATGATTGCATGGCTTCCATCAGGAATAATACTTATGACCACTCCAAGTACCGGGAGGAAGCCATGCAAA ATCGGATCCAAATACATCCAGTGAAGCTTTCGTCAGGATATAAGGACGTGATCTTGTGGTTTAGCTTCGCGGCCTCTTGTTTCATCCTCCT GGCGATTGTCATGGGACTTGTATTCATTTGTGTGAAGAACGGAAACATGCGCTGCACCATTTGCATA
2786	ATGAACACACAGATCCTTGTGTTCGCCCTTATCGCAATAATTCCCACAAACGCCGACAAAATATGCCTGGGTCACCACGCCGTAAGCAAT GGCACCAAAGTTAACACCCTCACCGAGCGTGGGGTGGAGGTGGTGAACGCCACCGAGACCGTTGAAAGGACAAACATTCCACGAATTT GTAGTAAGGGCAAACGGACTGTAGATCTGGGTCAGTGTGGGCTGCTGGGCACTATCACCGGCCCACCACAGTGCGACCAATTCCTAGAA TTCTCAGCTGATTTAATCATCGAAAGGCGCGAGGGGTCTGACGTCTGTTACCCGGGAAAATTTGTGAACGAAGAGGCCCTGCGACAGAT TCTGCGTGAGTCCGGGGGGAATCGACAAAGAGGCGATGGGATTTACCTATTCAGGCATACGGACCAATGGAGCTACTTCTGCATGCCGGA GGAGCGGCTCGTCTTTCTACGCTGAAATGAAGTGGTTGCTGAGTAATACCGACAACGCAGCCTTCCCCCAGATGACAAAGAGCTACAAG AACACACGCAAGTCACCCGCACTCATAGTCTGGGGGATTCATCACTCAGTCAGTACCGCGGAGCAGACCAAGCTCTACGGTAGCGGCAA CAAGCTGGTCACAGTTGGTTCTAGTAATTACCAGCAGTCCTTTGTTCCCTCCCCCGGCGCCCGTCCCCAAGTCAACGGGCTCTCGGGCA GAATCGACTTCCACTGGCTCATGTTGAACCCCAACGATACTGTGACATTCTCTTTTAATGGTGCTTTTATCGCGCCGGACAGGGCCTCTTT CTTACGCGGCAAGAGCATGGGTATACAGTCTGGAGTCCAGGTTGATGCTAATTGCGAAGGCGACTGTTATCATTCAGGAGGCACCATCA TATCAAACCTGCCCTTCCAAAATATAGATAGTCGGGCTGTGGGTAAATGCCCTAGGTACGTCAAGCAGAGAAGTCTTCTGCTCGCCACTG GCATGAAAAACGTACCAGAGATTCCTAAAGGAAGAGGCCTATTTGGAGCAATTGCCGGCTTTATCGAGAATGGGTGGGAAGGCCTTATAA ACGGGTGGTATGGCTTCAGGCACCAGAACGCCCAGGGGGAGGGTACCGCCGCCGATTACAAGAGCACTCAGAGTGCCATAGACCAGAT TACTGGAAAATTAAATAGACTGATTGAGAAAACAAACCAGCAGTTTGAACTGATTGACAATGAATTCAATGAGGTGGAGAAGCAGATTGGG AACGTGATCAATTGGACCCGGGACTCTATTACTGAGGTATGGAGCTATAACGCTGAGCTTTTGGTGGCCATGGAGAACCAACATACGATT GATCTGGCAGATAGCGAAATGGATAAGCTTTATGAGCGTGTTAAGAGGCAGCTCCGCGAAAATGCCGAAGAGGATGGTACGGGATGCTT CGAGATTTTCCATAAATGCCACGACCATTGCATGGCTAGCATTAGGAACAACACCTATGACCACAGTAAATATAGACAGGAGGCCATGCA AAATCGCATTCAGATAGACCCTGTCAAGTTGAGCTCGGGGTATAAGGACGTCATACTTTGGTTTAGTTTTGGGGCGAGCTGTTTTATTTTG CTGGCCATAGTTATGGGTCTGGTCTTTATTTGTGTAAAGAACGGGAATATGCGTTGCACCATTTGCATA
2787	ATGAACACCCAAATATTAGTGTTCGCCCTCATAGCAATCATTCCCACAAATGCTGATAAGATTTGCCTTGGTCATCATGCTGTGAGTAACG GGACCAAGGTGAACACTCTTACGGAGCGCGGGGTTGAAGTTGTTAACGCTACCGAGACCGTTGAAAGAACCAACATCCCACGAATCTGC AGCAACGGCGAAGCGGACCGTGGACTTGCGACAGTGTGGGTTGTTGGGGACTATCACCGGCGCCCCCTCAGTGCGACCAGTTCCTCGAAT TCAGCGCCGATCTGATCATAGAGCGACGGGAGGGGAGTGACGTTTGCTACCCCGGTAAATTCGTGAAGGAAGAAGCCCTGAGACAGATT CTACGGGAGTCTGGCGGCATAGATAAAGAGGCTATGGGTTTCACATATAGCGGCATCCGCACAAACGGAGCCACTTCTGCCTGCAGGAG ATCTGGCTCCTCTTTTTACGCTGAAATGAAATGGTTACTGTCCAACACCGACAATGCCGCCTTTCCCCAAATGACCAAGTCATACAAGAAT ACCCGGAAATCCCCAGCTTTGATCGTTTGGGGTATCCATCACTCAGTTTCTACCGCGGAGCAGACTAAACTCTATGGCTCTGGAAATAAG CTGGTGACAGTGGGCTCTTCCAATTATCAGCAGTCGTTTGTTCCGTCCCCTGGCGCTAGACCACAGGTGAACGGCCTGTCCGGGCGAAT CGATTTCCACTGGCTGATGCTGAACCCGAACGACACCGTCACATTTAGCTTCAATGGAGCCTTCATTGCTCCCGATCGAGCTTCTTTCCT GCGGGGCAAATCAATGGGCATACAATCGGGTGTGCAGGTGGACGCCAATTGTGAGGGAGACTGTTACCATTCAGGCGGCACCATTATTA GCAATCTACCATTCCAGAATATAGACTCCCGGGCTGTTGGCAAGTGCCCCCGGTATGTTAAACAGCGGTCCCTGCTCCTGGCTACCGGT ATGAAGAATGTTCCGGAAATCCCGAAGGGGCGCGGACTCTTCGGCGCTATCGCGGGTTTTATCGAGAACGGGTGGGAGGGCCTTATCG ATGGGTGGTACGGCTTCCGGCACCAGAATGCCCAGGGAGAAGGCACCGCCGCCGACTATAAGTCAACACAGTCCGCAATTGACCAAAT CACCGGTAAGCTGAATCGCTTAATTGAGAAAACCAACCAACAATTCGAACTGATTGATAACGAATTCAACGAGGTCGAAAAGCAGATCGG GAATGTGATCAACTGGACCCGCGATAGCATTACCGAGGTCTGGTCTTACAACGCAGAACTTCTCGTTGCGATGGAGAATCAACACACTAT TGATCTGGCTGACTCCGAGATGGACAAGCTGTACGAGCGCGTGAAGCGCCAGCTGCGGGAAAATGCAGAAGAGGATGGCACTGGCTGC TTCGAGATCTTCCACAAGTGCGATGATGATTGCATGGCATCGATCAGGAACAATACATACGACCACTCGAAGTACAGAGAGGAGGCTATG CAGAACCGAATACAGATAGATCCAGTTAAACTATCATCCGGGTACAAGGACGTTATCCTGTGGTTCTCATTTGGAGCATCCTGCTTCATTC TTCTCGCAATCGTCATGGGACTGGTTTTCATCTGTGTGAAGAACGGCAACATGCGGTGCACCATCTGCATC

2788	ATGAATACCCAGATATTAGTATTTGCCCTTATAGCCATCATCCCTACTAACGCGGACAAAATCTGCCTGGGACACCATGCCGTTTCTAACG GTACAAAAGTCAACACCTTAACGGAACGGGGGGTGGAAGTGGTTAATGCTACAGAGACCGTTGAACGCACCAACATCCCCCGCATTTGC TCCAAAGGGAAGAAAACTGTGGACCTAGGTCAGTGTGGCCTGTTGGGTACTATCACCGGCCCACCTCAATGCGACCAGTTTCTGGAGTT CAGCGCCGATCTGATCATCGAAAGAAGAGAGGGTAGCGACGTTTGTTATCCTGGAAAGTTTGTTAATGAAGAGGCTCTGAGACAGATTTT GCGAGAATCCGGGGGAATCGACAAGCAAGCCATGGGCTTTACATACTCTGGTATTAGGACCAACGGTGCCACTTCTGCCTGCAGACGGA GCGGTAGTTCTTTTTACGCCGAAATGAAATGGCTATTGTCTAATACGGATAACGCTGCATTTCCTCAGATGACGAAATCATACAAAAATAC AAGGAAAAGCCCCGCTCTGATTGTGTGGGGTATCCACCACTCCGTATCCACAGCAGAACAAACAAAGCTGTATGGATCTGGTAACAAGCT CGTGACGGTGGGCTCTAGCAACTATCAGCAGAGTTTTGTGCCATCACCTGGCGCAAGACCCCAGGTCAACGGCCAGTCCGGCCGAATC GACTTCCACTGGCTTATGCTAAACCCCAACGACACGGTGACTTTCAGTTTTAACGGCGCCTTTATCGCTCCTGATCGCGCTTCTTTCCTTA GGGGTAAATCTATGGGGATACAATCAGGAGTGCAGGTAGACGCTAACTGCGAAGGAGACTGCTATCACTCTGGAGGAACGATAATCTCC AATTTACCCTTCCAGAACATAGACAGTCGGGCCGTCGGGAAATGCCCCAGGTACGTGAAGCAGCGGTCTCTTCTGCTCGCGACTGGCAT GAAAAACGTTCCCGAGATACCCAAGGGAAGGGGCCTGTTCGGCGCAATTGCTGGGTTCATTGAAAACGGCTGGGAGGGCCTTATTGATG GTTGGTACGGATTCAGACACCAGAATGCACAGGGCGAGGGCACAGCCGCCGATTATAAGTCAACGCAGTCAGCGATAGACCAGATAACC GGGAAATTGAACCGTCTGATTGAGAAAACAAACCAGCAATTCGAATTGATTGATAACGAGTTTAATGAGGTGGAAAAGCAGATTGGAAAC GTCATTAACTGGACACGAGATTCAATTACAGAGGTGTGGAGCTATAACGCGGAGTTGCTGGTCGCTATGGAAAACCAACATACTATAGAT CTTGCCGATTCCGAGATGGACAAACTCTATGAGAGAGTCAAACGACAATTAAGGGAAAACGCCGAGGAAGATGGCACCGGATGCTTCGA AATCTTCCATAAGTGCGATGACGACTGTATGGCATCTATTCGCAACAACACGTACGATCACTCGAAATATCGGGAAGAGGCCATGCAGAA TAGGATCCAGATTGACCCTGTCAAATTGAGCAGCGGCTACAAGGATGTTATTCTGTGGTTCAGTTTCGGCGCCTCCTGTTTCATCTTACTC GCTATTGTCATGGGCTTAGTGTTCATCTGTGTCAAGAATGGCAATATGCGGTGCACTATATGCATT
2789	ATGAACACTCAGATTCTGGTGTTCGCCCTAATTGCCATTATTCCCACCAATGCCGACAAAATTTGCCTGGGCCATCATGCCGTCTCGAAC GGTACCAAAGTCAATACCCTGACAGAGCGCGGCGTGGAGGTGGTGAATGCAACTGAGACCGTCGAGAGGACTAATATCCCCAGAATTTG CTCAAAGGGAAAGAAAACAGTGGATCTGGGTCAGTGCGGTCTCTTAGGGACTATCACTGGTCCTCCTCAGTGCGACCAGTTTCTGGAATT TAGTGCTGATCTGATTATCGAGCGGCGCGAAGGGTCCGACGTTTGCTACCCCGGAAAGTTTGTTAACGAGGAAGCCTTGCGGCAAATCC TGCGGAAGTCCGGCGGCATCGACAAGGAGGCCATGGGCTTTACATACTCTGGCATCCGCACCAACGGAGCTACCAGCACCTGTAGGAG GTCTGGTAGTTCATTCTATGCTGAGATGAAGTGGCTGCTCAGTAATACCGACAATGCTGCTTTTCCTCAGATGACAAAATCATACAAGAAC ACGAGAAAAAGCCCAGCTATTATTGTATGGGGGATTCATCATTCTGTCAGTACCGCTGAGCAGACTAAGCTTTACGGTAGCGGAAATAAG CTTGTCACAGTGGGGTCCAGCAATTACCAGCAGTCGTTTGTTCCCTCCCCAGGAGCACGCCCCCAAGTCAACGGTCTATCGGGGACGAT CGACTTCCATTGGCTGATGCTCAATCCTAACGACACCGTGACATTCTCTTTCAATGGCGCCTTTATCGCACCGGACCGTGCCTGTTTTTTA CGCGGTAAGTCAATGGGGATACAATCTGGAGTCCAGGTGGACGCCGACTGTGAAGGGGACTGCTATCATTCGGGTGGTACTATTATCAG CAATCTGCCCTTCCAAAACATAGACTCAAGAGCTGTTGGAAAGTGCCCGAGATACGTGAAGCAAAGGTCACTCTTGCTTGCAACCGGGAT GAAGAACGTGCCTGAGATCCCGAAGGGCCGAGGGCTGTTTGGTGCTATAGCTGGCTTCATCGAGAATGGATGGGAGGGTTTAATTGACG GCTGGTATGGATTTCGGCACCAGAATGCACAGGGAGAAGGGACCGCCGCCGATTACAAAAGTACGCAGAGTGCTATCGATCAGATTACC GGTAAATTGAATAGGCTGATTGAAAAAACTAACCAGCAGTTCGAGCTGATTGATAACGAGTTCAACGAGGTCGAGAGACAGATTGGGAAC GTCATTAATTGGACGAGAGACTCAATCACGGAGGTCTGGAGCTACAATGCTGAGCTGTTAGTAGCAATGGAAAATCAGCACACTATTGAC CTCGCCGACTCCGAGATGGACAAACTGTACGAACGGGTGAAGCGGCAACTTCGCGAAAACGCTGAGGAAGACGGCACGGGATGTTTCG AAATCTTCCATAAATGCGACGATGACTGCATGGCCTCTATCCGTAATAACACATATGACCATAGTAAGTATAGGGAGCGAAGCCATGCAGA ATCGAATCCAGATCGACCCCGTGAAGCTGAGCTCCGGCTACAAAGACGTCATCCTCTGGTTTAGCTTTGGGGCGTCTTGTTTCATTCTCC TCGCGATCGTGATGGGTCTTGTCTTCATTTGCGTGAAAAACGGGAATATGAGGTGTACCATTTGTATC
2790	ATGAATACCCAAATACTTGTGTTTGCCTTGATTGCCATAATTCCTACCAATGCCGACAAAATCTGCCTGGGCCATCATGCCGTTAGTAATG GGACTAAGGTGAACACCCTGACAGAACGTGGCGTGGAGGTGGTCAACGCAACAGAAACCGTCGAGAGGACCAACATTCCTCGGATATG TTCGAAGGGAAAGAAGACGGTGGACCTGGGGCAATGCGGGCTGCTCGGCACCATTACGGGTCCACCTCAGTGTGATCAGTTTCTCGAG TTTTCTGCAGACCTCATCATTGAACGTAGAGAGGGCTCAGACGTGTGTTACCCAGGCAAATTTGTTAACGAGGAAGCCCTGCGCCAGATC CTGCGAGAATCTGGCGGCATTGACAAAGAGGCTATGGGATTCACGTACTCTGGTATCCGTACCAACGGCGCCACGTCCGCCTGTAGAAG GTCAGGCTCATCTTTTTACGCCGAGATGAAATGGCTCCTGTCTAATACGGACAATGCCGCATTTCCTCAGATGACCAAAAGTTACAAAAAC ACCCGGAAGAGTCCAGCTCTGATTGTGTGGGGCATCCATCACAGCGTGTCAACTGCGGAACAGACCAAGCTGTATGGGTCCGGCAATAA ACTGGTGACTGTGGGTTCCAGCAATTACCAGCAATCCTTTGTGCCCAGCCCTGGCGCTAGACCGCAAGTGAACGGGCAGAGCGGGAGG ATTGACTTCCACTGGCTGATGCTCAATCCCAACGATACCGTAACCTTTTCATTTAACGGCGCTTTCATCGCCCCCGATAGAGCTAGTTTCC TACGGGGAAAAAGCATGGGTATTCAAAGTGGTGTGCAGGTAGACGCAAATTGTGAAGGCGACTGTTACCATTCGGGTGGGACCATTATC TCCAACCTGCCCTTCCAGAACATTGATTCACGCGCCGTGGGAAAGTGCCCTAGGTATGTTAAACAGAGAAGCCTGCTGTTGGCCACAGG GATGAAGAATGTGCCAGAGATTCCAAAGGGACGGGGACTCTTTGGTGCGATCGCAGGCTTCATCGAGAACGGTTGGGAAGGGCTGATA GATGGATGGTACGGTTTCCGTCACCAGAACGCCCAAGGAGAGGGGACAGCCGCGGACTACAAATCAACGCAGTCTGCCATTGACCAAAT CACTGGCAAACTGAACCGGCTTATCGAGAAAACCAACCAGCAGTTTGAACTGATAGATAATGAATTCAACGAGGTTGAGAAGCAAATTGG GAACGTCATAAATTGGACTAGAGACTCTATCACAGAGGTGTGGTCATATAATGCAGAGCTGTTGGTGGCAATGGAAAACCAGCACACTAT CGACCTAGCAGATTCTGAGATGGACAAGCTATATGAACGCGTTAAGCGGCAGCTGCGGGAAAACGCAGAAGAAGACGGTACAGGTTGTT TCGAGATATTTCACAAATGCGACGATGATTGTATGGCATCGATCAGAAATAATACGTATGACCACTCCAAATACCGAGAAGAGGCTATGCA GAACCGAATCCAGATTGATCCTGTAAAGCTCAGTTCTGGCTACAAGGACGTCATTCTCTGGTTTAGCTTTGGAGCCTCCTGTTTCATTCTT TTGGCCATCGTTATGGGCCTCGTGTTTATTTGCGTGAAAAATGGTAATATGCGGTGTACCATATGTATC

2791	ATGAATACCCAAATTCTGGTTTTTGCGCTGATCGCGATCATACCGACCAACGCGGATAAAATTTGTCTCGGGCATCACGCCGTGAGCAAT GGGACCAAGGTAAACACACTGACGGAGCGGGGGGTGGAGGTGGTGAACGCCACAGAGACGGTCGAGCGGACAAATATTCCTCGGATAT GCTCCAAGGGCAAGAAGACTGTTGATTTAGGCCAGTGCGGCCTCCTCGGCACCATCACCGGACCGCCCCAGTGCGACCAGTTCCTGGA GTTTTCCGCTGACCTGATCATCGAAAGGAGAGAGGGAAGCGATGTGTGTTACCCTGGAAAGTTTGTTAATGAGGAGGCCCTAAGACAGA TTCTGCGGGAATCCGGCGGCATCGAGAAGGAGGCCATGGGATTTACTTACAGCGGGATTAGAGCGAACGGAGCTACATCTGCCTGTCG CAGGAGTGGATCTTCCTTCTACGCTGAGATGAAATGGCTGCTCTCGAATACGGACAACGCCGCGTTTCCTCAAATGACTAAATCTTACAA AAACACTCGGAAGTCCCCCGCGCTCATTGTGTGGGGAATCCACCACAGTGTGAGTACAGCCGAGCAGACTAAACTGTATGGCAGTGGGA ATAAGCTGGTCACAGTTGGGTCCTCCAACTACCAGCAGAGCTTCGTTCCCAGTCCGGGCGCACGCCCGCAGGTTAATGGCCTTAGCGGT AGAATCGATTTCCACTGGCTGATGCTGAATCCTAACGATACTGTGACTTTCAGCTTCAATGGCGCTTTTATTGCACCCGACAGAGCTTCTT TTCTGCGTGGCAAATCCATGGGCATTCAGTCCGGCGTGCAGGTGGACGCCAACTGCGAAGGCGATTGTTACCACTCTGGGGGCACCAT CATCTCCAACCTCCCTTTTCAAAACATAGACAGTCGCGCTGTGGGAAAATGTCCAAGGTACGTTAAACAACGGTCTCTGCTACTGGCTAC CGGCATGAAGAATGTGCCCGAAATCCCCAAGGGACGGGGCTTATTTGGCGCGATTGCCGGATTTATTGAAAATGGCTGGGAGGGCCTCA TTGACGGATGGTACGGTTTTCGACACCAGAACGCTCAGGGTGAAGGGACCGCTGCAGACTACAAATCCACGCAGTCCGCTATCGATCAG ATTACAGGAAAGCTGAACAGACTAATCGAGAAGACTAATCAGCAGTTCGAGCTAATCGATAATGAGTTCAACGAGGTCGAAAAGCAGATT GGGAACGTCATCAACTGGACTCGCGATAGCATCACGGAGGTATGGTCTTATAATGCAGAGTTGCTGGTGGCCATGGAAAATCAGCATAC AATCGACTTGGCTGACTCTGAGATGGATAAATTGTATGAGAGGGTAAAAAGGCAGTTGAGAGAGAACGCGGAGGAAGATGGCACTGGTT GCTTCGAGATCTTTCACAAGTGTGACGACGATTGCATGGCCAGTATTAGAAATAATACATATGACCACAGCAAGTACAGGGAGGAGGCCA TGCAGAATCGTATCCAAATCGATCCCGTGAAATTGTCAAGCGGATATAAAGACGTTATACTATGGTTCTCCTTTGCGCCTAGTTGTTTCAT TCTCCTGGCGATCGTCATGGGCCTGGTTTTCATCTGCGTTAAGAATGGGAATATGAGATGTACTATCTGCATC
2792	ATGAATACCCAGATTCTGGTGTTCGCTCTCATTGCAATAATCCCGACTAACGCTGATAAAATCTGTCTTGGCCACCACGCTGTGAGTAATG GAACGAAAGTCAATACGCTCACTGAGCGGGGAGTCGAAGTTGTTAACGCTACCGAGACCGTGGAGCGGACAAACATACCGCGGATTTGC TCAAAGGGCAAAAAGACAGTGGATCTGGGTCAGTGCGGCCTTCTCGGCACAATCACCGGGCCCCCACAGTGTGACCAATTTCTCGAGTT TTCAGCTGATTTGATAATTGAGCGCCGGGAGGGCAGTGATGTATGCTATCCAGCGAAGTTCGTGAATGAAGAGGCCCTCCGCCAAATTC TCCGGGAATCTGGCGGGATTGACAAGGAAGCCATGGGGTTCACTTACTCTGGAATTCGGACTAATGGTGCTACCAGTGCTTGCCGTAGA AGTGGTAGCAGTTTCTATGCCGAAATGAAATGGTTGTTAAGCAATACCGATAACGCAGCATTTCCCCAGATGACAAAGTCGTACAAAAACA CCAGGAAAAGTCCTGCTCTGATCGTGTGGGGTATCCATCACTCTGTGAGTACAGCAGAACAGACTAAACTGTATGGATCAGGCAATAAGT TGGTGACAGTAGGCTCTTCGAATTACCAACAGTCATTTGTCCCCAGCCCGGGAGCCCGGCCACAGGTCAACGGCCAAAGCGGACGCAT CGATTTCCACTGGCTTATGCTGAATCCTAATGACACCGTCACATTTTCCTTCAATGGGGCCTTTATTGCCCCAGATCGTGCATCCTTTCTG AGAGGGAAATCCATGGGTATACAGAGCGGGGTGCAGGTGGACGCAAACTGTGAGGGAGATTGTTACCATTCTGGCGGTACGATCATCA GCAACTTGCCATTCCAGAATATAGACTCCCGGGCAGTCGGTAAGTGTCCCCGGTACGTCAAACAAAGAAGCTTGCTGCTGGCCACCGGG ATGAAAAATGTTCCCGAGATCCCCAAGGGGCGTGGCCTCTTTGGCGCAATCGCAGGGTTTATCGAGAATGGGTGGGAAGGGCTGATCG ACGGCTGGTATGGATTCAGGCACCAAAACGCACAGGGAGAAGGGACAGCCGCCGACTACAAAAGCACACAGAGTGCGATCGATCAAAT CACAGGTAAATTGAACCGACTGATTGAGAAGACAAACCAACAGTTTGAGCTCATAGATAATGAGTTCAACGAGGTGGAGAAGCAGATAGG AAATGTGATCAACTGGACAAGGGATTCCATCACTGAAGTCTGGTCCTATAATGCCGAGCTGTTGGTTGCCATGGAAAACCAGCATACCAT TGACCTCGCCGATTCCGAAATGGATAAGCTGTACGAAAGGGTGAAGCGGCAGTTGCGGGAGAACGCAGAAGAAGATGGAACAGGATGC TTTGAGATATTTCACAAATGCGATGATGATTGCATGGCCTCGATACGCAACAATACCTATGACCATAGCAAGTACCGAGAGGAGGCGATG CAAAACAGGATCCAGATTGACCCAGTGAAGCTCAGCAGTGGGTACAAGGATGTCATTTTGTGGTTTAGCTTTGGAGCCTCATGCTTCATA CTCCTCGCCATCGTCATGGGGTTGGTATTTATCTGCGTCAAGAACGGCAATATGAGGTGTACTATCTGCATC
2793	ATGAATACTCAGATTCTGGTGTTTGCGCTTATCGCTATTATACCCACTAATGCCGACAAAATCTGTCTGGGCCATCACGCCGTTAGCAACG GCACCAAAGTGAACACACTCACAGAACGTGGTGTGGAAGTCGTCAATGCCACCGAGACAGTTGAGCGCACCAATATTCCTAGAATCTGTT CCAAAGGCAAAAAAACTGTCGATCTACGCCAATGCGGACTGCTCGGAACTATTACTGGACCACCGCAGTGCGATCAGTTCCTCGAGTTC AGCGCAGACCTGATCATAGAGAGGAGAGAAGGTAGCGACGTTTGTTACCCTGGAAAATTTGTCAACGAGGAGGCGCTTCGACAGATTCT CAGAGAGAGTGGAGGTATAGATAAGGAGGCCATGGGTTTCACGTATTCAGGTATCAGAACGAATGGGGCAACCAGTGCCTGTCGCCGGT CAGGCAGCAGCTTCTATGCAGAGATGAAGTGGCTCCTGTCAAACACAGACAATGCTGCATTCCCACAGATGACCAAGAGCTACAAGAAC ACGCGAAAGTCACCTGCATTAATAGTCTGGGGTATCCACCACTCTGTCTCCACTGCCGAACAGACCAAACTTTACGGATCTGGAAATAAG CTGGTCACTGTGGGGTCTTCTAATTATCAGCAGAGTTTTGTACCCTCGCCCGGGGCCCGGCCCCAGGTAAACGGACAGTCCGGACGGAT TGACTTTCACTGGCTCATGCTTAACCCAAACGACACAGTGACCTTCAGTTTTAACGGCGCTTTCATCGCTCCCGACAGGGCCTCATTCCT GAGGGGAAAGTCGATGGGAATCCAGAGCGGGGTTCAGGTTGATGCAAATTGCGAAGGGGATTGTTACCATTCCGGCGGAACCATTATCT CCAACCTGCCATTTCAGAATATAGATAGCCGCGCCGTGGGAAAGTGCCCTAGGTACGTCAAACAGCGGTCTCTGCTCCTGGCTACAGGC ATGAAAAATGTTCCGGAAATCCCAAAGGGGCGGGGACTCTTCGGAGCCATCGCAGGCTTCATCGAAAACGGCTGGGAGGGGTTAATCG ATGGATGGTATGGCTTCCGCCATCAGAACGCACAGGGGGAGGGCACCGCGGCCGATTATAAGAGTACCCAGAGCGCAATAGATCAGAT AACTGGTAAGTTGAATCGGTTAATTGAAAAAACTAACCAGCAGTTCGAACTAATCGATAATGAGTTCAATGAAGTCGAGAAACAGATAGGC AACGTTATCAATTGGACCAGGGACAGCATAACAGAAGTGTGGTCCTATAACGCCGAACTCCTCGTTGCCATGGAGAACCAGCATACCATT GACCTGGCTGATTCAGACATGGATAAACTATACGAACGGGTCAAGCGTCAATTGAGAGAGAATGCGGAACAGGACGGGACTGGGTGCTT CGAAATCTTCCATAAGTGTGATGACGATTGTATGGCGTCCATTCGCAATAACACCTACGATCATTCAAAATACCGTGAAGAGGCGATGCA GAATAGGATTCAGATCGACCCAGTGAAGCTCAGTAGTGGATATAAAGACGTAATTCTCTGGTTCTCTTTCGGAGCTAGCTGCTTCATTCTT CTGGCCATCGTGATGGGACTCGTTTTCATATGTGTTAAAAATGGCAACATGCGTTGCACAATCTGTATC

2794	ATGAACACTCAGATTTTAGTTTTCGCTCTAATTGCCATTATTCCTACAAATGCTGATAAAATATGCCTGGGACACCATGCTGTCTCCAACGG TACCAAAGTCAATACTCTTACGGAGAGGGGGGTGGAGGTAGTGAACGCCACCGAGACAGTCGAACGTACAAATATACCTCGGATTTGCT CCAAGGGAAAGAAGACAGTGGATCTGGGGCAATGTGGTCTGCTGGGAACTATCACAGGCCCACCGCAGTGCGACCAGTTTCTTGAGTTT TCAGCCGACCTGATCATCGAGCGACGCGAGGGCAGCGACGTGTGTTACCCCGGGAAGTTCGTAAACGAGGAAGCTCTCCGCCAGATCC TGCGCGAGAGTGGAGGTATTGATAAGGAGGCCATGGGCTTTACATATTCTGGCATCCGGACCAATGGGGCCACAAGCGCCTGTAGGAG ATCTGGATCCAGTTTTTACGCCGAAATGAAATGGCTGCTGAGCAACACAGACAATGCCGCCTTTCCTCAGATGACAAAGAGCTACAAAAA CACAAGGAAGTCACCCGCCCTTATTGTCTGGGGTATCCATCACTCTGTCAGTACAGCTGAGCAGACCAAGCTGTATGGTTCTGGCAATAA ATTGGTTACCGTCGGGAGCAGTAACTATCAACAATCATTTGTTCCAAGTCCAGGAGCCCGGCCCCAGGTCAATGGTCAAAGTGGAAGGA TTGACTTTCACTGGTTAATGCTTAATCCCAACGATACCGTGACCTTCTCATTTAATGGGGCCTTTATTGCTCCGGACCGTGCCAGCTTTCT CCGCGGTAAGTCCATGGGCATCCAGTCCGGCGTACAAGTGGATGCCAATTGCGAAGGCGACTGTTACCACTCTGGCGGCACCATCATCT CAAACCTTCCGTTCCAGAACATAGATAGTCGGGCCGTGGGCAAGTGCCCTAGGTATGTGAAGCAGCGCTCATTACTGTTGGCTACTGGC ATGAAAAATGTGCCAGAGATCCCTAAAGGTCGTGGTCTTTTTGGAGCTATCGCTGGGTTCATTGAAAATGGATGGGAGGGGCTGATCGAC GGTTGGTACGGATTCCGGCATCAGAATGCACAGGGCGAAGGCACAGCCGCTGATTACAAGAGCACCCAGTCAGCTATTGACCAGATCAC AGGGAAGCTGAACCGGCTAATCGAGAAAACAAACCAACAATTTGAACTTATCGATAACGAATTTAATGAGGTGGAGAAACAGATAGGCAA CGTGATCAACTGGACACGGGATAGTATCACTGAGGTGTGGAGTTACAATGCCGAACTCTTGGTGGCTATGGAGAATCAGCATACTATCGA TCTAGCGGACTCTGAGATGGACAAACTCTATGAACGAGTGAAGAGGCAGCTTCGCGAGAACGCTGAGGAAGATGGGACAGGGTGTTTTG AAATTTTTCATAAGTGTGATGATGATTGCATGGCATCTATCCGGAATAACACATACGACCATTCCAAGTACCGTGAAGAGGCTATGCAAAA CCGAATTCAGATAGACCCAGTGAAGCTCTCTTCCGGATACAAGGATGTAATTCTGTGGTTTTCATTCGGTGCATCCTGCTTCATCCTGCTT GCAATCGTGATGGGTTTAGTGTTCATCTGTGTGAAGAATGGGAATATGAGGTGTACAATTTGCATC
2795	ATGAACACGCAGATCCTGGTGTTCGCACTGATTGCAATTATTCCCACCAATGCCGATAAGATCTGTTTAGGACATCACGCCGTGTCCAAC GGTACGAAGGTGAACACATTGACCGAGCGGGGAGTCGAAGTGGTAAATGCAACCGAGACTGTTGAACGCACAAATATCCCAAGAATTTG TTCAAAGGGTAAAAAAACCGTGGACCTCGGACAGTGCGGGCTGCTTGGAACGATTACAGGACCTCCACAATGCGACCAGTTCCTGGAAT TTTCCGCTGACCTCATCATAGAACGACGGGAAGGGTCCGACGTGTGTTATCCGGGAAAGTTCGTTAACGAGGAGGCTCTGAGGCAGATC TTGAGGGAGAGTGGTGGCATCGACAAGGAGGCGATGGGCTTCACATACAGTGGCATCAGAACTAACGGAGCCACCAGCGCATGTCGAC GGTCTGGTAGCAGTTTTTATGCAGAGATGAAGTGGTTGCTGAGCAATACAGATAATGCCGCATTCCCCCAAATGACAAAGTCCTATAAGA ACACGCGGAAAAGTCCTGCCCTAATTGTGTGGGGAATCCACCACTCTGTTAGTACCGCTGAGCAGACCAAGTTATACGGCAGCGGGAAC AAGTTGGTGACAGTTGGATCTAGTAACTACCAGCAAAGCTTCGTTCCTAGCCCGGGGGCACGTCCCCAGGTGAACGGGCAGAGTGGCC GGATCGATTTTCATTGGCTGATGCTCAATCCAAACGACACTGTCACATTCAGTTTTAACGGAGCCTTTATCGCCCCTGATAGAGCCTCATT CCTTAGGGGAAAAAGTATGGGAATCCAGTCTGGTGTACAGGTGGATGCCAACTGTGAAGGTGACTGTTATCATTCAGGTGGAACCATCAT ATCAAACCTGCCCTTCCAAAATATTGACTCAAGAGCAGTGGGCAAGTGCCCCCGATATGTGAAGCAGCGATCCCTGCTGCTCGCAACTG GCATGAAGAATGTCCCGGAAATCCCTAAGGGGCGGGGCCTTTTCGGGGCAATCGCAGGCTTCATCGAAAATGGATGGGAGGGCCTGAT CGACGGATGGTACGGTTTCCGGCATCAGAATGCTCAGGGGGAGGGGACCGCGGCTGACTATAAGTCGACACAATCTGCCATAGATCAG ATCACGGGAAAACTGAACAGACTGATAGAGAAGACTAATCAGCAGTTCGAACTGATCGATAACGAGTTCAATGAGGTTGAAAAGCAAATC GGGAACGTAATCAACTGGACACGAGACAGCATTACTGAGGTGTGGTCCTACAATGCGGAATTGCTGGTGGCTATGGAAAATCAACACAC AATCGATCTTGCCGATTCAGAGATGGACAAGCTTTACGAGAGAGTAAAGCGACAACTGCGCGAGAATGCTGAGGAGGACGGGACTGGGT GCTTCGAGATCTTTCACAAGTGTGACGACGACTGCATGGCCTCTATCCGGAATAATACTTATGACCATAGCAAATATAGAGAGGAGGCCA TGCAAAACCGGATCCAGATTGATCCAGTCAAGCTCTCCAGCGGTTATAAGGATGTGATTTTGTGGTTCAGTTTTGGCGCTTCATGCTTCAT ACTCTTGGCCATCGTGATGGGCCTGGTGTTTATTTGTGTGAAGAACGGAAACATGCGGTGTACAATTTGCATC
2796	ATGAATACCCAGATCCTCGTTTTCGCTCTGATCGCAATCATTCCCACAAACGCAGATAAAATCTGTCTGGGGCATCACGCTGTGAGCAAC GGCACAAAAGTTAACACCTTAACCGAGCGCGGCGTGGAGGTGGTTAATGCTACCGAGACCGTCGAACGGACCAATATCCCCAGGATCTG TAGTAAAGGCAAGAAGACAGTCGACCTGGGGCAGTGTGGGCTCCTCGGGACTATCACAGGACCCCCTCAGTGTGATCAGTTCCTCGAGT TCTCAGCGGACCTTATTATAGAACGCCGAGAGGGTTCTGATGTGTGTTACCCCGGCAAATTTGTGAACGAGGAGGCTCTGAGGCAAATTC TGCGCGAAAGCGGCGGCATTGAAAAGGAAGCTATGGGGTTTACATACTCCGGAATAAGAGCCAATGGCGCGACTTCTGCTTGTCGAAGG TCCGGAAGCTCCTTTTATGCGGAAATGAAATGGCTATTGAGCAATACCGACAATGCTGCCTTCCCCCAGATGACCAAGAGTTATAAAAATA CCCGAAAAAGCCCCGCGCTTATCGTTTGGGGCATCCATCATTCTGTTTCTACAGCCGAACAGACTAAGCTTTATGGCAGCGGGAATAAGC TCGTGACCGTCGGCTCAAGTAACTATCAGCAGTCATTCGTGCCTTCCCCGGGGGCTAGACCTCAAGTGAATGGATTATCCGGACGAATC GACTTCCATTGGTTGATGTTAAACCCTAATGATACTGTGACCTTCTCCTTCAACGGGGCCTTCATTGCTCCAGATCGTGCTAGTTTCCTGC GGGGCAAGTCCATGGGTATTCAATCCGGAGTCCAGGTCGACGCAAACTGTGAGGGGGATTGCTACCACTCCGGAGGCACAATTATTTCT AATCTGCCATTTCAGAACATTGACTCCCGCGCCGTGGGGAAATGTCCAAGGTATGTGAAACAACGGAGCCTACTGCTCGCCACCGGAAT GAAGAACGTTCCTGAAATACCTAAAGGTCGGGGTTTGTTTGGGGCTATAGCCGGCTTCATAGAAAACGGATGGGAAGGCCTGATCGATG GCTGGTATGGCTTTAGGCATCAAAATGCACAGGGGGAGGGCACTGCGGCTGACTATAAGAGCACCCAGTCAGCGATTGACCAGATTACT GGCAAGCTGAATAGACTTATAGAAAAGACAAATCAGCAATTCGAGCTGATAGACAACGAATTCAATGAGGTGGAGAAACAAATTGGTAAC GTCATTAACTGGACTCGTGACTCTATTACGGAAGTGTGGAGTTACAATGCTGAGCTCCTCGTCGCCATGGAAAACCAACACACTATAGAT TTCGCCGATAGCGAGATGGACAAGCTGTACGAGACGCGTAAAACGCCAGTTGAGAGAAAACGCCGACGAAGATGGCACCGGTTGCTTTG AGATTTTCCACAAGTGCGATGATGACTGCATGGCAAGCATCAGAAATAACACATACGATCATAGTAAATACAGGGAGGAAGCCATGCAAA ACCGCATTCAGATCGATCCCGTCAAGCTCTCTTCGGGGTACAAAGATGTCATCCTTTGGTTCTCATTCGGCGCATCTTGCTTTATCCTGCT GGCTATAGTGATGGGGCTCGTGTTTATTTGCGTCAAAAATGGAAACATGCGCTGTACCATCTGTATT

2797	ATGAATACTCAGATCCTGGTGTTTGCGCTCATTGCGATCATCCCAACCAACGCCGACAAAATATGTCTGGGACATCACGCCGTTTCCAAC GGGACGAAGGTGAACACGTTGACAGAGCGCGGAGTGGAGGTCGTGAATGCTACTGAGACCGTGGAGAGGACTAATATCCCGAGGATCT GCAGTAAGGGCAAGAAGACTGTCGACTTAGGACAGTGTGGCCTGTTAGGCACAATCACAGGCCCCCCTCAATGTGACCAATTTCTGGAG TTTTCTGCCGACCTCATAATCGAGCGCAGGGAAGGCTCCGACGTGTGCTATCCGGGAAAGTTCGTCAACGAGGAAGCCCTCAGACAGAT CCTTCGCGAATCCGGCGGCATAGATAAGGAAGCTATGGGTTTCACATATTCAGGAATCCGCACGAATGGCGCAACTTCCGCTTGCCGGC GATCAGGATCCTCTTTCTACGCAGAGATGAAGTGGCTGCTGAGCAATACCGACAATGCCGCCTTCCCCCAAATGACAAAGTCATACAAGA ATACTCGTAAATCACCTGCCCTGATTGTGTGGGGAATCCATCACTCAGTGTCCACAGCGGAGCAGACCAAGCTGTACGGAAGCGGTAAC AAGCTTGTTACCGTTGGCTCTAGCAATTATCAGCAGAGCTTTGTTCCTTCACCTGGAGCCAGACCTCAAGTAAATGGGCAAAGCGGTAGA ATCGACTTTCACTGGCTGATGCTGAATCCCAATGACACAGTCACCTTTTCGTTTAACGGCGCGTTTATAGCACCCGATAGGGCCAGTTTTT TGAGAGGAAAGTCAATGGGAATCCAAAGCGGCGTGCAAGTCGATGCCAATTGTGAGGGCGACTGTTACCATAGTGGGGGAACCATTATA AGTAATTTACCCTTTCAGAATATTGACTCAAGGGCTGTGGGAAAATGTCCCAGGTATGTCAAGCAGAGAAGCCTGCTGCTGGCCACCGGT ATGAAGAACGTACCCGAGATTCCAAAAGGCAGAGGACTGTTTGGAGCAATTGCTGGGTTCATCGAGAACGGGTGGGAGGGGCTTATTGA TGGCTGGTACGGATTTAGACATCAAAATGCACAGGGGGAAGGCACTGCCGCTGATTATAAAAGTACGCAGTCCGCGATAGACCAGATTA CCGGAAAGTTGAACCGGCTGATCGAGAAAACTAATCAGCAGTTTGAGTTGATCGACAACGAATTTAACGAGGTGGAGAAACAAATCGGCA ATGTGATTAACTGGACTAGAGACTCCATTACCGAGGTGTGGTCATATAACGCCGAATTGCTCGTGGCCATGGAAAACCAGCACACCATCG ATCTCGCTGATAGCGAGATGGACAAACTCTACGAAAGGGTCAAGAGACAACTCCGAGAGAACGCCGAGGAAGACGGGACTGGGTGTTTT GAGATCTTTCATAAATGTGATGACGACTGTATGGCTTCTATCCGGAATAACACATACGACCACTCCAAATATCGGGAGGAAGCTATGCAAA ATCGTATTCAAATTGATCCCGTCAAGCTGAGCTCCGCGTATAAGGACGTTATCCTCTGGTTCTCCTTTGGCGCTTCCTGTTTCATCCTTCT GGCTATTGTCATGGGGCTCGTTTTCATATGCGTTAAGAACGGGAATATGCGGTGTACGATCTGTATA
2798	ATGAACACACAGATATTGGTGTTTGCTTTGATTGCAATCATTCCTACAAACGCGGATAAAATCTGCCTGGGCCACCATGCTGTGAGTAACG GCACCAAAGTGAATACTTTAACTGAACGCGGTGTCGAGGTTGTGAACGCGACTGAAACGGTGGAGCGCACGAATATTCCCCGGATCTGC AGCAAAGGCAAAAAGACTGTCGACCTTGGTCAGTGCGGGCTACTTGGGACTATCACTGGACCCCCACAGTGTGATCAATTCCTGGAATTT TCAGCCGATTTGATTATTGAGCGGAGACAGGGATCCGACGTGTGCTATCCAGGCAAGTTTGTGAACGAGGAGGCATTGCGGCAGATTCT CAGGGAATCTGGAGGCATTGATAAGGAAGCTATGGGTTTCACCTATAGCGGTATCCGGACCAACGGAGCTACCAGCGCGTGCCGGAGG AGCGGTAGCTCCTTTTACGCGGAGATGAAGTGGTTGCTCTCTAACACGGACAACGCCGCCTTCCCACAGATGACCAAGAGCTACAAGAA CACCCGCAAATCACCTGCCTTAATTGTGTGGGGCATTCATCATAGTGTCTCTACGGCCGAGCAGACAAAACTGTATGGATCTGGCAATAA GCTTGTCACCGTCGGCAGCTCCAATTACCAGCAGTCTTTCGTACCCAGTCCCGGTGCAAGGCCACAGGTCAACGGACAGTCGGGGCGG ATTGACTTCCACTGGCTGATGCTCAACCCTAATGATACTGTAACCTTCAGCTTCAATGGCGCATTCATAGCTCCCGATCGGGCATCATTCC TGCGAGGTAAATCTATGGGGATCCAGTCCGGCGTGCAGGTGGATGCCAACTGCGAGGGCGATTGCTATCACAGTGGTGGGACAATCAT CTCTAATTTACCCTTCCAGAATATTGATAGCCGAGCTGTGGGGAAATGCCCAAGGTACGTCAAGCAGCGTTCGCTTCTGTTAGCCACTGG CATGAAAAATGTGCCAGAGATACCCAAAGGCAGGGGGTTATTTGGAGCAATTGCCGGTTTTATTGAAAATGGCTGGGAGGGACTCATCG ACGGCTGGTATGGCTTCAGACACCAAAATGCCCAGGGTGAAGGTACCGCCGCCGACTACAAGTCCACTCAAAGTGCAATTGACCAAATA ACTGGAAAGCTGAACCGGCTGATCGAAAAGACAAATCAGCAGTTTGAGTTAATCGATAACGAGTTCAACGAGGTCGAGAAGCAGATTGGA AATGTCATTAACTGGACAAGGGATTCCATCACTGAGGTGTGGAGCTACAACGCCGAGCTGCTCGTCGCAATGGAGAACCAGCATACTATT GACCTGGCAGACTCTGAAATGGATAAATTATACGAACGCGTGAAACGCCAGCTCAGAGAAAATGCTGAAGAGGACGGGACCGGTTGTTT TGAGATTTTTCACAAGTGCCACGACCATTGCATGGCCTCCATTAGGAACAATACATACGACCATAGTAAGTATCGAGAGGAAGCCATGCA GAACAGGATACAGATAGACCCTGTTAAACTGTCCTCCGGGTACAAGGATGTGATCCTTTGGTTTTCTTTTGGGGCATCCTGCTTTATTCTG CTGGCAATCGTCATGGGCCTTGTTTTTATCTGCGTAAAGAATGGAAATATGAGATGCACAATCTGTATC
2799	ATGAACACCCAGATCCTTGTATTCGCATTAATAGCTATCATTCCAACAAACGCGGACAAAATTTGCCTGGGGCACCATGCAGTCTCCAACG GCACTAAGGTGAATACGCTCACCGAGCGGGGCGTGGAGGTTGTTAATGCGACGGAGACGGTTGAGCGGACGAATATCCCCCGAATTTG TTCGAAGGGAAAACGGACAGTGGACCTGGGTCAGTGTGGGTTGCTGGGTACCATTACAGGCCCCCCCCAGTGTGATCAGTTTCTGGAAT TTTCCGCCGACCTTATTATCGAGCGGAGGGAAGGGTCCGATGTCTGTTACCCAGGGAAGTTCGTTAACGAGGAGGCTTTGCGCCAGATA CTCCGGGAGAGTGGTGGTATTGACAAGGAGGCCATGGGGTTCACCTACAGCGGCATCAGGACTAATGGAGCCACCTCGGCCTGTAGAA GAAGTGGATCTTCTTTCTATGCCGAGATGAAATGGTTGCTATCTAACACTGACAACGCCGCCTTTCCACAGATGACTAAAAGTTACAAAAA TACACGGAAAAGCCCAGCCTTAATAGTCTGGGGCATCCATCATAGTGTGTCCACCGCCGAACAGACTAAACTATATGGGTCCGGAAGCA AGCTTGTGACGGTGGGCTCTTCCAACTACCAGCAGAGTTTTGTGCCCTCCCCCGGAGCACGGCCTCAGGTCAACGGCTTGAGCGGACG GATCGACTTTCATTGGTTAATGCTGAATCCAAATGATACAGTAACATTTTCATTTAATGGCGCCTTCATTGCTCCAGACAGAGCTAGTTTCT TACGGGGCAAATCAATGGGCATCCAGAGCGGGGTTCAGGTGGATGCCAACTGTGAAGGAGATTGTTATCACAGCGGCGGGACTATCATT TCTAATCTCCCATTTCAGAATATCGATTCACGGGCAGTTGGCAAGTGCCCGAGATACGTCAAGCAGCGATCTCTCCTTCTCGCAACCGGA ATGAAAAACGTCCCAGAGATTCCTAAAGGCAGGGGCCTGTTCGGCGCCATCGCCGGATTTATTGAGAACGGGTGGGAGGGGCTGATAG ATGGCTGGTACGGCTTTCGGCATCAGAATGCACAGGGCGAGGGTACGGCTGCGGACTACAAAAGTACTCAAAGCGCAATCGACCAGATT ACAGGAAAGCTGAACCGGTTAATTGAAAAGACAAACCAGCAATTCGAGCTCATCGACAATGAGTTTAATGAAGTGGAGAAGCAGATCGGC AATGTAATCAACTGGACCAGAGATAGTATCACTGAAGTGTGGTCCTACAACGCCGAGCTTCTGGTGGCTATGGAGAACCAGCATACTATC GATCTTGCGGATAGCGAGATGGATAAGCTGTACGAGCGTGTTAAGCGACAGCTAAGAGACAACGCGGAGGAGGACGGTACTGGCTGTT TCGAAATTTTTCACAAGTGCGATGACGATTGTATGGCGTCTATCCGCAATAATACTTATGATCATAGTAAGTATCGGGAAGAAGCTATGCA GAACCGCATCCAAATCGACCCCGTGAAGCTTTCAAGCGGTTACAAAGACGTTATCTTGTGGTTCAGCTTCGGGGCGTCTTGCTTTATCCT GCTGGCGATTGTAATGGGCTTGGTCTTTATCTGTGTGAAGTCCAGAAATATGAGATGCACCATTTGTATC

2800	ATGAATACCCAAATCCTCGTGTTCGCTTTAATCGCAATCATCCCGACCAATGCCGACAAGATCTGCCTGGGTCACCACGCTGTCTCCAAC GGCACTAAGGTGAACACCTTAACAGAGAGAGGCGTGGAAGTAGTTAATGCTACCGAAACCGTGGAAAGGACCAACATTCCGCGGATATG TTCTAAGGGAAAACGCACCGTAGATCTCGGCCAGTGTGGACTGCTGGGCACAATCACCGGTCCTCCACAATGTGATCAGTTTCTTGAGTT CTCGGCCGACCTCATCATCGAAAGGAGGGAAGGATCTGATGTGTGTTATCCCGGGAAGTTCGTGAAAGAAGAGGCCCTGCGTCAGATAT TGCGACACAGTGGCGGGATAGATAAGGAAGCAATGGGCTTCACATACACGCGCATCAGAACAAATGGCGCGACATCCGCCTGTAGAAG GTCTGGGTCTAGCTTCTACGCAGAGATGAAATGGCTGCTCAGTAACACGGACAATGCGGCCTTCCCCCAGATGACTAAAAGCTATAAAAA TACAAGGAAGAGCCCCGCCTTGATAGTGTGGGGAATCCATCATAGTGTGAGTACCGCTGAACAGACGAAGCTATACGGATCAGGCAACA AGCTGGTTACTGTCGGCAGTTCTAACTACCAGCAATCCTTTGTGCCTTCTCCCGGCGCCCGCCCACAAGTGAATGGACTATCTGGCAGAA TTGATTTCCACTGGCTGATGCTGAACCCCAATGACACTGTGACTTTCAGCTTTAACGGAGCCTTTATCGCCCCCGATAGAGCCAGTTTCCT GAGGGGCAAATCCATGGGCATCCAGTCAGGCGTGCAGGTCGATGCCAACTGCGAGGGAGATTGTTACCATTCGGGTGGCACCATAATTA GTAACTTACCATTCCAGAATATTGACAGCAGAGCCGTTGGGAAATGCCCCAGGTATGTCAAGCAGAGGTCCCTACTACTGGCCACAGGC ATGAAGAATGTTCCTGAAATTCCTAAAGGCCGTGGTCTTTTCGGAGCAATCGCAGGATTCATCGAAAACGGCTGGGAAGGACTGATTGAT GGGTGGTATGGTTTTAGGCACCAAAATGCACAGGGCGAGGGCACAGCTGCAGATTACAAGAGCACTCAGTCAGCTATTGATCAGATTAC CGGAAAGCTTAACAGGCTCATCGAAAAGACTAATCAGCAGTTTGAACTGATCGACAACGAATTCAACGAAGTAGAGAAACAGATTGGGAA CGTGATTAATTGGACCCGAGACTCTATCACAGAGGTGTGGAGTTACAACGCAGAGTTGCTGGTAGCAATGGAAAACCAGCACACAATCG ATCTCGCAGACAGTGAGATGGATAAACTTTACGAAAGGGTCAAACGTCAGTTAAGAGAGAACGCGGAAGAGGATGGCACCGGTTGCTTT GAAATTTTCCACAAATGCGATGACGACTGCATGGCGTCTATCAGAAATAACACTTACGACCATAGTAAGTATCGGGAGGAGGCTATGCAG AACCGAATTCAGATCGATCCAGTAAAGCTCAGCAGCGGCTACAAGGACGTCATCCTTTGGTTCTCTTTCGGGGCCAGTTGCTTCATCCTT CTCGCTATAGTCATGGGCCTCGTGTTTATCTGCGTAAAAAATGGGAACATGAGGTGCACCATTTGCATA
2801	ATGAATACACAGATATTGGTCTTCGCTCTCATCGCTATCATTCCCACCAACGCTGACAAAATCTGTTTAGGCCACCATGCCGTATCCAATG GCACTAAGGTCAACACACTTACTGAGCGAGGCGTAGAGGTCGTCAACGCTACAGAAACTGTTGAGCGAACCAACATCCCCAGGATTTGC TCTAAGGGGAAAAAGACTGTAGACCTTGGGCAGTGTGGCCTGCTGGGGACTATCACCGGGCCACCTCAATGTGACCAATTTCTGGAGTT CTCACCTGATCTGATTATCGAGAGGCGCGAGGGATCTGATGTCTGTTACCCAGGTAAGTTCGTCAATGAGGAGGCGCTCCGCCAGATCC TCCGCGAGAGTGGCGGGATCGATAAAGAGGCCATGGGCTTCACTTACTCCGGCATCAGGACCAATGGCGCTACATCCGCCTGTAGACG GAGCGGATCCAGCTTTTACGCCGAAATGAAGTGGCTGCTGTCCAACACAGACAACGCTGCTTTTCCACAGATGACAAAGTCTTATAAAAA TACCCGCAAATCTCCTGCCCTGATTGTGTGGGGCATTCACCACTCAGTGTCCACTGCCGAGCAGACTAAGTTGTACGGAAGTGGCAACA AGCTGGTGACGGTCGGCTCTTCAAACTATCAACAGAGCTTTGTACCTTCCCCAGGGGCCCGGCCACAGGTGAACGGGCAGTCAGGAAG GATTGATTTCCACTGGCTGATGCTGAATCCAAATGATACTGTCACCTTCAGCTTTAATGGAGCTTTTATTGCCCCGGATCGCGCATCCTTC CTGCGCGGGAAGAGTATGGGGATCCAGTCCGGGGTACAAGTCGACGCTAACTGCGAAGGCGATTGCTACCATAGCGGAGGAACCATCA TTAGTAACCTCCCCTTTCAGAATATTGATTCCCGTGCCGTTGGGAAATGTCCCAGATACGTCAAGCAGAGGTCCTTGCTGCTCGCAACCG GCATGAAGAATGTGCCAGAGATTCCAAAGGGGAGAGGTCTTTTCGGTGCTATCGCCGGCTTTATCGAGAACGGCTGGGAAGGGCTGATC GATGGCTGGTACGGATTCCGGCATCAAAACGCGCAGGGAGAGGGAACTGCGGCCGACTACAAGTCAACTCAGTCTGCCATCGATCAGA TTACTGGTAAGCTTAACCGTCTCATCGAGAAAACCAACCAACAATTTGAATTAATTGATAACGAGTTTAACGAAGTGGAGAAGCAAATTGG GAATGTCATAAACTGGACACGGGATTCTATCACCGAGGTATGGTCTTACAACGCGGAGTTGCTGGTCGCCATGGAGAACCAACATACCAT CGATCTGGCTGACAGTGAAATGGACAAGCTGTATGAGCGGGTGAAACGGCAGCTCCGGGAGAATGCAGAGGAAGACGGTACTGGCTGT TTCGAGATCTTCCATAAGTGCGACGATGACTGTATGCCCTCGATTAGGAACAACACCTATGATCACAGCAACTATCGCGAGGAAGCGATG CAGAACCGGATTCAGATCGACCCAGTTAAACTCTCGTCGGGCTACAAGGATGTAATCCTGTGGTTTAGCTTCGGAGCGTCGTGTTTCATC TTGTTGGCGATAGTAATGGGCCTTGTTTTTATCTGCGTGAAGAACGGAAATATGAGATGCACTATCTGTATT
2802	ATGAACACGCAGATCCTAGTGTTCGCACTTATCGCCATCATCCCTACCAATGCTGACAAAATTTGCCTGGGACATCATGCGGTTTCTAATG GAACCAAGGTGAATACCCTAACAGAACGGGGGGTGGAAGTTGTCAATGCCACAGAAACGGTTGAGCGTACTAACATTCCCCGCATTTGC TCCAAAGGAAAAAAAACTGTGGATCTGGGCCAGTGTGGACTTCTGGGCACTATTACCGGCCCTCCTCAGTGCGACCAGTTCCTGGAGTT CTCTGCGGACCTTATTATTGAGAGACGTGAGGGATCTGATGTGTGTTATCCAGGGAAGTTCGTAAACGAAGAGGCTCTTAGACAAATCCT GCGGAAGAGCGGTGGCATAGATAAAGAGGCAATGGGGTTTACATACAGCGGCATAAGGACCAACGGCGCCACATCCACTTGCAGGCGG TCCGGGTCGTCATTTTACGCCGAGATGAAATGGTTGTTAAGTAATACAGACAACGCAGCCTTTCCGCAGATGACTAAATCTTATAAAAACA CACGAAAATCTCCGGCGATCATTGTATGGGGCATACACCACAGTGTTAGCACTGCCGAGCAGACGAAACTATATGGCTCCGGAAATAAAC TGGTCACCGTAGGATCCTCCAACTACCAGCAGAGCTTCGTGCCCTCGCCTGGCGCACGGCCACAAGTTAATGGTCTCAGCGGTCGGATT GACTTTCACTGGCTCATGCTTAATCCTAATGACACAGTAACCTTCTCCTTTAACGGCGCCTTCATAGCTCCCGACAGGGCATGTTTTCTGC GCGGGAAGAGTATGGGGATCCAGTCCGGGGTCCAGGTAGATGCAGACTGCGAAGGGGACTGCTACCACAGTGGGGGTACCATCATTAG TAACCTTCCCTTTCAGAATATTGACAGCAGGGCCGTCGGGAAGTGCCCCCGCTACGTGAAGCAGCGGTCTTTGCTCCTGGCTACCGGTA TGAAGAACGTGCCCGAGATACCAAAAGGCAGGGGGCTCTTTGGTGCCATCGCCGGCTTTATCGAAAATGGATGGGAGGGCCTCATCGA CGGTTGGTACGGTTTCAGGCACCAGAATGCCCAGGGGGAAGGAACAGCTGCAGATTACAAGTCAACCCAGTCTGCTATTGACCAGATAA CAGGGAAACTGAACCGATTAATTGAGAAAACTAATCAACAGTTCGAGCTGATTGATAACGAGTTTAACGAGGTGGAACGCCAAATCGGGA ACGTGATTAACTGGACAAGGGATTCAATCACAGAGGTTTGGAGTTATAATGCCGAACTGCTGGTCGCCATGGAGAACCAGCACACTATCG ATCTAGCTGATTCTGAGATGGACAACCTCTATGAGCGCGTCAAGCGTCAGCTCCGCGAAAACGCTGAACAAGATGGCACTGGCTGTTTC GAGATCTTCCACAAGTGTGACGACGACTGCATGGCCTCCATAAGGAATAACACTTATGATCACAGCAAGTATCGAGAGGAGGCCATGCA GAACCGCATTCAGATTGATCCCGTTAAGTTAAGTTCTGGGTATAAAGATGTCATTCTTTGGTTCTCTTTTGGTGCTTCATGTTTTATACTGC TGGCTATCGTCATGGGACTCGTGTTCATCTGCGTCAAAAATGGGAATATGCGGTGTACCATATGCATT

2803	ATGAATACGCAGATCCTGGTGTTCGCTCTAATCGCTATTATCCCTACAAACGCAGATAAGATCTGTTTAGGCCACCATGCAGTTTCTAATG GCACTAAGGTCAACACTCTCACTGAACGCGGGGTGGAGGTAGTCAACGCTACAGAGACAGTTGAACGAACAAACATCCCCAGGATTTGC TCTAAGGGTAAAAAAACAGTGGACCTGGGACAGTGCGGACTGCTCGGGACCATTACTGGGCCACCTCAGTGTGACCAGTTCTTGGAGTT TTCCGCCGATCTAATCATCGAAAGACGAGAAGGATCCGATGTGTGCTATCCTGGCAAGTTTGTGAATGAGGAAGCACTGCGGCAGATCTT GCGGGAGTCTGGCCGGAATTGATAAGGAAGCAATGGGTTTTACCTACTCCGGTATTCGGACAAATGGAGCTACATCCGCATGTCGGCGGT CCGGAAGCAGTTTCTACGCCGAGATGAAGTGGCTATTGTCAAACACAGATAACGCTGCCTTCCCCCAGATGACTAAGAGCTATAAGAACA CCCGGAAGAGCCCTGCCCTGATCGTCTGGGGCATTCATCATTCGGTCAGTACAGCCGAACAAACTAAGTTGTACGGCTCCGGAAACAAG TTGGTTACAGTGGGGTCTTCCAACTACCAGCAGAGCTTTGTACCTTCCCCCGGAGCCCGACCCCAGGTCAACGGCCAGTCTGGAAGAAT TGATTTTCATTGGCTAATGCTGAATCCTAATGATACTGTGACATTTTCATTCAATGGAGCCTTCATAGCACCTGATAGGGCCTCTTTCCTGC GGGGAAAAAGCATGGGTATTCAGAGTGGCGTTCAGGTGGATGCTAATTGCGAGGGGGACTGCTATCATAGTGGTGGGACCATTATTAGC AACCTCCCCTTTCAGAATATAGATAGCAGGGCTGTCGGCAAATGTCCAAGGTACGTTAAGCAGCGGAGTCTACTGTTGGCCACCGGTAT GAAAAACGTCCCCGAGATCCCTAAGGGACGCGGCCTGTTCGGCGCCATCGCCGGATTTATCGAGAACGGGTGGGAAGGCTTGATCGAT GGGTGGTATGGATTCCGACATCAGAACGCCCAGGGCGAAGGTACTGCTGCCGACTATAAATCTACGCAGAGCGCAATAGACCAGATTAC GGGGAAACTGAACAGGCTGATCGAGAAGACCAATCAGCAGTTTGAGCTGATTGACAATGAATTCAACGAAGTTGAGAAACAGATCGGTAA CGTGATAAATTGGACTAGAGATAGTATCACCGAGGTATGGAGCTACAACGCCGAGCTGCTTGTCGCTATGGAAAACCAGCACACCATCG ACCTTGCTGACAGTGAGATGGATAAACTCTATGAACGAGTGAAGAGACAGTTGCGGGAAAACGCCGAAGAGGACGGTACCGGGTGCTTT GAGATATTTCACAAGTGCGATGATGACTGTATGGCTAGTATCCGCAATAATACATATGACCATAGTAAGTACAGAGAAGAAGCCATGCAAA ACAGAATACAAATTGACCCCGTGAAGTTAAGCTCAGGATACAAAGATGTCATCCTTTGGTTCTCCTTCGGCGCATCATGCTTTATACTTCT CGCAATCGTGATGGGTCTGGTTTTTATATGCGTCAAGAACGGCAATATGCGCTGTACGATCTGTATT
2804	ATGAACACCCAGATTCTGGTTTTTGCATTGATCGCCATTATTCCGACTAATGCCGACAAAATCTGTTTGGGACACCACGCCGTGAGCAAC GGCACGAAAGTCAACACACTCACAGAACGAGGCGTGGAAGTTGTCAACGCAACCGAGACGGTTGAGCGCACCAACATCCCAAGGATCT GCTCTAAGGGCAAGAAGACCGTGGACCTTGGGCAGTGTGGCCTGCTGGGGACAATAACCGGTCCACCACAATGTGATCAATTCCTGGA GTTTTCAGCCGACCTCATCATAGAACGCCGGGAAGGTTCTGACGTGTGCTACCCGGGCAAATTTGTCAACGAGGAGGCTCTCAGGCAGA TCCTGAGGGAGTCAGGCGGAATTGACAAGGAAGCCATGGGTTTCACGTACAGCGGTATTCGGACCAATGGGGCTACATCAGCCTGTCG CAGATCTGGAAGTTCTTTTTACGCTGAAATGAAGTGGCTGCTGTCAAATACCGATAACGCTGCATTTCCCCAGATGACAAAAAGTTATAAG AATACCAGGAAGTCCCCCGCCTTGATCGTGTGGGGCATACACCACTCAGTGAGTACAGCCGAGCAGACCAAGTTGTATGGATCGGGTAA CAAACTAGTCACCGTGGGCTCCAGCAATTATCAGCAGAGTTTTGTCCCCAGCCCCGGGGCTAGACCGCAAGTCAATGGTCAGTCCGGCC GTATCGATTTTCACTGGCTGATGCTAAATCCCAATGACACTGTAACCTTTTCCTTTAATGGGGCATTCATTGCACCCGACAGGGCATCCTT TTTGCGGGGTAAGAGTATGGGTATCCAGAGTGGCGTCCAGGTGGATGCAAACTGCGAGGGTGACTGCTACCATTCAGGAGGCACGATTA TATCAAATCTGCCCTTTCAGAACATTGACAGTCGCGCCGTTGGCAAATGTCCAAGGTACGTGAAGCAGAGAAGCCTCCTGCTCGCCACTG GCATGAAAAATGTGCCCGAGATTCCCAAAGGCCGGGGTCTTTTTGGCGCAATAGCTGGTTTTATCGAGAATGGTTGGGAAGGACTCATC GACGGCTGGTACGGATTTCGGCACCAAAACGCCCAAGGGGAAGGAACCGCCGCAGATTACAAGAGCACGCAGTCAGCTATCGACCAGA TCACCGGCAAGCTTAATCGGCTTATTGAAAAAACCAATCAACAATTCGAGCTGATTGACAATGAATTTAATGAAGTGGAGAAACAAATTGG GAACGTGATCAATTGGACCAGAGACAGCATAACTGAAGTGTGGAGTTATAACGCAGAACTGCTGGTCGCTATGGAAAATCAACACACTAT TGATCTGGCCGATAGCGAAATGGACAAGCTATACGAAAGGGTTAAACGGCAGCTTCGCGAGAACGCCGAGGAGGACGGCACCGGCTGC TTCGAGATTTTCCACAAGTGTGATGACGATTGTATGGCCTCCATCAGAAACAACACCTACGACCATAGCAAGTATAGACAAGAAGCCATG CAGAATCGCATCCAGATCGACCCCGTTAAGCTTTCATCCGGCTATAAGGACGTTATCTTATGGTTCTCTTTTGGAGCATCATGCTTTATCC TCCTGGCTATCGTTATGGGCCTTGTTTTCATCTGCGTTAAAAACGGGAATATGCGCTGCACCATCTGTATT
2805	ATGAATACACAGATCTTAGTGTTTGCTCTTATAGCAATCATACCCACAAACGCCGATAAGATCTGTCTGGGGCACCATGCCGTGAGCAAC GGAACTAAGGTTAACACATTAACTGAGAGAGGTGTGGAGGTAGTTAACGCGACCGAGACTGTAGAGCGGACTAATATCCCTCGCATTTG CAGTAAGGGCAAAAAGACCGTTGACCTTGGTCAGTGTGGGCTGCTTGGAACCATCACCGGACCTCCTCAGTGTGACCAGTTCCTCGAAT TTTCCGCTGACCTGATTATTGAGAGAAGGGAAGGGAGTGACGTGTGTTATCCTGGCAAATTCGTCAATGAAGAAGCGCTACGCCAGATCC TTAGAGAGAGTGGGGGAATCGATAAGGAGGCAATGGGCTTTACATACAGTGGCATAAGAACCAATGGCGCGACAAGTGCCTGCCGCCG GAGCGGCTCAAGTTTCTATGCTGAGATGAAATGGTTACTCAGTAATACTGATAATGCCGCATTTCCACAGATGACTAAATCTTATAAAAATA CACGGAAATCACCCGCACTGATTGTATGGGGTATCCACCACAGCGTCTCCACAGCGGAGCAAACAAAACTGTACGGCAGTGGGAACAAA TTGGTTACTGTTGGGTCCTCGAATTATCAGCAGTCTTTCGTCCCTAGCCCTGGAGCTCGCCCCCAAGTTAACGGTCAGAGCGGCAGGATA GATTTCCACTGGTTAATGCTAAACCCTAATGACACCGTAACCTTTTCGTTTAATGGAGCATTCATTGCGCCTGACCGCGCGAGCTTCTTAA GGGGAAAGTCAATGGGAATTCAATCTGGGGTCCAAGTCGACGCAAATTGTGAGGGCGACTGCTACCACTCAGGAGGAACTATCATCTCG AACCTGCCTTTTCAAAACATCGATAGCAGAGCCGTTGGCAAGTGCCCTCGCTATGTGAAACAGCGCTCACTTCTCTTAGCGACGGGAATG AAGAACGTTCCGGAGATCCCAAAGGGTCGTGGCTTGTTTGGCGCAATAGCAGGGTTCATAGAGAACGGATGGGAAGGTCTGATTGACGG TTGGTATGGCTTCAGACACCAGAATGCTCAGGGAGAAGGAACCGCAGCCGATTATAAATCAACACAGAGCGCCATAGATCAGATCACAG GAAAGTTAAATCGCCTGATCGAGAAGACGAATCAGCAGTTCGAGCTGATCGACAACGAGTTCAACGAAGTCGAGAAACAAATTGGCAAC GTGATCAACTGGACGAGGGATAGCATTACGGAGGTGTGGTCTTATAACGCTGAACTACTTGTGGCAATGGAGAACCAGCATACAATCGA CCTCGCTGACTCAGAAATGGATAAGCTGTACGAGAGAGTTAAACGGCAGTTGAGACAAAACGCAGAGGAGGATGGCACAGGCTGCTTTG AAATTTTTCACAAGTGCGATGATGACTGCATGGCTTCCATTCGGAACAATACGTATGACCACTCTAAATACAGGGAAGAGGCCATGCAGA ATCGCATCCAGATCGATCCTGTAAAGTTGTCCTCAGGCTATAAAGATGTGATCCTCTGGTTTTCCTTTGGGGCATCATGTTTCATTCTGCT CGCCATCGTGATGGGGCTGGTGTTCATTTGTGTTAAGAACGGGAATATGCGGTGCACTATCTGTATT

2806	ATGAATACGCAGATACTCGTGTTTGCCCTGATCGCCATAATCCCTACGAACGCAGATAAGATCTGCCTCGGACACCACGCTGTTAGCAAC GGAACCAAGGTTAACACTTTGACGGAGAGGGGAGTGGAGGTCGTTAATGCCACCGAAACCGTGGAACGTACCAACATCCCCCGGATCT GTTCCAAAGGTAAAAAGACTGTCGATCTGGGACAGTGCGGCTTACTGGGAACTATTACTGGGCCCCCGCAGTGTGACCAATTCCTGGAA TTCTCTGCAGATCTTATTATTGAGCGGAGGGAGGGGAGCGACGTCTGCTATCCGGGGAAGTTCGTCAACGAGGAGGCCTTGAGACAGAT CCTGACAGAATCTGGGGGCATCGATAAAGAAGCCATGGGATTTACTTATAGCGCCATAAGAACCAATGGCGCCACATCCGCCTGCCGTA GGAGTGGCTCAAGTTTTTATGCAGAAATGAAATGGCTACTCAGCAATACTGACAACGCTGCCTTCCCCCAGATGACAAAGAGCTATAAGA ATACCAGGAAGTCACCAGCCCTGATCGTTTGGGGCATACACCATTCCGTATCTACCGCGGAGCAAACAAAGCTTTATGGCTCTGGGAATA AGCTTGTAACTGTTGGCAGTAGTAATTATCAGCAGAGCTTTGTTCCATCCCCTGGCGCCCGGCCCCAGGTGAATGGTCAAAGCGGCAGG ATAGACTTTCACTGGCTGATGCTCAACCCCAACGATACTGTGACATTCTCCTTTAACGGAGCCTTCATTGCTCCCGACCGCGCCTCTTTCC TCAGGGGCAAAAGTATGGGGATTCAGTCAGGTGTTCAGGTGGACGCAAACTGCGAAGGCGATTGCTACCACTCGGGGGGGACAATCAT ATCCAACCTGCCCTTCCAGAACATTGACAGTCGCGCCGTGGGTAAATGCCCACGGTACGTGAAACAGCGGTCCCTGCTACTGGCTACTG GGATGAAAAACGTCCCTGAGATTCCCAAAGGCCGGGGACTGTTTGGCGCTATCGCGGGGTTCATAGAGAATGGGTGGGAGGGCTTGAT CGACGGGTGGTACGGTTTTAGACATCAGAACGCGCAAGGGGAGGGGACTGCTGCAGATTACAAGTCGACACAGTCCGCAATCGACCAG ATCACCGGTAAATTGAACCGGCTGATAGAAAAGACCAACCAGCAGTTTGAGTTGATAGACAATGAATTCAATGAGGTCGAAAAACAGATC GGTAATGTGATTAATTGGACTCGGGATTCAATTACGGAAGTATGGTCTTACAACGCGGAGTTGTTAGTTGCAATGGAAAACCAGCACACG ATCGATCTTGCAGACTCTGAAATGGATAAATTATACGAGCGGGTCAAACGCCAGCTTCGGGAAAACGCCGAAGAGGATGGTACTGGATG TTTTGAGATCTTTCATAAATGCGACGACGATTGTATGGCCAGCATTCGGAACAATACTTACGACCATTCAAAGTACCGGGAGGAAGCCAT GCAGAATAGAATCCAGATTGATCCAGTGAAACTCAGCTCGGGATACAAAGATGTGATTCTGTGGTTCTCCTTCGGGGCCTCTTGCTTCAT ACTGCTGGCTATCGTAATGGGGCTTGTCTTCATATGTGTGAAAAACGGGAATATGAGATGCACCATTTGTATC
2807	ATGAATACTCAGATACTCGTTTTTGCTTTGATAGCCATCATTCCAACTAACGCTGACAAAATCTGCCTAGGACACCATGCAGTGTCAAATG GCACAAAGGTGAATACTTTGACGGAGAGGGGCGTGGAGGTGGTGAACGCAACAGAGACAGTGGAGCGCACCAACATTCCACGCATTTG TAGTAAAGGAAAGAAGACAGTCGACCTCGGCCAGTGCGGGCTACTGGGTACGATTACTGGACCCCCCCAGTGTGATCAGTTTCTTGAGT TTAGCGCGGATCTGATAATTGAGCGCCGGGAGGGTTCCGATGTCTGTTACCCCGGAAAGTTTGTGAACGAAGAAGCCCTTAGCCAAATC CTGCGCGAGAGCGGCGGAATTGAGAAGGAAGCTATGGGGTTCACGTATAGCGGTATTCGGGCTAACGGAGCCACTAGTGCATGTCGCC GCAGCGGCTCCTCATTCTACGCTGAGATGAAGTGGCTGCTGTCCAATACCGACAATGCTGCTTTTCCGCAGATGACTAAGTCATACAAAA ACACTAGAAAGTCGCCCGCTTTGATTGTTTGGGGCATCCACCACAGCGTGAGTACAGCTGAGCAGACAAAGCTATATGGTTCAGGGAATA AATTAGTTACCGTGGGCAGCTCTAACTACCAGCAGTCCTTTGTGCCCTCTCCCGGCGCTCGGCCTCAGGTAAATGGGCTTAGCGGCCGG ATCGACTTTCATTGGCTGATGCTCAACCCTAATGACACAGTGACGTTCTCTTTCAATGGCGCGTTCATTGCCCCCGACAGGGCTTCTTTCC TACGGGGGAAAAGCATGGGTATTCAGTCAGGTGTTCAAGTAGACGCAAACTGTGAAGGAGACTGTTACCACAGCGGGGGCACAATCATT TCAAATCTTCCTTTCCAGAACATAGATAGCCGGGCAGTCGGTAAATGCCCTCGGTATGTGAAACAAAGATCCCTGTTGCTAGCAACAGGC ATGAAAAACGTCCCAGAGATCCCTAAGGGACGCGGACTTTTCGGCGCGATCGCGGGATTTATCGAGAACGGCTGGGAAGGCCTAATCG ATGGATGGTATGGGTTTAGGCATCAGAATGCTCAGGGGGAGGGAACGGCTGCAGATTATAAATCTACTCAGTCTGCTATCGACCAAATCA CTGGGAAGCTCAATAGGCTGATTGAAAAAACAAATCAACAGTTCGAGCTGATAGACAATGAGTTCAACGAGGTAGAGAAGCAGATAGGTA ACGTAATTAATTGGACTCGCGACTCCATTACAGAAGTGTGGTCCTACAATGCTGAGCTGCTTGTGGCAATGGAAAACCAGCACACCATAG ACCTGGCTGATTCAGAAATGGATAAACTGTATGAGAGGGTGAAGAGGCAGCTCCGGGAAAACGCTGAGGAAGACGGCACAGGTTGCTTC GAAATTTTTCATAAGTGCGATGACGACTGTATGCCTTCAATTCGAAACAATACATACGACCACTCCAAGTATCCAGAGGAAGCCATGCAGA ATCGTATCCAAATTGATCCCGTTAAACTATCATCCGGATATAAAGACGTCATCTTGTGGTTTTCCTTCGGGGCCTCTTGTTTCATCCTCCTC GCAATCGTTATGGGATTGGTGTTTATCTGCGTGAAAAACGGCAACATGCGGTGCACAATATGCATT
2808	ATGAACACCCAGATCTTGGTGTTCGCCCTCATTGCTATTATACCAACTAATGCGGACAAAATCTGCCTTGGACACCATGCCGTATCCAATG GAACCAAGGTCAATACACTGACGGAGAGGGGAGTGGAAGTAGTGAACGCCACGGAAACAGTGGAAAGGACAAATATCCCTAGGATTTGC TCCAAAGGCAAAAGAACAGTGGACCTGGGGCAATGCGGGCTTTTGGGTACTATTACAGGACCACCTCAGTGCGACCAATTCCTTGAGTT CAGCGCTGATCTGATTATAGAGAGGCGCGAGGGGAGTGATGTTTGCTACCCCGGTAAGTTCGTAAATGAGGAGGCTCTGCGGCAGATCT TACGCGAAAGCGGGGGCATTGATAAGGAAGCTATGGGGTTTACCTACAGCGGCATTCGCACTAATGGCGCGACATCTGCCTGTCGCCGT TCCGGGTCATCTTTCTATGCAGAGATGAAATGGCTGCTCTCCAATACTGATAATGCTGCTTTCCCTCAAATGACAAAGTCTTACAAGAATA CAAGGAAGTCTCCCGCACTGATTGTGTGGGGCATCCACCACAGCGTCTCAACCGCTGAGCAGACTAAACTGTATGGTTCTGGGAACAAG CTCGTGACCGTCGGATCGTCTAATTATCAGCAGAGCTTCGTCCCATCACCTGGCGCCAGACCCCAAGTGAACGGTCTGTCCGGCAGAAT TGATTTCCACTGGCTGATGCTGAATCCCAATGACACGGTCACATTCTCATTTAATGGCGCATTCATTGCCCCGGATAGAGCTTCCTTTCTC AGGGGCAAGTCTATGGGAATTCAGAGCGGTGTCCAGGTGGACGCGAATTGCGAAGGGGACTGCTATCACTCCGGCGGCACCATCATTT CTAACCTCCCTTTTCAGAATATTGACAGTCGCGCCGTCGGGAAATGCCCTCGCTACGTGAAGCAGAGATCACTGCTCTTGGCCACAGGAA TGAAGAACGTGCCCGAGATCCCGAAAGGGAGAGGGCTCTTCGGAGCCATCGCTGGGTTCATTGAAAACGGATGGGAAGGCCTGATAGA CGGCTGGTACGGCTTCCGGCACCAAAACGCCCAGGGGGAAGGAACAGCCGCTGACTATAAATCTACACAGAGCGCGATAGACCAGATC ACCGGCAAATTAAATCGTCTCATCGAAAAAACAAATCAGCAGTTTGAGCTTATCGATAATGAATTCAACGAGGTTGAAAAGCAAATCGGGA ATGTTATAAATTGGACCCGAGACTCCATTACCGAGGTGTGGAGCTATAACGCCGAGCTGTTGGTGGCCATGGAGAACCAACATACAATTG ATTTGGCCGACTCCGAGATGGACAAGCTGTATGAGCGGGTTAAACGGCAACTGCGTGAGAATGCAGAAGAAGACGGCACTGGCTGCTTC GAAATCTTCCACAAGTGCGATGATGATTGCATGGCTTCAATACGCAATAACACATACGATCACAGCAAATACAGGGAGGAAGCTATGCAG AATCGCATCCAGATTGATCCAGTGAAGCTGTCATCCGGATACAAAGATGTTATATTATGGTTCTCCTTTGGGGCTTCCTGTTTCATCCTGC TGGCAATCGTGATGGGCCTAGTGTTCATATGCGTCAAGAATGGAAACATGAGATGCACCATCTGTATC

2809	ATGAATACCCAGATACTTGTATTTGCCCTTATCGCCATAATCCCCACTAACGCCGACAAGATTTGTCTAGGGCACCATGCAGTATCTAATG GAACAAAAGTGAATACCCTCACAGAACCGGGGTGTTGAAGTGGTTAATGCCACCGAAACAGTCGAGAGAACTAACATTCCCAGGATTTGTA GCAAGGGTAAGAAGACAGTTGACCTTGGGCAGTGTGGCTTACTTGGAACTATTACCGGCCCACCACAGTGCGATCAGTTTCTGGAGTTC AGCGCAGATCTCATTATTGAGAGAAGAGAAGGAAGCGACGTGTGCTACCCTGGCAAGTTCGTGAATGAGGAGGCCCTTCGGCAGATCCT TAGGGAGTCCGGAGGCATTGATAAGGAGGCTATGGGTTTTACCTATTCTGGCATTCGTACCAACCGCGGCCACAAGTGCTTGTCGCCGCT CCGGTTCTTCCTTTTACGCTGAGATGAAATGGTTGCTCAGCAACACTGACAATGCTGCTTTCCCACAGATGACAAAATCCTACAAAAATAC CAGAAAATCGCCCGCCTTGATTGTGTGGGGGATCCACCATAGTGTGTCTACCGCTGAGCAGACCAAGCTGTATGGTTCCGGGAATAAGC TCGTGACCGTAGGTTCTTCCAATTATCAGCAGAGTTTCGTGCCTTCCCCCGGCGCCAGACCTCAGGTGAATGGCCAGAGCGGGAGAATA GACTTCCATTGGCTCATGCTGAACCCTAACGATACAGTGACCTTCAGTTTCAACGGCGCCTTCATCGCTCCAGACCGCGCCTCATTCCTT CGCGGCAAAAGCATGGGGATCCAGTCCGGGGTTCAAGTAGACGCCAACTGCGAGGGTGACTGCTATCACTCCGGAGGGACGATAATCA GCAATCTTCCCTTCCAGAACATCGATAGCCGCGCCGTCGGTAAGTGCCCACGCTACGTTAAACAGCGCAGCCTGCTCTTGGCTACCGGC ATGAAGAACGTCCCCGAAATCCCTAAGGGCCGGGGCCTGTTCGGGGCCATTGCGGGTTTTATTGAGAATGGCTGGGAGGGGCTGATAG ACGGGTGGTACGGATTTAGACACCAGAATGCTCAGGGGGAAGGGACCGCTGCCGATTATAAGAGCACCCAGTCTGCTATTGACCAGATT ACGGGTAAGCTGAACAGACTTATTGAAAAAACCAACCAGCAGTTTGAGCTTATCGACAATGAATTCAACGAAGTGGAGAAGCAGATCGGG AATGTCATCAACTGGACAAGAGATAGCATCACAGAGGTCTGGTCATATAACGCCGAACTGTTGGTGGCCATGGAAAATCAACACACTATA GATCTAGCCGACTCTGAAATGGATAAGTTGTACGAGCGGGTAAAACGGCAGCTGCGGGAAAACGCTGAGGAAGATGGCACCGGATGCT TCGAGATTTTTCATAAATGCGATGACGACTGTATGGCCTCTATCAGGAATAATACATCACGATCATAGTAAGTACCGGGAAGAAGCCATGCA GAACAGGATCCAGATAGATCCCGTTAAGCTGTCTAGCGGGTACAAGGATGTCATCTTATGGTTCAGCTTCGGCGCCAGCTGCTTCATCCT ATTGGCCATTGTCATGGGACTAGTCTTTATCTGCGTGAAAAACGGTAATATGCGGTGTACTATCTGTATC
2810	ATGAATACCCAGATCCTGGTTTTCGCGCTGATCGCAATCATACCTACTAACGCAGATAAGATTTGCTTGGGACATCATGCCGTGTCCAAC GGCACAAAAGTCAATACCCTCACAGAGAGGGGCGTTGAGGTGGTGAATGCTACCGAGACTGTGGAGAGGACCAATATTCCGAGAATTTG CTCAAAAGGAAAGCGTACAGTCGACCTGGGGCAATGTGGCCTACTCGGCACCATTACAGGGCCCCCCCAGTGTGACCAGTTTCTCGAAT TTTCTGCTGACCTCATCATCGAGAGACGCGAGGGGTCCGACGTGTGTTACCCTGGCAAATTCGTTAATGAAGAAGCCCTAAGACAAATTC TACGCGAGAGCGGAGGTATTGACAAAGAGGCTATGGGATTCACTTATTCTGGTATCCGCACCAACGGCGCTACCTCCGCCTGCCGGCGC TCCGGATCCAGCTTTTATGCTGAGATGAAATGGTTGCTGTCCAACACCGACAATGCTGCATTTCCTCAGATGACCAAATCTTATAAAAACA CAAGAAAGAGTCCTGCGTTGATTGTGTGGGGAATACACCACTCAGTAAGCACTGCCGAACAGACGAAGCTGTATGGTTCCGGAAACAAG CTCGTTACAGTGGGCAGTTCAAATTACCAGCAGAGCTTTGTTCCTTCCCCAGGGGAACGACCCCAAGTTAATGGACTGTCCGGGAGAATT GATTTCCATTGGCTGATGCTGAACCCCAACGACACTGTGACGTTCTCCTTTAATGGGGCATTTATCGCCCCGGATAGAGCTTCTTTTCTGA GAGGCAAGTCAATGGGGATCCAGTCAGGGGTGCAAGTCGATGCCAACTGCGAAGGAGATTGTTATCACTCAGGAGGCACGATTATTAGC AATCTGCCCTTCCAGAATATTGATTCTCGCGCCGTGGCAAATGTCCTCGATACGTGAAGCAGAGATCCCTGCTGCTGGCTACCGGCAT GAAAAATGTACCAGAGATTCCAAAGGGGAGAGGACTGTTCGGTGCCATCGCCGGTTTCATAGAAAATGGGTGGGAGGGCCTGATTGATG GCTGGTACGGTTTTAGGCACCAGAATGCTCAGGGAGAAGGGACCGCCGCAGATTATAAAAGCACTCAGTCTGCAATCGATCAGATAACA GGTAAACTGAACCGCCTGATTGAGAAGACAAACCAGCAGTTTGAACTGATCGATAATGAGTTTAATGAAGTGGAGAAGCAGATTGGCAAC GTTATCAACTGGACACGGGACTCCATCACCGAAGTGTGGAGCTACAACGCTGAGCTGCTGGTAGCCATGGAAAATCAGCACACAATCGA TCTGGCTGACTCAGAAATGGACAAGCTCTATGAGAGGGTCAAAAGGCAACTAAGAGAAAATGCCGAGGAAGATGGGACCGGCTGTTTCG AGATATTTCACAAATGCGACGATCACTGCATGGCTACCATTAGGAATAATACCTACCATCATAGCAAATATCGGGAAGAAGCTATGCAGAA CCGCATCCAAATAGATCCCGTCAAGTTGAGCTCTGGGTACAAGGATGTGATTCTGTGGTTCTCTTTCGGCGCCTCTTGCTTTATTCTGCTG GCAATCGTGATGGGTCTGGTGTTTATTTGTGTCAAAAACGGTAACATGCGCTGCACTATCTGCATA
2811	ATGAATACACAGATTCTAGTGTTTGCCTTGATAGCTATAATCCCAACCAACGCGGACAAGATCTGTCTCGGACACCATGCCGTGTCAAATG GTACTAAGGTGAATACACTGACCGAAAGAGGAGTGGAGGTCGTGAATGCGACAGAAACGGTTGAGCGAACCAACATTCCTAGGATCTGT AGTAAGGGAAAGCGAACTGTCGATCTGGGCCAGTGTGGACTCCTCGGGACCATTACAGGGCCTCCTCAGTGTGATCAGTTTTTGGAATT TTCCGCAGACTTGATAATCGAGAGGCGGGAAGGGTCTGACGTGTGCTACCCAGGCAAATTTGTAAATGAAGAGGCCCTAAGGCAGATCC TGAGGGAGAGTGGTGGAATCGATAAAGAAGCCATGGGATTTACATATTCAGGCATCAGGACCAACGGCGCAACATCAGCCTGCCGAAGA TCGGGATCTTCATTCTACGCCGAGATGAAATGGTTGCTATCCAACACTGACAATGCCGCATTCCCCCAGATGACTAAAAGCTACAAAAAC ACCCGGAAGTCCCCAGCTCTCATCGTCTGGGGGATCCATCACAGTGTGTCCACCGCCGAGCAGACCAAGCTGTATGGATCCGGCAGTAA ACTGGTTACTGTGGGGTCATCTAATTACCAACAGAGTTTTGTGCCATCGCCTGGCGCCAGACCTCAAGTGAATGGACTGTCCGGTAGAAT TGATTTTCACTGGCTCATGCTGAACCCTAATGACACGGTTACCTTTTCTTTCAACGGAGCATTTATAGCTCCGGATCGCGCCAGTTTTCTA CGGGGGAAAAGCATGGGAATCCAGTCGGGTGTGCAGGTAGATGCTAACTGCGAAGGTGATTGCTACCACTCCGGTGGTACAATCATCAG CAATCTGCCGTTCCAGAACATAGATAGTCGTGCAGTGGGTAAGTGCCCTAGGTACGTGAAACAAAGATCTCTGCTGCTTGCAACCGGTAT GAAGAACGTGCCTGAGATTCCGAAAGGGCGCGGCTTATTCGGAGCTATAGCTGGATTCATCGAGAATGGATGGGAGGGCCTGATTGATG GTTGGTACGGGTTTCGACATCAGAACGCCCAGGGAGAGGGTACCGCTGCTGACTACAAGAGCACCCAATCCGCTATCGACCAGATAACT GGCAAGCTGAATAGACTGATTGAAAAGACAAATCAGCAATTTGAATTGATCGACAACGAATTCAACGAAGTGGAAAAGCAGATTGGTAAC GTGATCAACTGGACCAGAGACTCTATCACAGAGGTATGGTCTTACAATGCCGAGCTCCTGGTTGCCATGGAAAACCAACATACCATTGAC TTCGCCCGACAGCGAAATGGATAAACTATACGAGCGGGTCAAACGACAGCTGCGCGAGAATGCCGAGGAGGATGGAACTGGCTGCTTCG AAATCTTCCACAAGTGTGACGACGATTGTATGGCCAGCATCCGGAACAATACATATGATCATTCAAAATATCGCGAGGAGGCTATGCAGA ATAGAATTCAGATTGACCCCGTGAAGCTGAGCAGCGGGTACAAGGACGTCATTCTGTGGTTCAGTTTCGGAGCCTCTTGTTTTATCCTCC TCGCGATCGTGATGGGATTAGTGTTTATTTGTGTGAAGAACGGCAATATGAGGTGTACGATTTGTATC

2812	ATGAACACGCAGATCTTGGTGTTCGCACTAATCGCGATAATCCCCACTAATGCAGACAAAATTTGTTTAGGCCATCATGCTGTATCCAATG GCACAAAAGTCAACACCCTGACGGAAAGAGGTGTGGAAGTCGTCAACGCTACCGAGACTGTGGAACGGACGAATATCCCAAGGATCTGT TCAAAGGGGAAGCGAACGGTCGACCTGGGACAGTGCGGCCTGCTGGGGACCATCACAGGACCGCCACAGTGCGACCAGTTTCTGGAG TTTTCTGCGGACTTGATTATCGAAAGACGGGAGGGAAGTGACGTATGCTATCCCGGCAAATTCGTGAATGAAGAGGCCTTGCGCCAGAT ATTACGCGAGTCCGGAGGGATCGACAAGGAAGCTATGGGGTTCACATACTCCGGCATCAGAACCAACGGCGCCACATCCGCATGCCGG CGCAGTGGATCCAGCTTCTACGCAGAAATGAAATGGCTCCTCAGCAATACGGATAACGCCGCCTTCCCACAAATGACCAAGTCATACAAA AATACCAGAAAGAGCCCAGCCTTGATAGTGTGGGGAATCCACCACTCTGTGTCTACCGCCGAACAGACTAAGCTGTATGGATCTGGCTC GAAATTAGTCACCGTCGGCTCCTCCAACTATCAGCAATCATTCGTGCCAAGCCCCGGAGCCCGCCCACAGGTTAATGGACTGTCAGGAA GGATTGACTTCCACTGGCTGATGCTGAATCCTAATGATACTGTGACTTTCTCATTTAATGGCGCTTTTATTGCCCCTGACAGAGCCAGCTT CCTTCGAGGGAAGTCCATGGGAATCCAGTCTGGCGTGCAAGTAGACGCAAACTGTGAAGGGGATTGCTATCATAGTGGTGGGACAATCA TTTCAAATCTGCCCTTCCAGAATATCGATAGTAGAGCAGTAGGAAAATGTCCCAGGTACGTGAAACAGAGATCACTGTTATTGGCTACAG GAATGAAAAATGTCCCAGAAATCCCGAAGGGGAGAGGCCTCTTTGGGGCCATAGCAGGCTTTATTGAGAATGGGTGGGAAGGCCTAATC GATGGATGGTACGGGTTCCGCCATCAGAATGCGCAGGGCGAGGGGACTGCAGCCGACTACAAAAGCACGCAATCCGCTATCGACCAGA TTACTGGAAAGCTCAACCGATTAATTGAAAAGACTAATCAGCAGTTCGAACTGATCGATAACGAGTTCAACGAGGTCGAAAAGCAGATCG GGAATGTGATCAACTGGACCAGGGATTCCATAACCGAGGTTTGGAGCTATAATGCAGAGTTGCTTGTAGCAATGGAGAACCAACATACCA TAGACCTGGCCGATTCGGAAATGGACAAGCTGTATGAGCGAGTGAAACGGCAGCTGCGGGAGAATGCAGAAGAAGACGGCACAGGTTG TTTTGAAATTTTTCACAAATGCGATGACGATTGCATGGCATCGATCAGAAATAACACCTATGATCATTCAAAGTATCGCGAGGAGGCCATG CAGAATAGGATCCAAATCGACCCTGTCAAACTCTCTAGTGGCTACAAGGATGTCATCCTCTGGTTCTCCTTTGGTGCCTCTTGCTTCATTT TGCTCGCCATTGTAATGGGATTGGTGTTCATTTGTGTTAAATCCAGAAATATGCGTTGTACCATTTGTATC
2813	ATGAACACCCAGATTCTGGTCTTCGCTCTGATCGCCATTATACCAACAAATGCGGATAAAATTTGCCTGGGACACCATGCTGTAAGCAAC GGCACGAAGGTGAACACACTCACCGAAAGGGGAGTCGAAGTAGTCAATGCCACAGAGACCGTTGAACGTACGAACATCCCTAGAATTTG TAGCAAAGGGAAACGGACTGTCGATTTGGGACAATGTGGACTGCTTGGGACTATCACAGGGCCCCCTCAGTGCGATCAGTTCCTGGAAT TCTCGGCCGATCTGATCATTGAACGGAGAGAAGGAAGCGACGTGTGCTACCCAGGTAAATTCGTGAACGAAGAGGCACTGCGCCAAATT CTTCGGGAGTCCGGCGGGATCGACAAGGAGGCAATGGGATTTACGTATTCTGGCATTAGGACCAACGGGGCTACATCTGCCTGCCGGC GAAGTGGATCCTCCTTCTATGCAGAAATGAAGTGGCTCCTTTCAAACACAGACAATGCCGCCTTTCCACAGATGACCAAGAGTTATAAGA ATACCAGGAAAAGTCCTGCCTTGATTGTGTGGGGGATTCACCACTCCGTGAGCACTGCCGAGCAGACCAAGCTTTACGGGAGCGGCAAT AAGCTGGTGACAGTGGGCTCTTCCAATTATCAACAATCCTTCGTGCCTAGCCCTGGGGAAAGACCCCAGGTGAATGGACTGTCCGGCCG GATCGACTTTCACTGGCTCATGCTGAATCCCAACGACACTGTAACCTTTAGCTTTAACGGCGCCTTCATCGCTCCCGATAGGGCGTCTTT CCTACGGGGGAAAAGTATGGGAATTCAATCAGGCGTCCAGGTAGACGCCAACTGCGAGGGTGACTGTTACCACAGCGGAGGTACAATAA TTTCCAACCTGCCTTTCCAAAACATAGATTCCCGCGCAGTGGGGAAATGCCCCAGGTACGTGAAGCAGCGAAGTCTCCTCCTTGCCACA GGAATGAAGAACGTGCCTGAGATTCCAAAAGGGCGGGGGTTGTTCGGGGCCATCGCTGGATTTATCGAAAACGGGTGGGAAGGCCTTA TCGACGGCTGGTACGGTTTTAGGCACCAGAACGCACAGGGTGAGGGGACTGCGGCCGACTACAAGTCGACACAAAGCGCTATAGACCA GATCACCGGGAAACTTAATCGGCTGATCGAAAAGACTAACCAGCAGTTTGAATTAATCGACAACGAATTTAATGAGGTGGAGAAGCAGAT CGGAAACGTGATTAATTGGACCAGGGATTCCATCACGGAAGTTTGGTCCTATAACGCGGAGTTGCTTGTTGCCATGGAAAACCAGCACAC CATTGACCTCGCTGACTCTGAGATGGACAAACTATATGAGAGAGTCAAAAGACAGCTGCGCGAAAATGCAGAGGAGGATGGCACGGGTT GTTTCGAAATCTTTCACAAATGTGATGACGATTGTATGGCCAGTATACGGAATAACACCTATCACCACTCAAAGTACCGAGAAGAAGCCAT GCAGAATCGCATCCAGATCGATCCTGTTAAGCTGTCCTCAGGATATAAAGACGTCATCCTGTGGTTCTCTTTTGGCGCAAGCTGCTTTATC CTTTTAGCGATCGTGATGGGCCTGGTTTTTATATGTGTCAAGAATGGAAATATGCGGTGTACAATTTGCATC
2814	ATGAACACTCAGATCTTAGTGTTTGCACTGATTGCCATAATCCCAACAAACGCCGATAAAATCTGCCTGGGCCATCACGCGGTGTCGAAT GGCACAAAGGTCAACACACTGACCGAAAGAGGCGTAGAGGTCGTGAACGCGACGGAGACCGTGGAGCGCACTAACATTCCTCGCATCT GTTCAAAAGGTAAGAAAACAGTGGATTTGGGACAGTGTGGCCTTCTCGGTACCATTACCGGCCCTCCCCAGTGCGACCAGTTCCTCGAA TTTTCTGCTGATTTAATTATTGAACGGCGGGAGGGGTCAGACGTGTGTTATCCCGGCAAGTTTGTGAACGAAGAAGCTCTTAGACAGATC CTCCGGGAGAGCGGGGGGATTGACAAGGAAGCAATGGGCTTTACATACTCAGGTATTCGAACAAATGGAGCCACTAGCGCCTGCCGAC GATCCGGGTCATCTTTCTACGCAGAGATGAAATGGCTGCTCAGCAACACTGATAATGCTGCCTTTCCACAGATGACAAAATCCTATAAAAA CACCAGAAAATCCCCTGCGTTAATTGTCTGGGGCATCCACCATTCGGTTTCTACAGCAGAGCAGACCAAGCTGTACGGCAGCGGAAATA AACTCGTGACTGTAGGATCGAGTAATTATCAGCAAAGCTTTGTTCCATCACCAGGGGCCCGTCCACAGGTCAATGGACAGAGTGGAAGG ATCGATTTCCATTGGCTAATGCTGAACCCAAACGATACTGTGACCTTTTCTTTCAACGGGGCCTTCATTGCTCCTGATCGGGCATCATTTT TGCGTGGGAAGTCAATGGGTATACAATCAGGGGTGCAAGTGGACGCTAATTGCGAAGGTGACTGTTACCATTCTGGAGGCACCATTATC TCTAACTTGCCTTTCCAAAACATTGACAGCAGAGCCGTGGGTAAGTGCCCTCGATACGTGAAGCAGCGGTCGCTTCTCCTTGCCACGGG CATGAAGAACGTGCCTGAAATCCCTAAAGGTCGTGGTCTATTCGGAGCAATTGCCGGTTTCATCGAGAATGGGTGGGAAGGCCTTATCG ATGGTTGGTACGGGTTTCGGCACCAGAATGCCCAGGGCGAGGGCACAGCTGCTGACTACAAGTCGACCCAGAGCGCTATCGACCAGAT CACAGGCAAGCTCAACAGGCTGATCGAAAAGACTAACCAGCAATTCGAATTGATCGATAACGAGTTTAATGAGGTGGAAAAGCAGATTGG GAATGTTATCAACTGGACCCGCGATTCCATTACTGAGGTATGGTCCTACAACGCTGAGCTCTTAGTCGCTATGGAGAACCAGCACACCAT AGATCTGGCTGATTCAGAGATGGATAAGCTCTATGAACGGGTTAAAAGGCAGCTTCGGGAGAACGCCGAGGAAGACGGAACTGGTTGCT TCGAGATCTTTCACAAATGTGACGATGACTGCATGGCTTCCATTAGAAATAACACGTATGACCATAGCAAGTATAGAGAGGAAGCAATGCA AAACCGGATCCAGATCGACCCAGTCAAATTGAGCTCGGGATACAAGGACGTTATCCTGTGGTTCTCATTTGGCGCCTCCTGTTTCATATT GTTGGCTATTGTGATGGGTCTGGTGTTCATCTGTGTCAAGAATGGGAACATGAGGTGCACCATTTGTATA

2815	ATGAACACACAGATTCTCGTGTTTGCGCTAATTGCAATCATACCCACTAATGCAGACAAAATCTGTCTGGGACACCACGCGGTTAGCAAT GGCACGAAGGTGAATACCCTGACAGAGAGAGGAGTGGAGGTTGTTAACGCGACCGAAACTGTTGAAAGGACTAATATCCCTCGTATCTG TTCAAAGGGAAAAAAGACCGTAGACCTGGGCCAGTGTGGACTGCTCGGCACGATCACAGGGCCACCTCAGTGTGACCAATTTCTGGAGT TCAGTGCTGACCTGATTATCGAAAGAAGAGAGGGATCAGACGTATGTTATCCCGGAAAGTTCGTGAATGAGGAAGCCCTTCGACAGATCT TAAGAGAATCAGGCGGGATCGATAAGGAAGCAATGGGCTTCACCTACTCTGGCATTCGGACCAATGGCGCAACTTCTGCATGCCGCAGA TCCGGCTCTAGTTTTTACGCGGAGATGAAGTGGCTCCTCTCCAATACAGACAATGCAGCATTCCCACAAATGACCAAGTCTTATAAAAACA CCCGCAAGTCCCCCGCTCTCATCGTTTGGGGTATCCACCATAGTGTGTCCACAGCCGAGCAAACAAAGCTATACGGAAGTGGCAACAAG CTGGTCACAGTGGGCAGTAGTAATTATCAGCAGAGCTTCGTGCCTAGCCCCGGGGCCAGGCCACAGGTTAACGGCCAGTCAGGGCGCA TCGACTTTCACTGGCTTATGCTGAACCCGAACGACACTGTCACATTCTCATTCAACGGCGCCTTCATAGCACCCGATAGGGCCTCATTTTT AAGGGGGAAGTCTATGGGAATTCAGTCCGGAGTGCAGGTCGACGCCAACTGTGAGGGCGACTGCTACCATAGTGGAGGAACCATCATA AGTAATCTGCCATTTCAAAACATCGATTCACGCGCGGTGGGAAAATGCCCCAGGTATGTCAAGCAGCGGAGCCTTTTGCTTGCTACCGG GATGAAGAATGTGCCCGAGATTCCCAAGGGGAGAGGTTTATTCGGGGCTATAGCCGGGTTTATTGAAAACGGTTGGGAGGGGCTAATAG ACGGATGGTACGGCTTCAGGCACCAGAATGCTCAAGGGGAAGGGACTGCCGCTGATTATAAGAGCACGCAGTCAGCTATCGACCAGAT CACTGGGAAATTGAATCGTTTAATCGAGAAAACCAATCAGCAGTTCGAACTGATTGATAACGAGTTCAACGAGGTCGAGAAACAAATAGG AAACGTTATAAACTGGACACGAGATTCCATAACCGAGGTGTGGTCCTATAATGCTGAACTACTCGTAGCCATGGAGAACCAGCATACCAT TGATCTGGCCGACTCTGAGATGGACAAGCTATATGAACGGGTGAAAAGACAGTTGCGCGAAAACGCGGAAGAAGACGGGACAGGTTGC TTTGAAATTTTTCACAAATGCGATGACGACTGCATGGCTAGTATACGGAACAACACATACGATCACTCTAAGTATCGCGAGGAGGCAATG CAGAATCGCATCCAGATCGATCCCGTCAAACTGTCTTCCGGATATAAAGATGTCATACTGTGGTTCTCTTTCGGCGCAAGCTGTTTCATTC TCCTAGCCATCGTCATGGGTCTGGTTTTCATATGTGTGAAGAACGGGAACATGCGGTGCACCATATGTATA
2816	ATGAATACGCAGATCCTGGTGTTTGCTTTAATAGCCATCATCCCTACTAACGCAGACAAAATTTGTCTCGGGCACCACGCCGTGTCTAATG GAACAAAGGTCAACACTCTGACAGAACGCGGAGTTGAAGTAGTGAATGCCACTGAAACTGTGGAGAGGACCAATATCCCTCGCATTTGTT CTAAGGGAAAGAAGACCGTCGACCTCGGTCAGTGCCGCCTATTGGGAACCATCACTGGCCCTCCTCAGTGTGACCAGTTTCTGGAGTTT AGTGCCGATTTGATCATCGAAAGAACAGGGAAGCGACGTTTGTTATCCCGGAAAGTTTGTCAACGAGGAGGCTCTCCGCCAAATTCTC AGGGAATCTGGAGGTATCGACAAGGAAGCAATGGGCTTCACTTATTCAGGCATTCGCACAAACGGGGCCACCTCTGCATGCCGCCGCAG CGGCTCTAGCTTTTACGCCGAAATGAAATGGCTGCTCTCCAATACTGACAACGCTGCCTTTCCCCAAATGACCAAGTCCTATAAAAACACG CGCAAAAGTCCGGCCCTCATTGTGTGGGGGATTCACCACTCCGTCTCCACAGCCGAGCAGACTAAGCTGTACGGTAGTGGTAATAAGTT GGTCACCGTGGGGTCAAGTAATTATCAGCAAAGCTTCGTCCCTAGCCCCGGGGCCAGACCCCAGGTGAACGGACAAAGCGGAAGGATC GATTTTCACTGGCTGATGCTTAACCCTAATGACACCGTTACATTCAGTTTTAATGGTGCGTTTATCGCCCCCGATCGGGCCTCCTTTCTTA GAGGCAAAAGTATGGGTATCCAGAGTGGAGTACAGGTCGATGCCAATTGCGAAGGCGACTGCTATCACTCTGGGGGAACAATTATCTCC AATTTACCGTTCCAGAACATCGACTCTAGGGCAGTCGGCAAATGCCCTAGATATGTAAAGCAGAGGTCTCTGCTGCTCGCCACGGGAAT GAAAAACGTCCCAGAGATCCCAAAAGGCAGAGGACTGTTCGGAGCCATAGCTGGCTTTATCGAGAACGGCTGGGAGGGACTGATCGAT GGATGGTATGGTTTCCGGCACCAGAATGCACAAGGTGAGGGAACTGCCGCAGATTATAAGTCCACCCAAAGTGCCATCGATCAGATTAC AGGCAAGTTAAACAGACTTATTGAAAAAACAAATCAACAGTTTGAGCTTATTGACAACGAGTTCAATGAAGTTGAGAAGCAAATCGGTAAC GTTATCAATTGGACTCGCGACTCCATCACCGAGGTGTGGTCTTACAACGCGGAACTACTTGTAGCGATGGAAAATCAGCATACCATCGAC TTGGCCGACTCGGAGATGGATAAGTTGTACGAGAGAGTGAAACGGCAGCTCAGGGAAAACGCGGAGGAGGACGGAACAGGTTGTTTCG AAATATTTCACAAGTGCGATGATGACTGTATGGCTAGCATTCGGAACAACACTTACGATCACTCCAAGTACCGTGAAGAGCCTATGCAGA ACAGGATTCAGATCGATCCGGTGAAACTCTCAAGCGGATACAAGGACGTTATACTTTGGTTCAGCTTCGGAGCTTCTTGCTTTATTCTTTT GGCTATCGTTATGGGACTAGTGTTTATATGTGTGAAAAACGGCAACATGCGCTGTACGATATGCATA
2817	ATGAACACACAGATCCTTGTGTTTGCACTGATTGCCATCATTCCCACCAATGCCGACAAAATCTGCTTAGGTCACCACGCTGTTAGCAACG GAACGAAGGTCAACACCCTGACCGAACGAGGCGTCGAAGTAGTTAATGCTACCGAAACTGTGGAGAGGACCAACATACCCAGGATCTGC AGTAAGGGTAAGAAGACAGTCGATCTAGGCCAGTGCGGACTACTCCGCCACAATTACCGGACCTCCCCAGTGCGACCAGTTTCTTGAGTT CAGTGCTGATCTTATTATAGAACGACGAGAAGGGAGTGATGTGTGCTATCCCGGCAAGTTTGTGAACGAAGAGGCGCTGCGACAAATCTT GAGGGAGTCAGGCGGGATTGACAAGGAAGCAATGGGGTTTACATATTCAGGTATTCGTACTAATGGAGCAACATCTGCCTGCCGGCGCA GTGGCTCCAGCTTTTACGCGGAAATGAAATGGCTGTTAAGTAACACCGACAATGCAGCATTTCCCCAAATGACAAAGAGTTATAAGAATAC ACGAAAGTCACCCGCACTGATTGTATGGGGCATCCACCATTCGGTGTCGACCGCTGAACAGACAAAATTGTACGGCTCAGGCAACAAAC TGGTGACTGTTGGTTCTTCTAACTACCAGCAGAGCTTTGTCCCGTCACCCGGAGCCCGACCACAGGTGAATGGACAGTCCGGACGGATT GACTTTCACTGGCTTATGCTCAATCCCAATGATACTGTGACATTTAGTTTCAACGGGGCCTTTATAGCTCCCGATCGGGCTTCTTTTCTTC GAGGAAAATCAATGGGCATTCAATCCGGAGTTCAGGTGGACGCCAATTGTGAAGGAGACTGTTACCATTCCGGAGGTACAATTATTTCCA ACCTGCCTTTCCAAAATATCGATAGTCGAGCCGTGGGAAAGTGTCCCAGATATGTAAAACAGAGAAGTCTATTGCTGGCTACAGGGATGA AAAATGTGCCCGAAATACCCAAGGGACGGGGATTGTTTGGTGCCATAGCTGGGTTTATCGAGAATGGGTGGGAAGGCCTTATCGACGGC TGGTATGGCTTTCGGCACCAAAACGCGCAGGGTGAGGCGACCGCGGCAGACTACAAGTCTACTCAGAGCGCAATTGATCAGATTACCG GCAAACTGAACCGACTAATAGAAAAAACAAACCAGCAGTTTGAGCTCATTGATAATGAATTCAACGAAGTGGAGAAGCAGATTGGGAACG TGATCAATTGGACGCGCGATTCCATCACTGAGGTGTGGAGTTACAATGCCGAACTCCTGGTGGCCATGGAAAACCAGCACACAATTGAC CTCGCTGACTCTGAGATGGACAAGTTGTATGAGCGCGTGAAGCGGCAGCTTAGAGAGAACGCCGAGGAAGACGGGACAGGGTGTTTTG AGATCTTCCATAAGTGCGATGACGACTGCATGGCATCCATCCGTAACAACACATATGATCACTCAAAGTATAGGGAGGAAGCCATGCAGA ACAGGATTCAGATCGACCCTGTGAAACTCTCATCAGGGTACAAAGATGTAATCTTGTGGTTTTCGTTTGGAGCTTCGTGCTTCATTCTGCT AGCGATTGTCATGGGCCTTGTGTTTATCTGCGTCAAGAACGGCAACATGAGGTGTACTATCTGTATA

2818	ATGAATACCCAGATTCTGGTGTTTGCGCTGATTGCCATAATCCCCACTAACGCTGATAAGATCTGCCTGGGGCACCATGCAGTGTCTAAT GGCACTAAGGTCAATACGCTCACGGAGAGAGGCGTGGAAGTGGTTAATGCTACCGAAACTGTGGAACGGACTAACATTCCAAGAATCTG TTCAAAGGGTAAGAAAACTGTGGACCTCGGCCAATGTGGCTTGCTGGGCACCATCACTGGGCCACCACAGTGTGATCAGTTTCTCGAGT TCTCTGCCGATTTAATCATCGAGCGCCGGGAAGGATCAGATGTGTGCTACCCGGGTAAATTCGTGAATGAGGAAGCACTGAGACAGATT CTGCGTGAATCAGGAGGGATTGATAAGGAGGCTATGGGGTTTACATACTCTGGCATCAGAACAAATGGGGCCACCAGCGCGTGCCGGC GCTCTGGGAGTTCTTTCTACGCTGAAATGAAATGGCTGCTTTCTAACACGGACAATGCCGCTTTCCCCCAGATGACAAAAAGCTACAAGA ACACTCGCAAGAGCCCTGCACTCATAGTGTGGGGCATCCACCACTCCGTCTCTACTGCAGAGCAGACCAAGCTGTACGGCAGCGGGAA CAAGCTTGTAACCGTTGGATCTTCAAACTACCAACAATCATTCGTTCCCTCTCCTGGAGCCCGGCCCCAAGTTAATGGGCAGTCCGGGAG AATCGACTTCCATTGGCTGATGCTAAATCCCAATGATACAGTTACCTTTAGCTTTAACGGCGCCTTCATCGCTCCGGATAGAGCGAGCTTC CTTCGCGGTAAGTCTATGGGAATCCAGTCCGGGGTGCAAGTCGACGCAAACTGCGAGGGAGACTGTTACCATTCCGGGGGTACTATCAT TTCAAATTTACCCTTTCAGAACATTGATAGCCGCGCCGTAGGCAAATGTCCTAGGTATGTGAAACAGAGAAGCCTGCTCCTGGCAACCGG AATGAAGAACGTGCCTGAGATCCCTAAGGGTAGAGGACTTTTCGGAGCTATCGCCGGGTTCATCGAGAACGGTTGGGAGGGACTCATTG ATGGGTGGTATGGTTTCCGCCACCAGAACGCCCAAGGGGAAGGCACTGCTGCTGACTATAAATCAACGCAGTCCGCCATAGACCAGATT ACAGGTAAATTAAATAGACTGATTGAGAAAACAAACCAGCAATTCGAGCTAATCGACAATGAGTTTAATGAGGTCGAGAAGCAGATCGGTA ACGTCATAAATTGGACACGGGATAGTATCACCGAAGTGTGGAGCTACAATGCTGAGCTCCTAGTTGCCATGGAGAATCAACATACGATCG ACCTGGCGGATTCTGAAATGGACAAGTTATACGAAAGAGTGAAGCGACAGCTAAGAGAGAATGCAGAGGAAGACGGCACTGGCTGCTTC GAAATATTTCACAAGTGTGATGATGACTGTATGGCAAGCATCAGGAATAACACCTACGACCACAGTAAGTATAGAGAGGAGGCAATGCAG AACAGGATACAGATCGATCCCGTCAAACTTTCCAGTGGCTATAAGGACGTTATCCTTTGGTTCTCGTTTGGGGCGAGCTGCTTCATCTTG CTGGCGATCGTGATGGGCCTCGTGTTTATTTGCGTGAAAAACGGAAACATGAGGTGTACAATTTGTATT
2819	ATGAATACACAAATTCTGGTCTTCGCACTGATCGCCATAATCCCCACTAACGCAGACAAGATCTGTCTCGGTCACCACGCCGTTTCGAAT GGTACTAAGGTAAATACGCTGACAGAGAGGGGGGTGGAAGTCGTGAACGCCACTGAGACCGTCGAAAGGACAAACATTCCCCGCATCT GTTCGAAGGGAAAGAAGACCGTCGATCTTGGGCAGTGCGGTCTCTTGGGTACGATCACAGGGCCTCCTCAGTGCGATCAATTCCTGGAG TTTTCAGCCGACCTGATAATCGAAAGAAGGGAGGGAAGCGATGTGTGTTACCCCGGGAAGTTCGTGAATGAGGAAGCCCTGCGACAGAT CCTAAGGGAGAGCGGGGGAATCGATAAAGAGGCAATGGGGTTCACCTATAGTGGAATCAGAACTAACGGCGCCACTTCCGCCTGCCGG AGATCGGGTAGCTCTTTCTATGCTGAAATGAAATGGCTGCTCAGTAATACAGACAACGCGGCGTTCCCACAGATGACCAAGTCCTATAAG AATACCCGGAAATCTCCTGCGCTTATTGTGTGGGGCATCCACCACTCTGTGTCCACCGCTGAGCAGACAAAACTTTACGGATCAGGAAAT AAGCTCGTCACCGTGGGAAGCTCCAATTATCAGCAATCCTTCGTGCCATCCCCCGGAGCAAGACCACAGGTGAATGGACAGTCCGGGAG GATCGATTTCCACTGGTTGATGCTGAACCCCAATGATACTGTCACGTTCAGCTTCAACGGTGCTTTTATTGCTCCCGATAGGGCCAGCTTT CTGAGAGGCAAGTCCATGGGCATCCAGAGCGGGGTCCAAGTGGACGCTAATTGCGAGGGTGATTGCTACCACTCTGGAGGCACCATCA TCAGCAACCTGCCTTTCCAGAATATAGATTCCCGGGCGGTGGGGAAGTGCCCAAGATACGTCAAGCAACGGAGCCTCCTCCTTGCGACT GGAATGAAAAACGTGCCAGAGATTCCAAAGGGGAGGGGCCTGTTTGGAGCGATTGCCGGATTCATCGAGAATGGCTGGGAAGGGCTCA TTGACGGATGGTATGGCTTTCGCCATCAGAACGCCCAGGGAGAGGGGACCGCGGCCGACTACAAGTCCACCCAAAGCGCCATTGACCA AATCACAGGAAAGCTGAATCGCCTAATCGAAAAGACTAATCAGCAGTTCGAGCTGATAGATAATGAGTTCAACGAGGTCGAGAAACAAAT CGGAAATGTCATCAATTGGACCCGCGACTCTATAACAGAAGTTTGGTCCTATAACGCGGAGCTGCTAGTCGCCATGGAAAACCAGCACAC TATTGACCTTGCCGATAGCGAAATGGACAAGCTGTATGAAAGGGTGAAAAGACAGCTGCGCGAGAATGCAGAAGAGGACGGTACCGGCT GTTTCGAGATTTTCCATAAGTGTGACGACGACTGCATGGCATCGATTCGAAACAACACATACGACCACTCTAAGTATACAGAGGAGGCCA TGCAGAACAGAATCCAGATCGATCCTGTGAAGCTGTCATCCGGGTACAAGGATGTCATTCTTTGGTTCTCATTCGGTGCGTCTTGTTTCAT CCTGCTCGCCATAGTCATGGGGCTGGTCTTCATTTGCGTGAAAAACGGGAACATGCGGTGTACAATTTGCATC
2820	ATGAATACCCAGATACTCGTGTTTGCCCTTATCGCTATAATCCCCACGAATGCAGACAAAATCTGCCTTGGGCATCACGCCGTGAGTAAT GGCACCAAAGTGAACACATTGACCGAGCGGGGGGTGGAGGTAGTCAATGCCACCGAAACAGTGGAACGCACCAACATTCCTCGGATCT GTAGCAAAGGAAAGAAAACAGTCGATCTGGGCCAGTGTGGCCTGTTAGGAACCATCACTGGACCGCCTCAATGCGATCAGTTTCTCGAA TTCAGCGCGGATCTCATTATTGAAAGAAGAGAGGGTTCTGACGTGTGTTACCCCGGCAAGTTTGTGAACGAAGAAGCACTCCGCCAGATA CTTAGGGAGTCCGGCGGGATCGACAAAGAGGCTATGGGGTTTACTTATAGTGGAATCCGCACGAACGGGGCCACGTCCGCCTGCAGAC GGAGCGGCTCCTCATTTTACGCGGAGATGAAGTGGCTGCTCAGCAACACAGACAATGCCGCCTTTCCCCAGATGACAAAGTCTTACAAG AACACCAGAAAGTCGCCAGCCCTCATAGTCTGGGGGATACATCACAGCGTTTCCACCGCCGAGCAGACTAAGCTGTATGGCAGCGGAAA TAAACTGGTGACTGTGGGCTCCAGCAATTACCAGCAAAGCTTTGTACCCAGCCCCGGAGCAAGGCCCCAAGTGAATGGCCAGTCTGGTA GAATCGATTTCCATTGGCTTATGTTGAACCCTAACGACACGGTTACATTCAGTTTCAATGGGGCATTTATCGCCCCCGACCGTGCCTCTTT CTTGAGGGGTAAATCTATGGGCATCCAGTCAGGTGTGCAGGTGGATGCTAATTGCGAAGGCGACTGCTATCACAGTGGCGGTACTATCA TATCCAATCTGCCTTTTCAAAACATCGATTCCCGTGCGGTGGGGAAGTGTCCTCGATACGTGAAACAGCGCTCATTGCTGTTGGCCACCG GAATGAAAAACGTGCCAGAGATCCCAAAGGGCAGAGGTCTGTTCGGAGCTATTGCTGGCTTTATCGAAAACGGCTGGGAAGGGCTGATT GACGGCTGGTATGGATTTAGGCACCAAAATGCTCAAGGCGAGGGCACAGCCGCCGACTATAAAAGCACTCAAAGTGCGATAGACCAGAT CACAGGCAAGCTGAACCGCCTGATTGAGAAAACAAACCAGCAGTTCGAACTCATTGACAACGAATTCAACGAAGTGGAGAAACAGATCG GAAATGTTATCAATTGGACAAGAGACAGTATTACAGAGGTGTGGTCCTATAACGCCGAGCTCCTGGTCGCTATGGAGAATCAGCATACCA TCGACCTGGCGGATTCAGAGATGGACAAACTTTATGAGAGAGTTAAAAGACAACTGCGTGAGAATGCTGAAGAAGATGCAACTGGATGCT TCGAAATTTTTCATAAGTGTGATGATGACTGTATGGCGTCAATCCGGAATAACACCTACGATCACTCCAAATATAGAGAAGAGGCAATGCA GAATCGTATTCAGATAGACCCTGTGAAATTGTCCAGTGGATACAAGGACGTAATTTTGTGGTTCAGCTTCGGGGCTTCCTGTTTCATCCTC CTCGCCATCGTGATGGGTCTCGTTTTCATATGCGTGAAGAACGGGAATATGAGGTGCACGATATGTATC

2821	ATGAACACCCAGATTCTGGTGTTCGCTCTCATCGCAATCATACCCACCAATGCTGACAAAATCTGTTTGGGGCATCATGCCGTCAGCAAC GGCACTAAGGTCAATACTCTTACAGAACGCGGGGTTGAGGTTGTCAATGCAACTGAAACCGTGGAAAGAACGAATATCCCACGTATTTGC AGTAAGGGCAAGAAAACCGTGGACCTCGGGCAATGCGGGCTGCTTGGAACGATTACCGGCCCACCCCAATGCGATCAGTTCCTCGAGT TTTCTGCAGATTTGATTATTGAGCGAAGGGAGGGCTCCGACGTCTGTTATCCTGGAAAGTTTGTGAATGAGGAGGCACTGCGCCAGATAC TCCGTGAAAGCGGCGGGATTGATAAGGAGGCCATGGGCTTTACTTACTCGGGAATACGCACAAATGGAGCCACTAGCGCCTGCCGGAG GTCTGGATCAAGCTTCTACGCAGAAATGAAGTGGTTACTAAGCAACACAGATAACGCAGCTTTCCCACAGATGACAAAAAGCTATAAGAAT ACTCGCAAGTCCCCTGCCCTGATAGTATGGGGCATCCACCACTCTGTCTCTACCGCCGAACAGACTAAGCTGTATGGAAGCGGGAATAA ACTGGTTACAGTGGGAAGCAGTAACTATCAGCAGAGTTTTGTCCCATCCCCAGGCGCTAGGCCACAAGTCAACGGACAGAGCGGGAGAA TAGATTTTCATTGGCTCATGTTAAATCCTAACGACACAGTGACTTTCTCGTTTAATGGCGCTTTTATCGCCCCTGATAGGGCCTCTTTTTTA AGAGGCAAGAGCATGGGAATCCAGTCTGGAGTGCAGGTCGATGCTAATTGCGAAGGCGACTGCTACCACAGCGGAGGAACAATCATCT CTAATTTGCCCTTCCAGAATATTGATAGTCGCGCAGTGGGCAAATGCCCCCGGTACGTAAAGCAGCGCTCACTCCTGCTGGCAACAGGG ATGAAAAATGTCCCTGAGATCCCAAAGGGCAGAGGACTGTTTGGTGCCATTGCCGGATTCATCGAGAACGGATGGGAGGGGCTGATCGA CGGATGGTATGGATTCCGGCATCAGAACGCCCAGGGCGAAGGGACAGCCGCTGATTATAAGTCCACCCAATCAGCCATTGATCAGATTA CCGGCAAACTAAATCGCTTGATTGAGAAAACAAATCAGCAATTCGAATTGATTGACAATGAGTTTAATGAGGTTGAGAAACAGATCGGAAA CGTGATTAATTGGACACGGGATTCTATAACCGAGGTCTGGTCATACAATGCTGAACTTCTGGTCGCGATGGAGAACCAGCACACAATTGA CCTGGCCGATAGTGAAATGGATAAGTTATACGAACGGGTCAAACGGCAGCTGAGGGAGAACGCCGAAGAAGACGGTACCGGTTGTTTC GAAATCTTCCATAAGTGTGATGACGACTGCATGGCCTCAATTCGGAACAACACCTACGATCACAGCAAGTATAGAGAAGAGGCTATGCAA AACCGCATACAGATCGACCCAGTAAAATTAAGTAGTGGCTACAAAGATGTGATTCTTTGGTTCTCCTTTGGAGCCAGCTGCTTTATTCTGC TGGCAATAGTTATGGGCCTCGTCTTTATCTGTGTCAAGAATGGAAACATGAGGTGTACTATTTGCATA
2822	ATGAACACGCAGATCCTTGTATTTGCGCTCATAGCAATAATCCCTACGAATGCAGATAAGATTTGCCTGGGACACCATGCTGTCTCTAACG GAACCAAGGTCAATACACTGACTGAACGTGGCGTTGAGGTGGTGAATGCAACTGAGACCGTTGAGCGAACCAACATCCCCCGCATTTGC TCTAAGGGAAAGAAGACAGTCGATCTCGGACAGTGTGGCCTGCTCGGCACAATTACGGGGCCCCCTCAGTGCGATCAGTTCTTGGAATT CTCTGCCGACCTCATCATCGAGCGCCGGGAAGGCTCTGACGTTTGTTATCCTGGCAAATTCGTGAACGAGGAGGCATTGAGGCAAATCT TGAGAGAATCTGGAGGTATAGATAAGGAAGCCATGGGCTTTACGTACTCTGGGATTAGGACCAATGGTGCAACTTCTGCGTGCCGTAGG AGCGGAAGTTCTTTCTACGCTGAGATGAAATGGCTGCTGTCGAATACTGATAATGCCGCTTTTCCCCAGATGACCAAGTCTTACAAGAATA CCAGGAAGTCTCCTGCCCTGATTGTGTGGGGTATCCACCACAGCGTGTCCACAGCCGAACAAACCAAACTTTATGGCTCAGGTAATAAGC TCGTCACTGTCGGTAGCTCTAATTATCAGCAGTCTTTTGTCCCTAGCCCCGGGGCTAGGCCACAGGTCAACGGCCAGAGCGGACGGATT GACTTCCACTGGCTGATGCTGAACCCGAACGACACAGTAACGTTTAGCTTCAATGGCGCCTTTATTGCCCCGGACCGAGCTTCATTCCTG CGGGGTAAGAGTATGGGGATCCAATCAGGTGTTCAGGTGGACGCCAATTGCGAAGGCGATTGCTATCATTCCGGGGGTACCATTATCTC CAACCTCCCATTTCAAAATATTGATAGCAGGGCCGTGGGCAAATGTCCCAGGTATGTTAAGCAGCGTTCTCTGTTACTGGCCACTGGCAT GAAGAATGTCCCTGAAATCCCAAAGGGCAGAGGACTGTTCGGCGCGATCGCGGGATTTATCGAAAACGGATGGGAGGGCTTGATCGAT GGCTGGTATGGGTTTCGACATCAGAACGCCCAAGGGGAGGGAACCGCCGCTGATTATAAAAGCACCCAGAGTGCCATTGATCAGATCAC GGGCAAACTGAACCGGCTCATTGAGAAAACAAACCAGCAGTTCGAACTCATCGATAACGAGTTTAATGAGGTGGAGAAACAAATCGGCAA CGTTATTAACTGGACTAGGGATTCTATCACGGAGGTCTGGTCCTACAACGCAGAACTGCTGGTGGCAATGGAAAATCAACATACTATTGA TCTGGCCGATTCCGAAATGGACAAGCTTTACGAGAGGGTCAAGAGACAGCTGAGAGAGAACGCTGAAGAGGATGGGACCGGATGCTTC GAAATCTTTCATAAGTGTGATGATGATTGCATGGCCAGTATCCGTAATAATACATATGATCATTCTAAATACCGCGAAGAGGCGATGCAGA ATCGAATCCAAATTGACCCAGTTAAACTGTCTAGTGGCTACAAAGACGTCATCCTGTGGTTTTCGTTCGGAGCTTCTTGCTTCATTCTGTT GGCAATAGTGATGGGCCTCGTATTTATATGTGTGAAAAACGGGAATATGAGATGCACAATTTGTATT
2823	ATGAATACCCAGATCCTGGTGTTCGCCCTTATTGCCATCATTCCGACCAACGCTGACAAAATCTGTCTCGGACACCACGCTGTCTCCAAT GGGACAAAGGTGAATACCTTGACCGAGCGAGGGGTCGAGGTCGTCAACGCGACAGAAACAGTGGAGAGGACCAACATTCCCCGGATAT GCTCAAAGGGCAAAAGAACGGTGGATCTGGGGCAATGCGGGCTGCTGGGCACTATAACCGGACCCCCCCAATGTGACCAGTTTCTTGA ATTTTCTGCCGATCTGATAATCGAACGCCGGGAAGGATCCGACGTGTGCTATCCTGGAAAGTTTGTGAATGAGGAAGCCCTCCGGCAGA TTCTGAGGGAATCCGGAGGGATAGATAAGGAAGCAATGGGGTTCACTTATAGCGGTATCCGCACAAATGGGGCGACTAGTGCATGTCGT CGAAGTGGAAGCAGTTTCTATGCCGAGATGAAATGGCTGCTAAGCAATACAGACAACGCCGCATTCCCACAGATGACCAAGTCCTATAAA AATACAAGAAAGAGTCCCGCGCTTATCGTGTGGGGCATACATCACTCCGTCAGCACCGCCGAACAGACCAAGCTCTATGGCTCAGGCAA CAAGTTGGTGACAGTCGGCAGTAGCAATTACCAGCAGTCCTTCGTGCCTAGCCCAGGCGAAAGGCCTCAGGTGAATGGTTTGAGCGGC CGAATAGACTTCCATTGGTTAATGCTCAATCCTAACGATACAGTCACGTTCAGTTTCAATGGTGCCTTTATAGCGCCTGATCGCGCATCAT TCCTGCGTGGAAAATCTATGGGCATTCAGAGCGGAGTCCAGGTCGACGCCAACTGCGAAGGCGACTGTTATCACTCCGGGGGCACCATT ATTTCCAATTTGCCCTTCCAGAACATTGATAGCAGAGCAGTTGGCAAGTGCCCGCGCTACGTAAAACAAAGGTCTCTGCTTTTGGCTACA GGTATGAAAAATGTCCCAGAAATTCCCAAGGGCCGCGGATTGTTCGGGGCGATCGCTGGTTTTATTGAGAACGGTTGGGAAGGCCTCAT CGATGGGTGGTATGGTTTTCGGCATCAGAACGCCCAGGGCGAAGGCACTGCCGCGGATTATAAGTCCACCCAGTCAGCGATCGATCAG ATCACAGGAAAGCTGAACCGACTTATCGAAAAAACAAACCAGCAGTTTGAACTAATTGACAATGAATTCAACGAGGTGGAGAAACAGATT GGCAATGTGATTAACTGGACCCGTGATAGCATCACAGAGGTCTGGAGCTATAATGCAGAACTGTTGGTTGCTATGGAGAATCAGCACACC ATCGACCTGGCCGACTCTGAAATGGACAAACTGTACGAGCGAGTGAAACGGCAGCTGCGTGACAACGCTGAGGAAGACGGCACCGGTT GTTTCGAAATCTTCCATAAATGCGACGACGACTGTATGGCTTCTATTAGAAACAATACCTACGATCACTCCAAGTACCGGGAGGAAGCAAT GCAGAATCGAATCCAGATCGACCCAGTTAAGTTGTCCAGCGGATACAAGGACGTGATTCTATGGTTTAGTTTTGGAGCCTCCTGCTTCAT TCTTCTGGCCATTGTGATGGGTTTGGTTTTTATCTGTGTAAAAAATGGAAATATGCGCTGTACTATTTGTATT

2824	ATGAACACTCAGATTCTTGTGTTCGCTCTGATCGCAATTATACCCACAAACGCGGATAAAATCTGCCTTGGACATCACGCAGTGTCGAAC GGAACTAAGGTGAACACTCTCACCGAGCGGGGCGTCGAAGTAGTCAATGCTACAGAGACCGTCGAACGCACTAATATTCCACGTATCTG CTCCAAGGGGAAAAGGACCGTCGACCTGGGACAATGTGGCTTGCTAGGGACTATTACCGGCCCACCGCAGTGTGATCAATTCCTCGAGT TCTCAGCCGACCTGATCATCGAGCGACGCGAGGGCAGCGATGTCTGTTACCCCGGGAAATTTGTTAATGAGGAAGCATTAAGGCAGATC CTGCGCGACTCTGGCGGGATCGACAAGGAGGCCATGGGCTTTACATATTCCGGGATCCGGACAAACGGCGCCACAAGTGCCTGTCGAC GGTCAGGATCATCATTCTATGCCGAGATGAAATGGCTTCTCAGCAATACCGATAACGCTGCATTTCCACAGATGACAAAGTCGTATAAAAA CACAAGAAAGTCACCCGCTCTGATTGTTTGGGGAATTCACCACAGTGTCTCAACCGCAGAACAGACCAAGTTGTACGGGAGTGGAAACA AGTTGGTAACCGTGGGATCTTCAAACTATCAGCAGAGCTTCGTGCCAAGCCCCGGGGAACGCCCCCAAGTGAATGGACTGTCTGGACG GATTGACTTCCACTGGCTGATGCTGAACCCCAACGACACAGTGACATTCAGTTTTAATGGAGCATTTATCGCCCCGGACAGAGCAAGCTT TCTCAGGGGTAAGAGTATGGGAATTCAGAGTGGTGTACAGGTTGACGCTAACTGCGAAGGTGACTGTTATCATTCTGGTGGGACTATCAT CTCCAACCTGCCTTTCCAGAACATTGACTCCAGAGCAGTCGGCAAGTGCCCCAGGTACGTGAAGCAAAGGTCTTTGCTGCTCGCAACCG GCATGAAGAACGTGCCTGAAATACCTAAGGGCCGTGGCTTGTTTGGCGCAATCGCGGGATTTATCGAAAATGGGTGGGAGGGGTTGATT GACGGTTGGTACGGCTTCCGTCATCAGAATGCCCAAGGGGAGGGCACCGCCGCAGACTATAAAAGCACCCAAAGTGCAATCGATCAGAT CACCGGCAAATTGAACCGGCTGATTGAGAAAACGAATCAGCAGTTCGAACTGATCGACAACGAATTTAACGAGGTCGAAAAGCAGATTGG AAATGTGATCAACTGGACCCGGGATTCCATCACCGAAGTCTGGTCATATAACGCGGAACTGCTTGTAGCAATGGAGAATCAACACACTAT CGATCTGGCGGACAGTGAGATGGATAAGCTCTACGAGAGGGTTAAGAGACAACTGCGCGAGAATGCTGAGGAAGATGGTACCGGGTGT TTCGAGATTTTTCACAAATGTGATGATGACTGCATGGCTAGTATTCGCAACAACACATATGATCACTCCAAATACCGCGAAGAGGCTATGC AGAACCGCATCCAAATAGATCCCGTAAAGCTTAGCTCGGGTTACAAAGATGTAATCTTATGGTTTAGTTTCGGAGCGAGCTGTTTTATTCT ACTCGCCATTGTGATGGGGCTTGTCTTCATATGTGTGAAGAATGGAAATATGCGCTGCACAATCTGTATA
2825	ATGAATACCCAGATTCTTGTCTTCGCACTCATAGCTATTATTCCTACGAACGCGGACAAGATTTGTCTGGGGCATCACGCCGTCTCCAATG GGACAAAAGTGAACACCCTGACTGAACGAGGGGTTGAAGTGGTCAATGCAACTGAAACAGTGGAACGGACAAACATTCCAAGAATCTGT TCCAAAGGAAAGCGAACAGTGGACCTGGGACAGTGCGGCCTGCTGGGTACCATTACTGGCCCCCCACAGTGCGACCAGTTCCTGGAAT TTAGCGCAGATCTTATCATCGAGCGGCGGGAAGCAAGTGACGTGTGTTATCCCGGGAAGTTTGTAAACGAGGAGGCCCTCCGGCAGATC CTTCGTGAGAGCGGCGGGATTGATAAAGAGGCTATGGGCTTTACCTATTCCGGAATAAGAACCAACGGGGCCACTTCTGCGTGTCGTAG GAGCGGCAGCAGCTTCTACGCCGAAATGAAGTGGCTGTTGTCTAACACCGATAATGCCGCCTTCCCTCAGATGACCAAGTCCTATAAGAA CACCCGGAAATCCCCTGCCCTGATTGTTTGGGGGATTCACCATTCCGTATCTACCGCCGAACAGACCAAGTTGTACGGTTCCGGAAACAA GCTCGTCACAGTTGGCTCCTCCAATTACCAGCAGAGTTTCGTACCTTCTCCCGGCGCAAGACCACAGGTCAACGGACTAAGCGGGCGGA TTGACTTTCACTGGCTCATGTTGAATCCAAATGACACTGTAACCTTTAGTTTCAACGGTGCCTTTATAGCACCAGATCGGGCCAGCTTTCT GCGTGGAAAGTCTATGGGCATTCAAAGTGGTGTGCAAGTGGATGCTAACTGTGAGGGAGATTGCTACCATTCAGGTGGGACAATCATTA GCAATCTTCCGTTTCAGAATATTGATTCACGGGCTGTGGGCAAGTGCCCCCGTTACGTGAAGCAGAGAAGCCTGCTTCTTGCCACGGGA ATGAAGAATGTTCCCGAAATACCAAAGGGTAGGGGTCTCTTCGGAGCAATAGCCGGATTTATCGAGAATGGCTGGGAGGGGCTTATCGA TGGTTGGTACGGCTTTCGCCATCAGAACGCTCAGGGAGAGGGGACCGCAGCAGACTACAAATCCACCCAGTCTGCTATCGACCAGATCA CTGGCAAGCTGAATCGGATAATCGAGAAGACTAACCAGCAGTTTGAGCTGATCGACAATGAGTTTAACGAAGTGGAGAAACAGATTGGCA ATGTGATCAATTGGACAAGAGATTCCATTACGGAAGTTTGGTCATATAATGCCGAGCTCCTGGTTGCAATGGAGAACCAACATACAATCGA CCTGGCGGACAGTGAGATGGACAAGCTGTACGAGCGTGTGAAGCGCCAACTCCGGGAAAATGCTGAGGAAGACGGCACGGGCTGCTTT GAAATCTTCCACAAGTGTGACGACGACTGCATGGCAAGCATAAGGAACAACACCTATGATCATTCTAAGTACAGAGAGGAGGCGATGCA GAATCGCATCCAGATTGACCCAGTTAAGCTTTCCAGCGGGTACAAAGACGTCATCTTGTGGTTTTCCTTTGGTGCCTCTTGTTTCATTTTG CTCGCTATCGTAATGGGACTGGTATTCATCTGCGTGAAAAATGGAAATATGAGATGCACCATCTGCATT
2826	ATGAACACACAAATTCTGGTGTTCGCTCTCATCGCAATAATTCCTACCAACGCCGATAAGATTTGCCTGGGGCACCATGCGGTCAGTAAC GGAACCAAAGTCAATACACTCACAGAAAGAGGCGTTGAGGTGGTGAATGCCACAGAAACTGTTGAACGTACAAATATTCCGCGAATCTGT TCTAAAGGCAAGAAAACCGTTGATCTGGGTCAATGCGGCCTTCTGGGAACCATTACAGGGCCACCTCAGTGCGACCAGTTTCTCGAATTT TCTGCTGATCTCATTATCGAGAGGCGGGAAGGCAGTGACGTTTGCTACCCTGGTAAGTTCGTAAATGAGGAGGCCCTCAGGCAGATCCT GAGGAAGTCCGGCGGTATTGATAAGGAAGCCATGGGCTTCACTTATTCTGGGATCAGGACCAACGGGGCCACTTCAACATGTAGAAGGT CCGGAAGCTCTTTCTATGCAGAGATGAAATGGCTCCTGAGCAACACAGACAACGCCGCCTTTCCACAGATGACCAAGAGCTATAAGAATA CTCGCAAATCTCCAGCCATCATAGTGTGGGGTATCCACCACAGCGTGTCCACGGCCGAGCAGACAAAGCTGTATGGGAGCGGCAACAA GCTGGTGACTGTCGGTAGCTCGAATTACCAGCAGTCTTTCGTGCCTAGCCCCGGGGCCCGGCCGCAGGTCAATGGGCTTTCCGGGCGA ATCGACTTTCACTGGTTGATGTTAAACCCAAACGACACGGTGACATTCAGCTTTAACGGCGCCTTTATTGCACCTGACCGCGCATGTTTCC TGAGGGGCAAGTCTATGGGTATCCAGTCTGGTGTACAAGTCGACGCCGACTGTGAAGGAGACTGCTACCATTCCGGCGGGACAATAATC TCAAACCTACCCTTCCAGAACATCGACAGTAGGGCGGTGGGCAAGTGTCCTAGGTACGTGAAACAGCGGAGCCTGCTGTTGGCCACAG GCATGAAGAATGTCCCCGAAATTCCAAAGGGGCGGGGTCTCTTTGGAGCAATTGCAGGATTCATTGAGAATGGTTGGGAAGGTCTCATC GATGGATGGTACGGGTTCCGCCATCAGAACGCCCAGGGTGAGGGGACCGCAGCCGACTATAAGAGTACCCAGTCAGCGATCGACCAAA TCACCGGCAAGCTAAATCGGCTGATTGAGAAGACAAACCAACAGTTTGAATTAATTGACAACGAGTTTAATGAGGTGGAAAGACAGATCG GCAACGTGATCAACTGGACACGCGATTCCATCACAGAAGTCTGGTCTTATAATGCAGAGCTGCTAGTGGCCATGGAGAATCAGCACACC ATTGATCTAGCCGACTCTGAAATCGCACAAGTTGTATGAACGGGTAAAACGTCAGTTACGACAAAATGCCGAGGAAGCATGGCACAGGGTG TTTCGAGATTTTTCACAAATGTGACGACGATTGTATGGCAAGTATTCGTAACAACACATACGACCACAGCAAATACAGAGAGGAGGCCAT GCAGAACCGCATCCAGATAGACCCAGTTAAACTGAGCTCCGGGTACAAGGATGTGATCCTCTGGTTTAGTTTCGGCGCAAGTTGTTTCAT CCTACTCGCGATAGTTATGGGCCTGGTATTCATTTGCGTCAAGAACGGGAACATGAGGTGTACCATCTGCATT

2827	ATGAATACCCAGATTCTCGTGTTTGCTTTGATTGCTATCATCCCCACGAATGCTGATAAAATTTGCCTAGGCCATCATGCTGTGTCTAACG GGACCAAGGTGAACACCCTCACTGAGAGAGGCGTGGAGGTCGTGAACGCGACCGAGACTGTTGAGCGCACAAACATTCCCAGGATTTG TAGCAAAGGAAAAAGAACCGTTGATCTCGGCCAATGCGGCCTCCTGGGTACCATCACGGGGCCTCCACAGTGTGACCAGTTCTTGGAAT TCAGTGCAGACCTTATCATCGAGCGCAGGGAGGGCTCTGATGTATGCTACCCTGGAAAATTCGTGAACGAGGAGGCTTTGAGACAAATC CTGAGGGACTCTGGAGGCATCGATAAGGAGGCCATGGGCTTTACATACAGCGGCATTCGCACTAACGGAGCCACTTCCGCATGCAGGC GGTCCGGCAGCTCATTCTACGCTGAAATGAAGTGGCTCCTGAGTAATACGGATAACGCCGCCTTTCCGCAGATGACCAAAAGTTATAAGA ATACAAGGAAATCACCTGCTCTCATTGTCTGGGGTATCCACCACTCGGTGAGCACCGCTGAGCAGACCAAACTGTACGGGTCAGGCAAT AAACTTGTGACCGTGGGCTCAAGCAATTACCAGCAGTCCTTTGTGCCTTCACCCGGCGAAAGACCTCAAGTCAACGGACTGTCTGGTCG GATCGACTTTCACTGGCTGATGTTAAACCCAAATGATACAGTGACATTCTCCTTCAACGGAGCATTCATAGCCCCCGATAGGGCCAGCTT CCTCCGGGGGAAATCCATGGGAATTCAGTCAGGCGTCCAGGTTGACGCTAACTGCGAGGGTGACTGTTATCATTCCGGAGGTACTATAA TAAGCAATCTTCCATTCCAGAATATAGATTCACGAGCGGTCGGCAAGTGTCCGCGGTACGTTAAACAGCGTAGCTTACTGCTCGCAACCG GGATGAAAAACGTTCCAGAAATTCCTAAGGGCAGAGGGCTGTTTGGGGCCATCGCCGGCTTCATCGAGAACGGTTGGGAAGGGCTCAT CGATGGCTGGTACGGTTTTAGACACCAGAATGCCCAGGGCGAGGGGACCGCAGCAGATTACAAGAGCACACAGTCAGCCATAGATCAA ATTACTGGGAAGCTCAATCGGTTGATAGAAAAGACTAACCAGCAGTTCGAATTGATCGATAATGAATTTAACGAGGTCGAGAAACAAATCG GCAATGTTATTAATTGGACTAGAGACTCCATCACAGAAGTGTGGTCCTACAATGCCGAACTGCTGGTGGCTATGGAGAACCAGCACACAA TCGACCTTGCAGACAGTGAGATGGATAAACTGTATGAGAGAGTGAAAAGGCAGCTCCGCGAGAACGCTGAGGAAGATGGAACCGGCTG TTTTGAAATTTTCCATAAGTGCGACGATGATTGCATGGCTTCTATCCGGAATAATACTTACGACCACAGTAAGTACCGAGAAGAAGCCATG CAGAATCGAATTCACATAGACCCCGTCAAGCTGAGCTCCGGCTATAAAGATCTGATTCTGTCGTTCAGCTTTGGCGCCTCTTGTTTCATT CTTTTGGCCATCGTGATGGGCCTTGTGTTCATTTGTGTAAAAAATGGCAATATGAGGTGCACAATATGTATT
2828	ATGAATACCCAGATACTGGTGTTCGCCCTGATTGCAATCATCCCAACTAACGCTGATAAGATTTGTCTCGGTCACCACGCTGTGAGTAAC GGAACTAAAGTTAACACACTGACAGAGAGGGGAGTGGAGGTTGTTAACGCGACAGAGACAGTGGAGCGCACTAATATACCACGGATATG TAGTAAGGGAAAAAGAACTGTGGACTTGGGTCAATGCGGACTGTTGGGGACGATAACTGGCCCCCCTCAGTGTGATCAGTTTCTGGAGT TCTCAGCCGATCTGATCATCGAGAGGCGAGAGGGGAGTGATGTGTGTTATCCTGGTAAGTTTGTGAATGAGGAGGCTCTCAGGCAAATC CTCAGGGAATCTGGAGGGATCGATAAGGAGGCTATGGGCTTCACTTACTCTGGAATAAGAACTAACGGCGCCACTAGCGCCTGTAGGCG ATCCGGAAGTTCGTTCTACGCTGAAATGAAGTGGTTGCTCAGCAACACTGACAACGCAGCATTTCCCCAAATGACCAAAAGCTATAAGAA TACTCGCAAGTCTCCCGCCCTGATAGTGTGGGGGATTCACCATAGTGTCTCTACAGCCGAGCAGACTAAATTATACGGCAGTGGCAACAA GCTGGTGACGGTGGGGAGTTCAAATTACCAGCAGAGCTTCGTGCCCAGTCCGGGAGCTCGCCCTCAAGTTAATGGTCTTTCCGGCAGAA TTGATTTCACTGGCTGATGCTTAACCCTAACGACACCGTGACCTTTTCCTTCAACGGCGCTTTCATTGCCCCGGACCGAGCTAGTTTCCT TAGGGGGAAGTCTATGGGCATACAGTCTGGTGTGCAAGTTGATGCGAATTGCGAGGGGGATTGTTACCACTCTGGAGGTACAATCATCA GTAATTTACCCTTCCAGAACATTGATAGTAGAGCCGTTGGGAAGTGTCCACGGTATGTTAAACAGCGCAGCCTGCTTCTGGCCACAGGGA TGAAGAATGTCCCCGAAATCCCCAAAGGCAGAGGATTGTTCGGCGCAATCGCGGGGTTTATTGAAAACGGTTGGGAAGGCCTCATTGAC GGCTGGTACGGGTTCCGGCACCAGAATGCTCAAGGAGAGGGCACCGCCGCCGATTATAAGTCTACCCAGAGCGCCATCGACCAGATCA CAGGCAAATTGAACCGAATTATCGAGAAGACCAACCAACAGTTCGAGCTTATCGACAACGAATTTAATGAAGTAGAAAAGCAGATCGGAA ACGTTATTAACTGGACTCGGGACAGTATTACTGAAGTGTGGTCCTATAATGCTGAGCTGTTGGTGGCAATGGAGAACCAGCACACCATTG ACCTGGCCGACAGCGAGATGGACAAGCTTTACGAAAGAGTGAAGCGACAGCTTCGGGAAAACGCCGAGGAAGATGGAACCGGCTGTTT TGAAATCTTCCATAAGTGCGATGATGACTGTATGCCCAGTATACGCAACAACACTTATGACCATTCAAAATACCCCCAAGAAGCCATGCA GAATCGGATCCAGATCGACCCTGTGAAGCTGTCGAGCGGGTATAAGGACGTGATACTGTGGTTCAGCTTCGGAGCCAGCTGCTTCATTC TGCTGGCTATTGTCATGGGCCTGGTGTTCATTTGCGTCAAAAATGGTAATATGAGGTGTACGATCTGCATC
2829	ATGAACACTCAGATCCTCGTGTTTGCCTTGATCGCAATAATCCCCACCAATGCCGATAAGATATGTCTCGGACACCACGCTGTGTCCAAC GGGACCAAGGTTAATACTCTGACAGAGCGGGGCGTTGAAGTGGTGAATGCCACCGAAACCGTGGAACGGACAAATATTCCACGAATTTG CAGCAAGGCAAAGAAGACCGTGGACCTTGGTCAGTGTGGGCTGCTGGGGACTATCACCGGGCCTCCCCAATGCGACCAGTTCCTCGAG TTCTCTGCAGATCTGATTATTGAGCGGCGCGAGGGTAGTGATGTGTGCTACCCTGGCAAATTCGTGAATGAGGAGGCACTGAGACAGAT CCTGAGAGAGAGTGGCGGAATCGATAAAGAAGCAATGGGGTTTACCTATTCCGGGATACGGACTAACGGGGCAACATCGGCTTGCAGG CGAAGTGGCTCATCTTTCTACGCCGAAATGAAGTGGCTCCTGAGCAACACCGATAATGCCGCTTTCCCACAGATGACTAAGTCCTATAAG AACACCCGGAAGAGCCCGGCCTTGATTGTCTGGGGGATCCACCATTCTGTCAGTACAGCTGAACAAACTAAGTTGTACGGTTCCGGAAA CAAGCTGGTGACCGTGGGATCTAGCAATTATCAGCAGTCATTCGTGCCTTCACCTGGCGCACGACCACAGGTCAATGGACAGAGTGGTA GAATCGACTTTCACTGGCTGATGCTTAATCCGAATGACACCGTGACCTTCTCCTTCAACGGAGCCTTTATTGCCCCTGACCGAGCTAGCT TTCTCAGGGGAAAGTCTATGGGCATACAGAGCGGCGTGCAGGTCGACGCAAATTGTGAAGGGGACTGTTACCACTCAGGCGGGACCAT TATAAGTAATCTGCCTTTTCAGAACATTGACAGCAGAGCGGTGGGCAAATGTCCGCGTTACGTGAAGCAGAGATCACTGCTCTTGGCCAC CGGCATGAAGAATGTCCCTGAGATCCCGAAAGGCAGGGGTCTGTTTGGGGCTATTGCGGGCTTCATTGAAAACGGCTGGGAAGGTCTCA TCGATGGTTGGTACGGATTCCGGCACCAGAATGCTCAGGGGGAGGGCACCGCCGCCGATTATAAGAGTACCCAGTCTGCCATCGACCA AATCACCGGCAAGCTTAACCGTCTTATCGAGAAAACAAATCAGCAGTTCGAGCTGATCGATAACGAATTTAATGAGGTAGAGAAACAGATT GGTAATGTGATTAACTGGACACGCGACTCCATTACCGAGGTGTGGTCGTATAATGCAGAATTACTGGTGGCCATGGAGAATCAACACACC ATTGATCTGGCCGATAGCGAGATGGACAAGCTGTATGAGAGAGTGAAGAGACAACTGAGGCAAAACGCGGAGGAGGATGGAACAGGAT GTTTCGAGATCTTCCACAAGTGTGACGACGATTGCATGGCGAGTATCCGGAACAACACCTACGATCATTCAAAATATCGAGAGGAAGCTA TGCAAAACCGAATTCAGATCGATCCAGTAAAATTATCTTCAGGGTATAAGGATGTCATACTATGGTTCAGCTTCGGGGCGTCGTGCTTCAT CTTGCTGGCTATCGTCATGGGGCTCGTCTTTATTTGTGTGAAAAATGGAAATATGCGGTGCACGATCTGCATT

2830	ATGAACACGCAAATCCTGGTATTTGCGCTGATCGCTATCATTCCCACTAATGCTGACAAAATCTGTCTGGGCCATCACGCTGTGTCAAACG GAACCAAGGTGAATACGCTGACCGAACGCGGCGTCGAGGTGGTCAATGCCACGGAAACAGTCGAAAGAACAAACATTCCCAGAATCTGC TCAAAGGGTAAAAAGACGGTAGATCTCGGGCAGTGTGGGCTTCTTGGAACAATCACTGGCCCACCACAATGCGATCAATTTCTCGAATTC AGCGCCGACCTGATCATCGAGAGGAGGGAAGGTAGTGATGTGTGCTACCCCGGGAAGTTTGTGAATGAGGAGGCATTGCGCCAAATAC TGAGACAGAGCGGAGGGATTGACAAGGAGGCCATGGGATTCACCTACAGCGGGATCAGAACTAATGCAGCCACGAGCGCTTGCACGAG GAGTGGAAGCAGCTTTTATGCTGAAATGAAGTGGCTACTGTCCAATACGGATAACGCTGCGTTCCCACAAATGACAAAGTCGTACAAGAA TACGCGAAAGAGCCCAGCCCTCATTGTCTGGGGTATCCATCACAGCGTGAGTACCGCAGAACAAACCAAACTTTACGGGTCAGGTAACA AACTGGTAACCGTTGGAAGCTCCAATTATCAGCAATCGTTCGTTCCTTCACCTGGGGCTCGCCCACAAGTCAATGGACAGAGCGGCAGA ATTGACTTCCATTGGCTCATGCTCAATCCAAACGATACAGTCACCTTTTCATTTAACGGAGCTTTCATCGCTCCTGATCGGGCTAGCTTTCT AAGGGGCAAGTCTATGGGTATTCAGTCCGGCGTACAAGTCGACGCCAACTGTGAGGGGGACTGCTACCATAGCGGAGGCACTATCATAT CAAATCTCCCCTTCCAGAATATCGACTCCAGGGCTGTTGGTAAGTGCCCAAGATACGTGAAGCAGAGATCCCTCCTTCTGGCTACAGGCA TGAAAAATGTGCCAGAAATTCCAAAGGGACGGGGCCTATTTGGAGCTATTGCAGGCTTCATCGAGAACGGATGGGAAGGACTGATCGAT GGGTGGTACGGATTTAGGCACCAGAACGCCCAGGGAGAGGGAACAGCTGCTGACTACAAATCTACCCAGTCTGCAATTGATCAGATCAC CGGTAAACTGAATAGACTGATTGAGAAAACTAACCAGCAGTTCGAGCTAATTGATAATGAGTTCAATGAAGTGGAGAAGCAGATCGGCAA CGTGATTAACTGGACCAGAGATTCAATCACGGAGGTCTGGTCTTACAACGCCGAGTTGCTGGTGGCTATGGAAAACCAGCACACCATAG ACCTGGCCGACTCTGAAATGGACAAATTGTACGAGAGGGTAAAAAGGCAGCTCCGAGAAAACGCCGAGGAGGACGGTACTGGCTGCTTT GAGATATTTCACAAGTGCGACGATGATTGCATGGCCTCCATTCGAAACAACACGTACGACCACAGTAAATACAGGGAGGAAGCCATGCA GAACCGCATCCAGATTGACCCTGTCAAGCTGAGTAGTGGCTACAAGGACGTTATCCTCTGGTTTAGTTTTGGAGCATCATGTTTTATACTG TTGGCAATAGTCATGGGGCTTGTCTTCATCTGCGTGAAAAATGGCAACATGCGTTGTACAATCTGCATT
2831	ATGAATACCCAAATTCTGGTGTTCGCTCTAATTGCCATCATACCAACAAACGCAGACAAGATTTGTCTGGGCCACCACGCCGTGTCAAAC GGAACGAAAGTGAACACTCTGACTGAGCGTGGCGTGGAGGTAGTGAACGCTACCGAAACTGTGGAGCGGACTAATATCCCAAGGATCTG TAGCAAGGGAAAAAAGACAGTGGACCTGGGTCAGTGCGGATTGCTAGGAACCATAACCGGTCCACCTCAGTGCGATCAGTTTTTGGAGT TTAGCGCCGATCTCATCATCGAAAGAAGGGAGGGTTCTGATGTTTGTTACCCAGGAAAATTTGTTAACGAAGAAGCCCTTCGCCAGATCC TTAGAGAATCAGGCGGCATCGATAAAGAGGCAATGGGGTTCACCTATTCAGGAATACGCACAAATGGCGCCACGAGCGCGTGCAGGCG CAGCGGGAGTTCATTTTATGCTGAAATGAAATGGCTCCTCTCAAACACTGATAACGCTGCATTTCCACAAATGACGAAAAGTTATAAGAAT ACGAGAAAGTCTCCAGCCCTCATTGTCTGGGGCATTCATCACTCAGTGTCGACTGCCGAGCAAACCAAGCTGTACGGCAGCGGTAACAA ATTGGTAACTGTCGGCAGTAGTAACTACCAGCAAAGTTTTGTGCCTAGCCCCGGCGCTCGGCCACAAGTGAACGGACAGAGCGGGCGC ATCGATTTTCACTGGCTGATGCTAAATCCAAATGACACAGTCACCTTTTCGTTTAACGGGGCCTTCATCGCCCCTGACCGGGCTTCTTTCC TGAGAGGAAAAAGCATGGGAATCCAGTCCGGAGTCCAGGTTGACGCCAACTGCGAAGGAGACTGTTATCATTCCGGGGGCACTATTATA AGCAATCTGCCTTTTCAGAACATAGATAGTCGCGCCGTAGGCAAGTGTCCCCGCTACGTCAAACAGAGAAGCCTTCTGCTGGCTACCGG TATGAAGAACGTCCCCGAAATCCCTAAAGGCCGAGGACTATTCGGCGCTATCGCCGGCTTCATTGAAAATGGCTGGGAGGGCTTGATAG ATGGGTGGTATGGCTTTCGCCATCAGAATGCACAGGGTGAAGGGACTGCAGCCGATTATAAATCTACTCAGTCCGCCATCGATCAGATTA CCGGGAAGCTCAATAGACTCATTGAGAAGACAAATCAACAGTTCGAACTGATCGATAACGAGTTCAACGAGGTAGAGAAACAGATCGGGA ATGTCATTAACTGGACCCGAGATTCTATCACAGAAGTTTGGTCTTACAACGCGGAGCTGCTGGTCGCAATGGAAAATCAGCATACCATCG ATCTGGCCGATTCAGAAATGGACAAGCTTTATGAGAGAGTGAAGCGCCAGTTGCGCGAGAATGCGGAAGAGGACGGCACCGGATGCTT CGAGATTTTTCACAAATGTGATGATGATTGTATGGCTTCGATTCGGAATAACACCTACGATCACTCCAAATATCGCGAAGAGGCTATGCAG AACAGAATACAAATCGATCCCGTCAAACTTAGCAGCGGCTACAAAGACGTCATTCTGTGGTTTTCCTTCGGGGCATCCTGTTTTATACTGC TGGCCATAGTGATGGGTCTGGTGTTTATATGTGTAAAGAACGGCAACATGCGCTGTACCATTTGCATC
2832	ATGAACACCCAGATCCTCGTGTTCGCCTTAATCGCTATCATTCCAACCAATGCAGACAAGATATGTCTGGGTCATCATGCCGTTAGCAATG GCACCAAGGTCAACACCCTGACCGAGCGCGGAGTCGAGGTTGTTAATGCAACCGAAACTGTGGAGCGGACAAATATTCCCCGGATCTGC AGCAAGGGCGAAGAGGACAGTCGATCTGGGCCAATGCGGACTGCTGGGCACAATCACCGGACCCCCTCAGTGCGATCAATTTCTGGAGT TCAGCGCCGACCTGATCATCGAGAGAAGGGAGGGGAGTGACGTATGTTACCCAGGGAAGTTTGTGAACGAGGAGGCCCTGAGACAGAT TCTGAGAGAAAGCGGCGGCATCGATAAAGAGGCCATGGGATTTACCTATAGCGGCATTCGCACCAATGGAGCCACTTCAGCTTGTAGGA GATCCGGGAGTAGCTTTTACGCTGAGATGAAATGGCTGCTTTCCAATACTGATAATGCCGCTTTCCCACAGATGACTAAGTCTTATAAGAA TACTCGCAAGAGTCCTGCCCTGATAGTCTGGGGCATCCATCATTCCGTCTCAACCGCTGAACAGACCAAACTCTACGGGTCTGGGAACAA GTTGGTGACCGTGGGCTCCTCGAATTATCAACAATCATTCGTGCCGTCCCCCGGCGCTCGGCCGCAGGTTAACGGTCTTTCAGGCCGCA TCGACTTCCACTGGCTAATGTTGAACCCAAACGATACTGTTACTTTTAGCTTTAACGGCGCATTTATTGCCCCCGACCGGGCCTCCTTCCT GCGAGGCAAGTCCATGGGTATCCAGTCTGGAGTCCAAGTGGATGCTAACTGTGAAGGCGACTGTTACCATAGTGGAGGGACAATTATTA GTAACCTGCCCTTCCAGAACATTGACAGTAGGGCCGTGGGCAAATGCCCACGGTACGTGAAGCAGCGGTCCCTGCTGCTGGCAACTGG AATGAAGAACGTGCCAGAGATCCCTAAAGGGCGGGGCCTCTTCGGCGCTATCGCCGGGTTTATAGAGAATGGTTGGGAAGGGCTGATC GATGGATGGTATGGGTTCCGCCACCAGAATGCTCAGGGTGAGGGGACAGCTGCTGACTACAAGTCTACACAGAGTGCTATTGACCAGAT CACCGGAAAACTTAACAGGATCATCGAGAAAACGAATCAGCAATTTGAGCTCATTGATAACGAGTTCAATGAAGTCGAGAAGCAAATCGG GAACGTGATAAATTGGACACGCGACTCCATCACTGAGGTCTGGAGCTATAACGCCGAGCTTCTGGTGGCAATGGAGAATCAGCATACTAT CGATCTCGCGGATTCCGAAATGGATAAGCTCTACGAGAGAGTTAAGAGGCAGCTCCGCCACAATGCCGAGGAAGATGGCACCGGATGC TTCGAGATATTCCATAAGTGCGACGACGACTGCATGGCGTCAATAAGGAATAATACTTATGATCATTCTAAGTACAGGGAGGAGGCCATG CAGAATCGCATTCAGATTGATCCTGTTAAACTAAGCTCAGGCTATAAGGACGTAATATTATGGTTTTCTTTTGGCGCTTCCTGTTTTATCCT CCTGGCTATAGTTATGGGGCTGGTGTTTATTTGTGTTAAAAATGGAAATATGAGATGCACAATTTGCATC

2833	ATGAATACCCAAATTCTGGTCTTCGCTCTGATAGCTATAATCCCAACGAATGCCGATAAGATATGTCTGGGTCATCACGCTGTCTCCAATG GGACCAAAGTGAACACATTGACTGAACGCGGCGTGGAGGTGGTTAACGCCACTGAGACCGTAGAGCGGACTAATATACCCCGGATCTG CAGTAAGGGAAAAAAGACAGTCGACTTGGGTCAGTGCGGCCTGCTAGGGACCATCACAGGGCCTCCTCAATGTGACCAGTTCCTGGAGT TCAGCGCAGACCTTATAATCGAAAGAAGAGAGGGGTCGGACGTCTGTTACCCCGGCAAGTTTGTGAACGAGGAAGCACTGAGGCAGATC CTGACAGAATCCGGCGGAATTGATAAGGAAGCCATGGGTTTCACCTACAGTGGCATACGCACCAATGGAGCAACATCCGCTTGTAGACG GAGCGGTTCTTCCTTTTACGCCGAGATGAAGTGGCTGCTGTCCAACACGGATAATGCAGCCTTCCCTCAAATGACTAAGAGTTATAAGAA TACGAGAAAGTCCCCCGCCTTGATTGTGTGGGGGATCCACCATAGCGTTAGCACAGCCGAACAGACTAAGCTTTACGGTTCAGGCAACA AGCTCGTAACCGTGGGATCCTCCAATTATCAGCAATCCTTCGTGCCTTCTCCTGGCGCCCGCCCTCAGGTTAACGGTCAAAGCGGTCGC ATAGATTTTCATTGGCTCATGCTGAACCCGAATGATACAGTCACGTTTTCTTTCAATGGGGCATTTATCGCACCAGACCGGGCCTCCTTTC TTCGCGGGAAGTCTATGGGAATACAATCCGGCGTACAGGTTGATGCTAACTGCGAGGGCGACTGTTATCATTCAGGGGGTACCATAATC TCCAACTTGCCTTTCCAAAATATTGATTCCCGGGCTGTCGGAAAGTGCCCGAGATATGTGAAGCAACGCAGTTTACTCCTGGCCACGGGA ATGAAAAACGTTCCCGAAATCCCTAAGGGAAGGGGACTATTTGGTGCTATTGCAGGGTTTATTGAGAATGGCTGGGAGGGCTTAATCGAC GGCTGGTATGGCTTCCGCCACCAAAACGCACAGGGGGAGGGGACAGCAGCAGACTATAAAAGCACGCAGAGCGCAATCGACCAAATTA CCGGCAAGTTGAATAGGCTCATTGAAAAAACAAACCAGCAATTTGAGCTCATCGACAACGAGTTCAATGAAGTTGAGAAACAGATAGGAA ATGTGATCAATTGGACTCGCGACAGCATTACCGAGGTATGGTCATACAACGCAGAGCTCTTAGTGGCGATGGAGAACCAGCACACCATC GATCTAGCCGACAGTGAGATGGACAAGCTGTACGAACGAGTCAAGAGACAGTTACGAGAGAATGCCGAGGAGGACGGAACCGGCTGCT TTGAGATTTTCCACAAGTGTGACGATGACTGTATGGCTAGCATTCGCAATAATACCTATGACCACAGTAAATACCGGGAGGAGGCTATGC AGAATCGTATTCAGATCGATCCTGTGAAGCTGTCTAGTGGATACAAGCACGTAATCCTGTGGTTTTCTTTCGGTGCTTCTTGTTTCATCCT GTTAGCAATAGTAATGGGTCTGGTATTTATCTGCGTTAAGAACGGAAATATGCGGTGTACTATTTGCATC
2834	ATGAACACACAGATACTGGTCTTTGCTCTTATCGCTATCATTCCCACCAATGCCGATAAAATCTGCTTGGGGCACCACGCCGTGTCCAAC GGGACGAAAGTTAATACACTCACTGAGCGAGGCGTGGAGGTCGTCAACGCCACAGAGACGGTGGAGCGCACCAACATCCCGAGGATCT GCAGCAAGGGCAAACGAACCGTCGATCTTGGGCAGTGCGGGCTGTTGGGCACCATTACGGGACCTCCCCAGTGTGATCAGTTCTTGGA GTTTTCTGCTGACCTCATCATTGAGAGAAGGGAAGGCAGCGACGTGTGCTATCCCGGCAAGTTCGTGAAAGAAGAGGCCCTGCGGCAGA TACTGCGGGAGTCAGGCGGTATTGATAAAGAAGCGATGGGCTTCACGTACAGCGGTATCAGGACTAACGGCGCTACCAGCGCTTGCCG ACGGAGCGGTAGCTCTTTCTACGCCGAGATGAAATGGTTACTTTCTAACACCGATAACGCTGCGTTTCCACAGATGACGAAATCCTACAA GAATACTAGAAAAAGTCCAGCCCTGATCGTTTGGGGAATTCACCACTCTGTCTCCACAGCTGAACAGACCAAGTTGTATGGCTCCGGCAA CAAGCTGGTTACAGTAGGCTCCAGCAATTACCAGCAGTCTTTTGTCCCCTCCCCTGGAGCTCGTCCACAGGTCAATGGACTGTCAGGCC GAATCGACTTCCACTGGCTGATGTTGAATCCAAACGATACAGTGACTTTCTCATTTAATGGCGCTTTTATTGCCCCAGATCGGGCCAGTTT CCTCAGAGGTAAGTCCATGGGAATCCAGTCGGGGGTCCAAGTCGATGCCAACTGCGAAGGCGACTGTTACCACAGCGGCGGTACCATT ATCTCCAACCTGCCCTTCCAAAACATCGACTCTCGGGCAGTTGGAAAATGTCCAAGGTACGTCAAGCAGAGATCCCTGCTGCTGGCCACA GGCATGAAGAACGTGCCCGAGATTCCCAAAGGACGGGGCCTATTCGGAGCGATCGCAGGATTCATTGAGAACGGATGGGAGGGCTTGA TTGATGGCTGGTACGGCTTTAGACACCAAAATGCTCAGGGGGAGGGCACTGCCGCCGACTACAAATCTACTCAGTCGGCGATTGACCAG ATTACTGGAAAGTTGAACCGGCTCATTGAAAAAACCAACCAGCAGTTCGAATTGATTGATAATGAGTTCAATGAAGTGGAAAAACAGATCG GGAATGTGATTAACTGGACCAGAGACTCCATCACAGAGGTTTGGAGCTACAACGCGGAATTACTAGTTGCCATGGAAAATCAGCACACCA TTGACCTCGCTGACAGTGAGATGGATAAACTGTACGAGCGCGTGAAAAGACAGCTACGCGAGAATGCGGAAGAAGACGGCACAGGATG CTTTGAGATCTTTCACAAGTGCGACGATGATTGCATGGCTTCGATTCGGAATAATACCTACGATCACTCCAAATATAGAGAGGAGGCTATG CAAAATCGAATTCAGATCGACCCCGTCAAGTTGTCCTCCGGCTACAAGGACGTTATCTTGTGGTTCTCCTTTGGCGCATCATGCTTCATTT TGCTGGCCATAGTCATGGGCCTGGTATTCATTTGTGTCAAGAATGGAAATATGAGGTGCACCATATGCATC
2835	ATGAATACCCAGATTCTTGTGTTCGCTCTGATCGCGATTATTCCTACCAACGCTGATAAAATCTGCCTCGGACATCACGCTGTTAGCAACG GAACCAAGGTGAACACACTGACAGAGCGCGGGGTCGAAGTTGTCAACGCTACTGAGACGGTGGAACGGACCAACATACCACGGATCTG CTCTAAGGGCAAGCGAACTGTGGATCTCGGGCAGTGTGGCTTGCTGGGAACCATTACGGGCCCTCCTCAGTGTGATCAGTTTCTCGAGT TCTCTGCAGATCTGATAATCGAGAGAAGGGAAGGATCAGATGTCTGCTACCCTGGGAAATTCGTGAAAGAAGAAGCACTGAGACAAATCT TACGCGAGTCAGGGGGCATAGATAAGGAAGCCATGGGCTTCACATACTCTGGGATCCGTACCAACGGCGCCACGTCAGCTTGTCGTAG GTCTGGCTCTTCTTTCTATGCCGAGATGAAATGGCTCCTGTCGAATACTGACAATGCTGCGTTCCCCCAGATGACTAAGAGCTACAAAAAT ACAAGAAAGTCTCCCGCGCTGATAGTTTGGGGGATTCACCACTCTGTCTCCACTGCCGAACAGACAAAGTTATACGGCAGCGGTAACAA GCTAGTGACAGTTGGGAGCTCAAACTATCAACAATCATTTGTACCATCCCCCGGTGCCAGGCCACAGGTAAACGGCCTTTCCGGACGTAT CGACTTTCACTGGCTGATGCTGAATCCCAATGATACCGTAACTTTTTCCTTCAACGGGGCTTTTATTGCCCCAGATAGGGCATCCTTTCTG CGAGGTAAGAGTATGGGCATACAGAGCGGCGTGCAAGTGGACGCTAATTGTGAGGGGGACTGCTACCATTCCGGGGGAACCATCATCT CGAATTTACCGTTTCAGAACATCGATTCACGCGCTGTTGGGAAGTGTCCTAGGTATGTAAAACAGCGATCCCTTCTTCTGGCCACAGGCA TGAAGAACGTACCAGAGATACCTAAAGGACGGGGACTGTTCGGGGCCATTGCAGGGTTCATCGAAAACGGCTGGGAAGGCCTGATCGA CGGCTGGTATGGCTTTCGGCATCAGAACGCCCAGGGAGAGGGTACTGCAGCCGATTATAAGAGTACACAGTCTGCAATCGACCAAATTA CAGGAAAGCTGAATAGACTCATAGAGAAGACAAATCAGCAGTTTGAACTCATTGACAATGAGTTCAATGAAGTGGAAAAGCAAATCGGGA ACGTGATTAATTGGACCAGGGATAGCATAACAGAAGTCTGGTCCTATAACGCTGAACTACTTGTTGCAATGGAGAACCAGCACACTATCG ATCTGGCAGACTCTGAAATGGACAAACTGTATGAGAGGGTTAAACGCCAACTGAGAGAGAATGCTGAGGAAGACGGTACTGGCTGTTTT GAGATTTTCCACAAATGTGACGACGATTGTATGGCGTCCATCAGGAACAACACCTATGACCATTCTAAATATAGAGAGGAAGCCATGCAG AATCGGATACAGATTGATCCTGTGAAACTTAGCTCTGGATATAAGGACGTGATCCTGTGGTTTTCATTTGGAGCGTCATGCTTTATCCTGC TGGCCATCGTGATGGGATTGGTCTTCATATGTGTCAAGAATGGCAATATGCGGTGTACCATATGTATT

2836	ATGAATACACAGATACTCGTGTTTGCCCTCATTGCCATCATCCCTACCAATGCAGATAAGATCTGCCTCGGCCACCATGCTGTTAGCAACG GAACCAAGGTTAACACCCTCACTGAACGAGGAGTGGAGGTCGTTAACGCCACAGAAACCGTGGAACGGACGAACATCCCCCGCATCTG CAGCAAAGGCAAGCGCACCGTGGATCTGGGGCAGTGCGGCCTGCTGGGCACGATTACAGGGCCCCCCCAATGCGACCAGTTTTTGGAG TTCAGCGCAGACCTGATCATTGAAAGGCGTGAGGGGAGTGACGTGTGTTATCCCGGTAAATTTGTGAACGAAGAGGCCCTCAGACAGAT CCTGCCGGGAGTCTGGTGGGATCGATAAAGAGGCAATGGCGTTCACTTATTCTGGAATCCGGACCAACGGGGCCACCTCTGCGTGCAGG CGTTCCGGCAGCTCATTCTATGCAGAGATGAAATGGCTACTTAGTAACACGGACAACGCAGCTTTCCCCCAGATGACTAAAAGTTACAAG AACACTCGAAAGAGTCCAGCACTCATTGTGTGGGGGATTCATCATTCCGTGTCTACCGCCGAGCAGACTAAACTATACGGTAGTGGGAAT AAACTAGTAACCGTCGGCTCTTCTAACTACCAGCAGTCTTTTGTGCCCTCTCCAGGTGAACGCCCCCAGGTCAACGGGCTGTCTGGTCG CATCGACTTTCATTGGCTGATGCTCAACCCTAATGACACAGTCACGTTCAGTTTCAATGGTGCTTTCATTGCACCCGATAGGGCCTCATTT CTTCGGGGCAAGTCCATGGGCATCCAAAGTGGAGTTCAGGTGGACGCCAATTGCGAAGGTGACTGCTATCACAGTGGCGGCACAATCAT TTCTAACCTGCCGTTTCAGAATATTGACTCCCGTGCTGTAGGGAAGTGTCCCCGATACGTGAAGCAGCGGTCCTTGCTTCTGGCAACCGG TATGAAGAATGTCCCTGAGATTCCTAAAGGCCGGGGGCTGTTCGGAGCTATTGCAGGGTTCATCGAAAACGGCTGGGAGGGGCTCATCG ATGGTTGGTACGGATTCAGACATCAAAATGCACAAGGGGAAGGAACCGCAGCCGACTATAAGAGCACACAAAGCGCTATCGATCAGATC ACCGGGAAATTAAACCGCCTGATTGAGAAAACCAATCAGCAGTTCGAGCTTATTGACAACGAATTCAATGAGGTGGAAAAACAGATAGGG AATGTGATCAACTGGACCAGAGATTCTATCACCGAGGTTTGGTCTTATAACGCGGAACTGCTGGTTGCTATGGAGAATCAGCATACCATC GACCTGGCTGACAGCGAGATGGATAAGTTATACGAGCGTGTCAAACGACAGCTGAGGGAGAATGCGGAGGAGGATGGAACCGGGTGTT TCGAGATCTTCCATAAATGTGACGATGACTGCATGGCCAGTATTCGGAACAATACATATGACCATTCCAAATACCGCGAAGAGGCAATGC AGAATCGAATACAGATAGATCCTGTGAAACTGAGCAGCGGGTATAAAGACGTCATCTTGTGCTTTAGTTTTGGGGCATCCTGCTTTATTCT GCTCGCCATAGTGATGGGGTTAGTGTTTATATGTGTTAAAAACGGAAATATGCGGTGTACAATTTGTATC
2837	ATGAATACCCAGATTCTTGTGTTTGCCCTGATCGCTATAATACCTACTAATGCAGATAAGATCTGTTTAGGACATCACGCCGTCTCCAATG GAACTAAGGTAAACACGCTGACCGAGCGGGGCGTCGAGGTGGTGAACGCAACCGAAACTGTTGAGCGCACCAATATTCCCCGGATATG CTCAAAGGGGAAGCGAACCGTTGACCTAGGCCAATGCGGCTTGCTGGGGACAATTACCGGACCACCCCAATGTGACCAGTTTCTGGAAT TTTCAGCCGACCTAATCATCGAGCGACGCGAGGGCAGCGACGTGTGTTATCCCGGTAAGTTTGTCAATGAGGAAGCCCTGAGGCAGATA CTTAGAGAATCGGGCGGGATCGACAAGGAGGCCATGGGGTTCACATATAGTGGTATCCGCACTAATGGAGCTACATCCGCCTGTCGGAG GAGCGGGAGCTCCTTCTACGCTGAAATGAAGTGGCTCCTCAGTAACACGGATAACGCTGCTTTCCCCCAGATGACTAAAAGCTATAAAAA CACTAGAAAGAGCCCAGCACTCATTGTGTGGGGTATTCACCACTCAGTGTCTACGGCCGAACAAACAAAATTATATGGATCCGGGAACAA GCTGGTAACCGTCGGAAGTAGCAATTATCAGCAGTCCTTTGTGCCAAGCCCGGGAGCCCGGCCCCAGGTGAACGGCCTATCTGGGCGG ATCGATTTTCATTGGCTTATGCTCAACCCTAACGACACCGTTACCTTTAGCTTCAACGGGGCTTTCATTGCACCCGACCGTGCATCATTCC TGCGTGGGAAGAGCATGGGGATACAGAGCGGCGTACAGGTGGACGCGAACTGCGAAGGCGACTGTTACCATTCAGGCGGCACCATTAT TTCCAATTTGCCGTTTCAGAACATCGACTCACGCGCCGTAGGAAAATGTCCCCGGTACGTCAAACAGCGCAGCCTCCTACTAGCCACCG GGATGAAAAACGTGCCCGAGATCCCCAAGGGACGGGGACTATTCGGAGCGATAGCCGGGTTCATTGAGAATGGATGGGAGGGTCTGAT CGATGGCTGGTACGGGTTCCGGCATCAAAACGCACAGGGCGAGGGGACTGCAGCCGATTACAAATCTACCCAATCGGCAATCGACCAG ATTACAGGCAAACTGAATAGGATCATAGAGAAAACCAATCAGCAGTTCGAACTGATTGACAATGAGTTTAATGAAGTAGAGAAGCAGATTG GGAATGTGATAAATTGGACTCGAGACAGCATTACCGAAGTCTGGTCTTACAATGCCGAGTTACTCGTCGCTATGGAGAACCAGCACACCA TTGACCTGGCCGACTCAGAGATGGACAAACTCTATGAGCGCGTAAAGAGACAATTGAGGGAAAACGCGGAGGAAGACGGAACAGGCTG TTTCGAAATATTTCACAAGTGCGACGACGACTGTATGGCATCCATCCGGAATAATACCTATGATCACAGCAAGTACAGGGAGGAGGCTAT GCAAAATCGGATCCAGATTGATCCTGTAAAACTGAGCTCGGGATACAAGGATGTTATCCTGTGGTTTAGTTTCGGCGCCAGCTGCTTTATT CTCCTTGCCATCGTCATGGGCCTGGTCTTCATCTGCGTCAAAAATGGAAATATGAGATGCACAATTTGCATT
2838	ATGAACACTCAAATTCTCGTGTTTGCTCTAATTGCCATCATCCCGACAAATGCAGATAAGATTTGCCTGGGTCACCATGCGGTGTCTAATG GAACAAAGGTTAATACCCTGACCGAAAGGGGCGTGGAGGTCGTGAATGCCACCGAGACGGTGGAAAGAACCAACATTCCAAGAATTTGC TCCAAAGGAAAGAAAACCGTCGACCTGGGTCAGTGCGGATTGCTGGGAACAATCACCGGGCCACCCCAGTGCGACCAGTTCCTGGAATT CAGCGCCGACCTCATTATTGAGCGCAGAGAAGGTTCTGACGTGTGTTATCCAGGGAAGTTCGTGAACGAGGAGGCACTGCGGCAAATCC TGCGTGAAAGCGGCGGCATTGACAAGGAGGCGATGGGCTTTACATACTCCGGGATCCGTACAAACGGGGCCACTAGTGCGTGCAGAAG ATCTGGAAGCAGCTTTTATGCTGAAATGAAATGGTTGCTCTCCAACACGGACAACGCTGCCTTCCCCCAGATGACAAAGAGCTACAAAAA TACTCGAAAAAGCCCCGCCCTAATAGTGTGGGGCATCCACCATTCTGTGAGCACCGCCGAACAGACCAAGCTGTATGGCAGCGGTAACA AATTGGTTACCGTAGGCTCTTCTAACTATCAGCAGAGCTTCGTCCCATCGCCTGGTGCCAGGCCCCAGGTCAACGGCCAAAGCGGTCGG ATTGATTTTCACTGGTTAATGCTTAATCCAAATGATACAGTGACCTTTAGTTTTAACGGAGCTTTCATCGCCCCCGATCGGGCGAGCTTTC TCCGGGGGAAATCCATGGGAATCCAGTCTGGGGTGCAGGTGGACGCTAATTGCGAGGGCGACTGCTACCACTCCGGGGGGACCATCAT AAGTAACCTCCCTTTCCAAAACATCGATTCACGAGCTGTGGGAAAATGCCCCCGTTACGTAAAACAGCGTTCTCTCTTATTGGCCACAGGT ATGAAAAATGTGCCAGAAATTCCTAAGGGCCGGGGGCTCTTCGGCGCCATAGCCGGATTCATTGAGAATGGGTGGGAAGGTCTGATCGA CGGGTGGTACGGTTTCCGACACCAGAACGCACAGGGAGAGGGTACGGCTGCAGATTACAAGTCTACTCAGTCAGCTATTGACCAGATCA CCGGCAAGCTGAATAGGCTTATCGAGAAAACAAACCAACAGTTTGAACTGATCGACAATGAGTTTAACGAGGTCGAGAAGCAGATCGGCA ACGTTATTAACTGGACTAGAGACTCCATCACCGAAGTGTGGAGCTACAATGCCGAGCTGCTAGTGGCTATGGAGAATCAACATACCATTG ATCTGGCCGATAGTGAGATGGACAAGCTGTATGAACGAGTCAAGCGACAACTACGCGAAAATGCGGAAGAACATGGGACAGGATGTTTT GAGATTTTCCACAAATGCGATGACGATTGTATGGCCAGCATCCGGAACAACACATACGATCATAGCAAATACAGAGAGGAGGCCATGCAA AATCGCATTCAAATAGACCCTGTCAAGCTTAGCTCCGGGTATAAAGACGTGATTCTATGGTTCTCGTTTGGGGCCTCATGTTTCATTCTGC TGGCCATCGTCATGGGCCTCGTGTTTATTTGCGTAAAGAATGGCAACATGCGCTGTACCATCTGTATT

2839	ATGAACACTCAGATTCTGGTGTTTGCTCTGATCGCGATTATCCCCACTAATGCCGACAAGATATGTCTCGGTCATCATGCGGTTTCTAACG GGACGAAGGTGAACACATTGACAGAGCGAGGGGTGGAGGTGGTTAATGCGACTGAGACTGTAGAAAGGACCAATATTCCGCGTATTTGT AGCAAGGGTAAGAAAACTGTTGACTTGGGACAATGTGGTCTGCTCGGAACCATTACTGGCCCCCCTCAGTGTGATCAATTCCTGGAGTTC AGCGCCGATCTGATTATAGAGCGAAGAGAGGGTAGCGATGTTTGCTATCCCGGGAAATTTGTAAACGAGGAGGCCCTCCGCCAAATCCT ACGAGAGTCAGGTGGGATTGACAAGGAGGCTATGGGGTTCACCTATAGTGGAATCCGGACCAATGGGGCAACTTCTGCCTGTCGGAGAT CTGGCTCGAGCTTCTACGCGGAGATGAAGTGGTTGCTGAGTAACACTGATAACGCTGCGTTTCCCCAAATGACCAAGTCGTATAAAAATA CAAGAAAGAGTCCTGCTCTGATCGTGTGGGGAATCCATCACTCTGTAAGCACGGCCGAGCAAACTAAACTTTATGGATCGGGCAATAAAC TTGTCACAGTGGGCAGCAGCAACTACCAGCAATCATTTGTGCCTTCTCCGGGCGCGAGACCACAGGTCAATGGCCAGTCTGGGAGGATC GATTTTCACTGGCTCATGCTAAATCCCAACGACACAGTGACCTTTAGCTTTAATGGCGCATTTATTGCACCTGATAGAGCGTCATTCCTTA GGGGAAAAAGCATGGGGATACAATCAGGGGTCCAGGTCGACGCTAATTGCGAGGGCGACTGCTACCACTCAGGGGGTACCATCATATC TAATCTTCCATTCCAGAACATCGACTCGCGAGCCGTTGGAAAGTGCCCCAGGTACGTGAAGCAGCGGAGTCTGCTCCTGGCAACTGGCA TGAAGAACGTGCCTGAGATCCCCAAAGGCCGTGGCTTATTTGGAGCTATCGCAGGGTTTATCGAGAATGGCTGGGAGGGATTGATCGAT GGTTGGTATGGCTTTCGGCACCAAAATGCCCAGGGAGAGGGAACCGCCGCCGATTATAAGTCAACTCAGAGTGCGATAGATCAGATTAC AGGTAAGCTAAATAGACTGATTGAGAAGACTAACCAACAGTTCGAGCTTATCGACAACGAATTTAATGAAGTGGAAAAGCAAATAGGAAAC GTTATCAACTGGACACGCGATAGCATCACAGAAGTTTGGTCTTACAACGCCGAGCTCTTGGTGGCTATGGAGAATCAACATACAATCGAC CTGGCAGATAGCGAAATGGATAAGTTATACGAACGGGTCAAGCGACAGCTGAGGGAGAACGCTGAAGAGGATGGAACTGGCTGTTTTGA GATCTTCCACAAGTGCGATGATGACTGTATGGCAAGCATACGAAACAACACTTACGATCATTCAAAGTACCGGGAGGAGGCCATGCAAAA CAGGATTCAGATTGATCCAGTGAAGTTGTCCTCTGGCTACAAGGATGTGATACTGTGGTTTTCCTTCGGAGCAAGCTGCTTTATCTTACTG GCAATTGTGATGGGACTCGTGTTTATATGCGTGAAAAATGGAAATATGCGCTGTACAATCTGCATC
2840	ATGAACACGCAGATACTTGTGTTTGCCCTTATTGCCATTATTCCCACTAACGCGGACAAAATATGCCTGGGCCATCACGCTGTCTCCAATG GGACGAAAGTGAACACATTGACCGAGCGAGGCGTGGAAGTGGTAAATGCGACCGAAACCGTAGAGAGAACCAACATTCCTAGAATTTGC TCCAAGGGCAAACGGACAGTAGATCTGGGGCAGTGCGGACTGTTGGGTACAATTACAGGACCTCCCCAGTGCGATCAGTTCCTGGAGTT TTCCGCTGATCTGATTATAGAAAGGCGAGAGGGTTCCGACGTGTGCTACCCTGGTAAGTTTGTTAAAGAGGAAGCTCTCCGTCAGATTCT CCGAGAGTCTGGCGGGATCGATAAAGAGGCGATGGGGTTTACCTACTCTGGAATAAGGACCAACGGAGCAACCTCCGCCTGCAGGAGA TCAGGGTCGTCCTTCTATGCTGAAATGAAATGGTTGCTGTCTAATACCGACAATGCAGCCTTCCCCCAGATGACCAAAAGCTACAAGAAT ACACGCAAATCTCCCGCCCTGATTGTGTGGGGCATACACCATAGTGTGAGTACTGCAGAACAGACCAAGCTTTACGGATCCGGCAATAA GCTGGTGACTGTAGGCTCTAGCAACTACCAACAGAGCTTTGTACCATCCCCCGGTGCACGCCCTCAGGTGAACGGACTCTCGGGGCGG ATAGATTTCCACTGGTTGATGTTGAATCCCAACGACACCGTGACGTTCAGTTTCAACGGTGCTTTCATCGCGCCTGATAGGGCCTCTTTCT TAAGAGGGAAATCAATGGGCATCCAGAGCGGGGTTCAGGTAGACGCTAACTGTGAGGGTGACTGCTATCATAGCGGCGGCACTATTATA TCTAACCTGCCCTTCCAGAATATCGACTCCCGGGCCGTGGGAAAATGCCCAAGGTATGTGAAACAGAGGTCACTGCTACTGGCGACCGG GATGAAGAACGTTCCAGAAATACCTAAAGGTCGTGGGCTGTTCGGCGCAATTGCCGGTTTCATTGAAAACGGCTGGGAGGGCCTGATCG ACGGATGGTATGGGTTCAGACACCAGAACGCCCAGGGAGAGGGCACAGCTGCCGACTATAAGTCCACACAAAGCGCCATTGATCAGAT CACAGGCAAGCTTAACAGACTGATCGAGAAGACAAACCAGCAATTTGAGTTGATAGACAACGAGTTCAACGAAGTGGAGAAGCAAATCG GGAACGTCATTAATTGGACCCGGGATTCTATCACCGAAGTATGGTCATACAACGCCGAGCTGCTCGTGGCTATGGAGAATCAGCATACCA TTGATCTTGCCGATTCCGAGATGGACAAGCTGTATGAAAGAGTCAAACGACAGTTGCGAGAAAACGCTGAAGAGGACGGAACAGGATGC TTTGAAATCTTTCACAAATGTGACGACGACTGTATGGCAAGCATTCGGAACAATACCTATGATCACTCAAAGTACCGAGAGGAGGCCATG CAGAATAGGATTCAGATCGACCCAGTCAAGCTCTCCTCCGGGTACAAAGATGTAATCCTGTGGTTTTCGTTCGGCGCCAGTTGTTTCATC TTACTGGCCATAGTCATGGGCCTGGTCTTTATCTGCGTTAAGAACGGCAACATGCGCTGCACTATTTGCATC
2841	ATGAACACCCAGATACTTGTTTTTGCCCTGATTGCCATCATCCCAACCAACGCCGACAAGATCTGCTTGGGGCACCATGCTGTTAGCAAC GGAACAAAAGTGAACACCCTGACCGAAAGAGGCGTCGAGGTAGTAAACGCCACTGAAACGGTCGAGCGAACCAATATACCTAGGATATG TTCAAAGGGAAAAACGACGGTTGATTTAGGTCAGTGTGGACTTCTGGGGACAATCACCGGACCACCCCAATGTGACCAGTTCCTTGAGTT CAGCGCGGACCTGATTATTGAGAGACGTGAAGGATCTGACGTCTGCTACCCAGGAAAGTTTGTGAATGAGGAAGCACTCCGCCAGATTC TCAGGGAATCCGGCGGCATTGATAAAGAGGCAATGGGATTTACCTACTCCGGCATTCGGACTAACGGTGCTACATCTGCATGTAGACGG TCCGGCTCTTCATTCTATGCAGAGATGAAATGGCTGCTTTCAAATACTGATAACGCCGCCTTTCCTCAGATGACCAAATCCTACAAGAACA CCCGGAAATCCCCCGCCCTCATTGTTTGGGGGATCCACCACTCAGTGTCAACCGCCGAGCAGACCAAGCTCTACGGAAGTGGAAATAAG TTGGTAACTGTGGGAAGCTCTAACTACCAGCAAAGCTTTGTGCCAAGTCCTGGAGAGAGGCCCCAGGTGAACGGACTTAGCGGTAGGAT CGATTTTCATTGGCTTATGCTCAACCCCAACGATACGGTCACCTTTTCATTCAATGGGGCCTTTATTGCACCGGATAGAGCCAGCTTTCTG AGAGGGAAGTCTATGGGCATCCAGTCGGGCGTACAGGTGGATGCAAACTGTGAGGGGGATTGCTATCATAGTGGAGGCACCATCATAA GTAATCTGCCCTTCCAGAATATCGATAGCCGGGCCGTGGGCAAATGTCCAAGGTATGTGAAACAGAGGTCACTGCTGCTCGCTACAGGC ATGAAAAACGTGCCTGAAATACCCAAAGGACGGGGGCTTTTCGGCGCCATCGCCGGCTTCATTGAGAACGGATGGGAAGGTTTGATCGA TGGATGGTATGGATTTAGGCACCAAAACGCCCAGGGGGAAGGCACCGCCGCGGACTACAAGAGTACTCAGTCTGCAATCGACCAAATCA CTGGCAAACTGAATCGGCTGATTGAGAAGACCAATCAGCAATTCGAACTCATTGACAACGAGTTTAATGAAGTGGAGAAACAGATCGGTA ACGTGATAAATTGGACCAGAGACTCGATCACGGAAGTATGGAGCTACAATGCTGAACTGTTGGTCGCTATGGAGAATCAACACACTATCG ACCTGCCCGACAGCGAGATGGATAAACTCTATGAAAGAGTGAAACGGCAGCTCAGAGAGAACGCAGAGGAGGACGGAACCGGCTGCTT TGAAATATTCCATAAATGCGACGACGATTGCATGGCCAGCATCAGGAACAATACCTATGATCATAGCAAGTACCGGGAGGAAGCCATGCA GAATAGAATTCAGATCGACCCAGTGAAACTGTCTAGCGGATATAAGGACGTGATTCTGTGGTTCTCATTCGGCGCCTCATGCTTCATTCTA TTGGCGATTGTGATGGGACTTGTGTTCATCTGCGTCAAAAATGGCAATATGCGGTGCACGATATGTATC

2842	ATGAACACCCAAATTCTGGTCTTTGCTCTGATCGCAATTATCCCAACCAATGCTGACAAAATTTGCCTCGGACATCATGCGGTATCCAACG GCACAAAGGTAAACACTCTCACTGAACGCGGAGTTGAGGTGGTAAATGCCACGGAAACAGTTGAGAGGACTAATATACCTAGAATTTGTA GCAAAGGAAAAAAAACGGTGGACCTCGGGCAGTGTGGCCTGCTTGGAACAATTACAGGCCCGCCCCAGTGCGACCAGTTCCTAGAATTC TCCGCCGATCTGATCATCGAACGCCGCGAAGGAAGCGATGTGTGTTATCCAGGCAAGTTTGTGAATGAGGAGGCCCTGAGACAAATCCT GCGGGAGAGCGGCGGAATTGATAAACAGGCAATGGGATTCACTTACAGTGGCATCCGGACAAACGGGGCCACGTCGGCCTGCAGACGT AGCGGGAGTTCCTTCTATGCCGAAATGAAGTGGCTGCTGTCCAACACTGACAACGCGGCCTTTCCACAAATGACTAAATCTTATAAAAAC ACACGGAAGTCACCAGCTCTCATCGTGTGGGGTATCCACCATTCGGTCTCTACAGCTGAGCAAACTAAGCTTTACGGAAGTGGCAACAAA CTGGTGACAGTGGGCAGCTCTAACTACCAGCAGTCATTTGTCCCGTCTCCCGGGGCCAGACCACAGGTTAATGGGCAGTCAGGCCGGA TCGATTTTCATTGGTTGATGTTAAACCCAAACGACACAGTCACATTTAGTTTTAATGGGGCATTTATAGCCCCTGATAGAGCTAGTTTCTTG AGGGGAAAGAGTATGGGGATCCAGTCAGGTGTACAGGTCGACGCCAACTGCGAGGGTGACTGTTACCATTCCGGGGGTACCATCATTA GTAACCTGCCTTTCCAGAACATCGACAGTAGAGCAGTGGGTAAATGCCCAAGGTACGTGAAACAACGCAGCTTACTGCTGGCTACAGGA ATGAAAAACGTGCCCGAGATTCCAAAAGGGAGGGGTTTGTTCGGTGCTATTGCCGGTTTTATTGAAAACGGATGGGAGGGTCTCATCGAT GGATGGTATGGGTTTAGGCACCAGAATGCACAAGGGGAAGGAACTGCCGCCGATTATAAATCCACCCAGTCCGCTATTGACCAGATTAC CGGTAAACTGAATCGTTTGATTGAGAAAACAAACCAGCAGTTTGAACTAATCGACAACGAATTCAATGAAGTGGAGAAGCAGATTGGTAAC GTCATTAACTGGACCCGTGACAGTATCACTGAAGTTTGGTCTTATAATGCCGAACTCCTAGTAGCGATGGAGAACCAGCACACCATAGAC CTCGCCGATAGTGAAATGGACAAACTGTATGAGAGAGTGAAGCGGCAGTTACGCGAAAACGCTGAGGAGGACGGGACTGGCTGCTTCG AGATTTTTCACAAGTGCGACGACGACTGCATGGCATCAATCCGCAACAATACATACGATCATTCCAAGTATCGTGAAGAAGCCATGCAGA ACCGGATCCAAATAGACCCTGTCAAACTTAGCAGCGGGTATAAAGATGTTATTCTGTGGTTTAGCTTTGGAGCCAGTTGTTTCATACTGCT GGCCATTGTTATGGGCCTCGTCTTCATCTGCGTGAAAAACGGGAACATGCGATGTACGATCTGCATT
2843	ATGAACACTCAAATTTTAGTCTTCGCCCTCATTGCGATAATCCCTACTAACGCAGACAAGATCTGTCTGGGACACCACGCCGTGTCTAATG GCACCAAAGTGAACACATTGACGGAGAGAGGAGTAGAGGTGGTGAACGCCACCGAGACCGTCGAGAGAACCAATATACCTAGGATCTG CAGCAAGGGAAAGAAGACTGTCGATCTGGGCCAATGCGGGTTGCTCGGGACTATTACTGGACCTCCGCAATGTGACCAGTTCCTTGAAT TCTCTGCTGACCTGATAATCGAACGACGAGAGGGATCCGATGTTTGCTATCCAGGGAAGTTTGTGAATGAGGAGGCTCTCAGACAAATCC TGCGGGAGTCCGGCGGCATCGACAAAGAAGCCATGGGTTTCACCTATAGTGGAATTCGGACCAATGGCGCAACCAGCGCATGCCGGCG TAGTGGAAGCTCCTTTTATGCCGAGATGAAGTGGTTACTTTCCAATACCGATAACGCAGCATTCCCTCAGATGACCAAAAGTTACAAGAAC ACTCGTAAATCTCCTGCTCTTATTGTATGGGGGATACACCACAGTGTGAGCACCGCCGAGCAAACTAAGCTTTACGGATCAGGGAATAAG CTTGTAACAGTGGGAAGCTCTAACTACCAACAGTCCTTTGTTCCATCTCCCGGGGCGAGACCTCAAGTAAACGGGCAATCTGGTAGGATA GACTTTCATTGGCTGATGCTGAATCCGAATGATACAGTTACCTTCTCCTTTAATGGAGCCTTTATCGCCCCCGACAGAGCGAGTTTTCTGA GGGGGAAGTCTATGGGGATCCAGTCCGGCGTTCAGGTAGACGCGAATTGCGAAGGGGATTGCTATCACAGCGGCGGTACAATTATCTCT AACCTCCCTTTCCAGAATATCGACTCCCGAGCCGTTGGGAAGTGCCCTCGGTATGTTAAGCAACGGAGCCTGCTGCTGGCTACCGGGAT GAAAAACGTGCCTGAGATTCCTAAAGGCCGGGGCCTATTTGGGGCCATCGCTGGTTTTATAGAGAACGGCTGGGAAGGTCTGATCGATG GCTGGTATGGTTTTAGACATCAGAATGCCCAGGGTGAAGGGACCGCCGCCGACTACAAGAGTACCCAATCTGCCATCGACCAGATCACC GGGAAGCTGAACCGTCTCATCGAGAAGACGAACCAGCAGTTCGAACTGATTGATAATGAGTTCAATGAAGTGGAGAAGCAGATCGGTAA CGTTATCAACTGGACGCGCGACAGCATAACCGAGGTGTGGAGTTATAACGCTGAACTCTTGGTTGCCATGGAGAACCAGCATACCATCG ATCTCGCCGATAGTGAGATGGATAAGCTGTATGAACGGGTGAAAAGGCAACTGCGCGAGAACGCCGAGGAAGACGGCACCGGCTGCTT TGAAATCTTTCACAAATGCGATGATGATTGTATGGCATCTATCAGGAACAATACCTACGATCACAGCAAATACCGCGAGGAAGCCATGCA GAATAGAATCCAGATAGATCCCGTGAAGCTGTCCAGCGGGTATAAGGACGTCATCTTGTGGTTCAGTTTTGGGGCTTCGTGTTTTATTCT GTTGGCTATCGTCATGGGCCTCGTGTTCATCTGCGTCAAGAACGGGAACATGCGCTGCACCATCTGCATA
2844	ATGAACACTCAAATTTTAGTCTTTGCATTAATTGCCATTATCCCAACCAACGCAGACAAAATCTGCCTGGGACATCACGCCGTGTCAAACG GCACCAAAGTGAACACGCTGACCGAACGTGGGGTCGAAGTGGTGAATGCCACAGAGACTGTCGAGCGCACAAATATCCCGCGCATTTGT TCCAAGGGTAAGCGGACGGTGGACTTAGGTCAGTGTGGACTTCTACGTACTATTACCGGACCTCCACAGTGTGACCAGTTCCTGGAGTT CTCTGCCGATCTCATCATTGAGCGTCGCGAAGGAAGTGATGTCTGTTACCCAGGAAAATTCGTGAATGAAGAAGCCCTCCGGCAAATTCT GCGCGAGTCCGGGGGCATCGACAAGGAGGCGATGGGCTTCACCTATAGTGGCATCCGAACAAACGGCGCCACCTCCGCATGCCGGAG GAGTGGCAGCTCCTTCTACGCCGAGATGAAGTGGCTCTTATCCAACACAGACAATGCCGCTTTCCCGCAGATGACAAAGAGTTACAAAAA CACGCGGAAAAGTCCCGCTTTGATTGTGTGGGGTATCCACCACTCTGTCTCTACAGCGGAGCAGACAAAGTTATACGGGTCTGGCAGTA AACTGGTTACAGTGGGCTCAAGCAACTACCAACAGTCTTTCGTCCCCTCCCCTGGCGCAAGACCCCAGGTAAATGGCCTCTCTGGAAGG ATTGACTTCCACTGGCTGATGCTCAACCCAAATGATACCGTGACATTTAGCTTCAACGGTGCATTTATCGCCCCAGACAGAGCTAGTTTCC TTCGGGGCAAATCAATGGGGATTCAGAGCGGAGTGCAGGTGGATGCTAATTGTGAAGGCGATTGCTACCACTCCGGCGGTACTATTATC TCAAACCTGCCTTTTCAGAATATCGATAGTAGAGCCGTGGGCAAGTGCCCGCGTTATGTGAAACAGCGCTCTCTTTTGCTCGCTACAGGA ATGAAAAACGTACCAGAGATCCCCAAGGGGCGTGGCCTATTTGGAGCTATCGCTGGTTTTATTGAAAACGGGTGGGAGGGCCTGATAGA CGGTTGGTATGGCTTCAGACACCAGAATGCTCAGGGAGAGGGCACCGCCGCCGACTATAAATCAACACAGAGTGCGATCGACCAGATTA CCGGAAAACTGAATCGCCTCATAGAGAAGACAAACCAACAGTTCGAGCTCATCGACAACGAATTTAACGAGGTAGAAAAACAGATCGGTA ACGTTATCAACTGGACACGGGACAGCATCACGGAGGTGTGGTCTTACAATGCCGAGCTTTTGGTGGCTATGGAGAACCAGCACACAATC GACCTCGCCGATAGTGAAATGGATAAACTTTATGAGAGCGTAAAGCGTCAGCTGCGAGAGAATGCTGAAGAAGATGGGACAGGTTGCTT TGAGATCTTCCACAAGTGCGATGACGATTGCATGGCCTCTATTAGAAATAACACTTACGATCATAGCAAATATAGAGAGGAAGCTATGCAG AATAGGATCCAGATAGATCCCGTAAAGCTGTCCTCTGGATATAAGGACGTGATCCTCTGGTTCAGTTTTGGGGCATCATGTTTTATTCTGC TCGCAATCGTTATGGGTTTAGTTTTCATTTGTGTCAAATCACGGAATATGAGATGTACAATCTGTATT

2845	ATGAATACACAGATCCTGGTGTTCGCACTGATCGCTATCATCCCCACAAACGCTGACAAAATCTGTCTGGGGCACCACGCGGTGAGTAAC GGCACTAAGGTGAATACACTGACCGAACGCGGGGTAGAGGTGGTGAATGCTACAGAGACCGTTGAGCGCACTAACATCCCCAGAATATG TAGTAAAGGTAAGAAAACCGTCGACCTGGGCCAGTGTGGCCTGCTGGGCACAATTACCGGTCCACCCCAATGCGATCAGTTCTTAGAGT TCTCAGCAGATCTAATTATCGAGCGCCGCGAGGGGAGCGACGTCTGCTACCCTGGGAAATTTGTCAACGAAGAGGCCCTTCGGCAAATC CTTCGAGAGAGTGGCGGCATTGACAAGGAGGCAATGGGGTTTACTTATAGCGGAATCCGCACCAATGGAGCAACCTCTGCATGCAGGA GAAGCGGCAGTTCCTTCTATGCCGAAATGAAATGGCTGCTGAGCAATACAGATAACGCTGCCTTCCCACAGATGACCAAGAGTTACAAGA ATACTAGGAAGTCCCCAGCCCTGATTGTTTGGGGAATTCACCACTCCGTTAGTACCGCCGAGCAAACTAAACTGTACGGATCTGGAAACA AATTAGTTACAGTCGGAAGTTCTAACTACCAACAGTCATTTGTTCCATCTCCTGGTGCTAGGCCGCAGGTCAATGGCCAATCCGGCCGAA TAGACTTTCACTGGCTTATGCTTAATCCGAACGACACAGTGACCTTTAGCTTCAATGGCGCCTTCATTGCCCCTGACAGAGCAAGCTTCCT GAGAGGCAAGAGCATGGGGATTCAGAGCGGTGTCCAGGTGGACGCTAACTGTGAGGGTGACTGCTATCACTCCGGTGGCACTATAATC TCTAACCTACCCTTTCAGAATATCGACAGTCGGGCTGTGGGCAAGTGCCCCCGGTATGTTAAACAGCGTAGCCTTTTGCTGGCTACTGGG ATGAAAAACGTGCCCGAGATACCCAAGGGGCGCGGACTCTTCGGGGCCATCGCGGGCTTCATCGAGAATGGTTGGGAAGGTCTCATCG ATGGCTGGTACGGATTCAGGCATCAGAACGCACAGGGTGAGGGGACAGCGGCTGACTACAAGTCGACCCAGAGTGCTATCGATCAGAT AACTGGCAAGCTCAACCGGCTCATAGAGAAAACGAACCAGCAATTCGAGCTGATTGACAACGAATTCAACGAGGTTGAGAAGCAGATTG GCAATGTGATCAACTGGACGAGAGACAGCATTACCGAGGTGTGGAGTTACAACGCCGAGCTACTGGTAGCAATGGAGAATCAACATACC ATTGATTTGGCGGATTCTGAGATGGACAAGTTGTACGAGAGGGTCAAGCGGCAGCTGCGCGAAAACGCAGAAGAGGACGGCACTGGCT GCTTCGAGATTTTTCACAAATGTGATGACGATTGCATGGCGTCTATCAGGAATAACACATATGACCACAGCAAATACCGGGAGGAAGCGA TGCAGAACCGGATCCAAATTGATCCAGTCAAGCTGTCAAGCGGCTACAAGGATGTGATCTTATGGTTCAGTTTCGGAGCAAGTTGCTTTA TTCTCCTTGCTATCGTTATGGGACTTGTCTTCATCTGTGTTAAAAACGGTAATATGCGGTGTACCATTTGTATT
2846	ATGAACACCCAGATCCTGGTATTTGCTCTGATTGCCATCATTCCTACCAATGCCGACAAAATCTGTCTGGGCCACCACGCAGTTTCGAATG GCACTAAAGTGAATACCCTGACCGAGCGCGGCGTCGAGGTCGTGAATGCCACAGAGACCGTCGAACGCACAAATATACCTCGGATTTGC AGCAAGGGAAAGCGGACAGTGGACCTCGGGCAGTGCGGACTGCTCGGGACTATAACTGGACCGCCTCAGTGCGATCAATTTTTGGAGT TTTCCGCTGATTTGATCATCGAGAGAAGAGAAGGAAGCGACGTCTGCTATCCAGGAAAGTTCGTGAACGAGGAGGCCCTGAGGCAAATC TTGAGGGAGAGTGGTGGCATCGACAAAGAGGCTATGGGTTTCACCTACTCAGGTATACGCACCAACGGCGCAACAAGTGCTTGTCGGCG GTCAGGATCTTCCTTCTACGCCGAGATGAAATGGTTGCTAAGTAATACAGACAATGCTGCGTTCCCTCAGATGACAAAGTCCTATAAGAAT ACACGGAAGTCACCTGCCCTGATCGTGTGGGGCATTCACCACAGCGTTAGCACAGCTGAGCAGACCAAGTTATATGGCAGTGGTAACAA ACTCGTAACGGTGGGGAGTTCCAATTACCAACAGAGCTTCGTGCCGTCTCCTGGAGCTCGTCCACAGGTCAACGGACTGAGTGGGCGC ATTGACTTCCATTGGCTTATGTTAAACCCTAATGACACCGTGACGTTTAGCTTCAACGGCGCTTTTATCGCCCCTGACCGGGCCTCATTTC TGAGGGGAAAGAGCATGGGGATCCAGTCCGGAGTGCAGGTGGATGCTAATTGCGAAGGGGACTGCTATCACAGCGGAGGCACCATCAT TTCGAATCTCCCATTCCAGAACATCGACTCTAGGGCTGTGGGCAAGTGCCCTCGATATGTGAAACAGCGGAGCTTGCTGCTTGCTACCG GGATGAAGAACGTGCCGGAGATCCCCAAGGGCCGTGGCTTATTTGGCGCTATCGCGGGTTTCATCGAAAATGGTTGGGAGGGCCTAATA AACGGCTGGTACGGTTTCCGCCACCAGAACGCGCAGGGCGAGGGGACCGCCGCCGATTACAAATCCACACAGAGCGCAATCGATCAAA TAACTGGAAAGCTGAATCGCTTGATCGAAAAGACCAATCAGCAGTTTGAGCTAATCGATAACGAATTTAATGAAGTGGAAAAGCAAATCGG AAACGTGATCAATTGGACCAGAGACTCGATAACAGAGGTGTGGAGTTACAATGCAGAGCTGCTGGTAGCCATGGAAAATCAGCACACGA TTGACCTGGCAGATTCAGAGATGGATAAGCTGTACGAGAGGGTGAAGAGGCAGCTCCGTGAGAATGCTGAAGAGGATGGAACCGGGTG TTTCGAGATTTTTCACAAATGCGACGACGATTGTATGGCCTCCATCAGGAATAACACATACGACCATAGCAAATACAGACGAGGAGGCAAT GCAAAATCGGATTCAAATTGATCCCGTGAAACTTTCTAGCGGGTACAAGGATGTTATCTTGTGGTTCAGTTTCGGCGCCTCCTGTTTTATT CTGCTGGCAATTGTGATGGGTCTGGTGTTTATTTGTGTAAAGAACGGGAATATGAGGTGCACAATTTGCATC
2847	ATGAACACTCAGATACTGGTTTTCGCTTTGATCGCTATCATACCCACAAATGCCGACAAGATATGCTTGGGTCATCACGCCGTGTCCAACG GCACTAAAGTGAACACATTGACCGAGAGAGGAGTAGAAGTGGTCAATGCTACAGAGACCGTGGAACGCACTAATATCCCGCGGATATGT AGTAAGGGCAAAAAAACCGTGGACCTCGGCCAGTGTGGCCTGCTGGGCACCATTACCGGCCCGCCACAATGTGATCAGTTCTTGGAATT CTCTGCAGACCTTATCATAGAGCGCCGGGAAGGATCTGACGTGTGTTATCCAGGCAAGTTCGTTAACGAGGAGGCCCTTAGGCAGATTT TGCGTGAGAGCGGAGGCATCGATAAAGAAGCTATGGGGTTTACATATAGCGGTATTCGAACAAACGGAGCCACCTCTGCATGCAGGCGG TCCGGGTCATCATTCTACGCGGAGATGAAGTGGCTGCTCTCAAATACGGACAATGCGGCATTTCCCCAGATGACAAAAAGTTACAAAAAC ACAAGGAAATCCCCAGCCCTGATCGTGTGGGGAATTCACCACAGTGTTAGCACCGCTGAGCAGACCAAACTCTATGGCTCTGGCAACAA ATTAGTTACCGTCGGATCAAGTAATTACCAGCAGTCTTTTGTCCCGAGCCCAGGGGCTAGACCTCAGGTGAATGGACAGTCCGGAAGGA TCGACTTCCACTGGCTGATGCTTAATCCAAACGATACAGTTACTTTCTCATTCAATGGCGCCTTCATCGCTCCCGATAGAGCTTCTTTCTT GCGAGGCAAATCAATGGGCATCCAGAGCGGGGTGCAGGTCGATGCAAATTGTGAGGGTGATTGCTATCATTCAGGCGGTACCATCATAT CCAATCTCCCTTTCCAAAACATTGATTCCCGGGCAGTGGGAAAATGTCCCCGCTATGTGAAACAGAGGTCACTTCTATTAGCCACAGGGA TGAAAAATGTGCCAGAGATTCCCAAAGGAAGGGGCCTCTTTGGCGCAATCGCAGGTTTCATCGAGAATGGTTGGGAGGGTCTGATCGAC GGCTGGTACGGCTTCAGGCACCAGAACGCTCAGGGGGAGGGCACCGCAGCCGATTATAAAAGCACCCAAAGTGCTATTGATCAGATCA CAGGCAAACTGAACCGGCTTATAGAGAAGACAAATCAGCAGTTTGAGTTAATAGACAACGAATTCAACGAGGTGGAGAAACAGATTGGAA ACGTTATAAACTGGACTCGGGACTCTATTACCGAGGTTTGGTCTTACAACGCCGAACTCCTAGTGGCCATGGAGAATCAGCATACCATCG ATCTCGCTGATTCTGAAATGGACAAGCTGTACGAGAGGGTTAAACGGCAGCTGCGACAGAATGCAGAGGAAGACGCAACAGGCTGCTTC GAGATCTTCCACAAATGCGATGACGATTGTATGGCTTCCATCCGCAATAACACCTACGACCACAGCAAGTATCGGGAAGAAGCCATGCAG AACCGCATTCAAATTGACCCGGTAAAACTTTCCAGCGGATACAAGGACGTCATCCTGTGGTTTAGTTTCGGTGCGAGCTGCTTCATCTTA CTAGCTATCGTGATGGGCCTCGTGTTCATCTGCGTTAAAAACGGCAACATGCGTTGTACCATATGTATT

2848	ATGAACACACAGATCCTCGTTTTCGCTCTGATAGCAATTATCCCAACAAACGCAGACAAAATCTGTCTGGGGCACCATGCCGTGAGTAAC GGCACTAAGGTCAATACACTGACAGAGCGCGGCGTGGAGGTTGTCAATGCTACAGAAACTGTTGAGAGAACGAACATACCTCGGATTTG CAGCAAGGGCAAAAAGACCGTCGACCTGGGCCAGTGTGGCTTGCTCGGAACCATCACCGGCCCACCCCAATGCGACCAGTTCCTTGAA TTCAGTGCGGACCTCATCATCGAAAGGAGAGAAGGAAGTGATGTGTGCTATCCCGGGAAGTTCGTGAACGAAGAGGCCCTGAGGCAGAT TCTTCGGGAGTCTGGTGGTATCGAAAAGGACGCCATGGGATTCACCTACAGCGGCATTCGCGCCAACGGTGCAACTTCTGCATGTCGCC GCTCTGGGTCTTCTTTTTACGCCGAAATGAAGTGGCTGCTTTCTAATACAGATAATGCGGCATTCCCACAAATGACCAAGTCATATAAAAA CACGAGAAAGAGTCCCGCCCTTATAGTGTGGGGGATTCACCACTCCGTCTCCACAGCTGAACAGACAAAATTATACGGCAGTGGGAATA AGCTTGTTACTGTCGGCTCCAGCAATTATCAGCAAAGCTTCGTGCCATCCCCAGGGGCCAGACCACAGGTAAACGGGCTCAGCGGACGC ATCGATTTTCATTGGTTGATGCTAAACCCAAACGACACCGTAACATTTAGTTTCAACGGTGCCTTCATAGCGCCCGATCGGGCATCGTTCC TGCGGGGAAAATCCATGGGCATTCAAAGCGGCGTGCAGGTGGACGCGAATTGCGAAGGTGACTGTTATCATAGCGGCGGGACCATTAT AAGCAATCTGCCTTTTCAGAACATCGACTCACGGGCCGTGGGCAAATGCCCTCGTTACGTGAAACAACGAAGCCTGCTGCTGGCGACCG GAATGAAAAATGTCCCGGAAATACCTAAAGGAAGGGGGCTCTTCGGGGCTATAGCCGGTTTTATCGAGAACGGCTGGGAGGGCCTGATC GATGGATGGTACGGGTTCAGGCACCAGAACGCCCAAGGGGAAGGCACGGCCGCGGACTATAAGTCAACTCAGTCTGCCATTGACCAGA TTACTGGGAAGCTCAATCGCCTGATCGAAAAGACCAACCAGCAATTTGAGTTGATCGACAATGAGTTTAATGAAGTGGAAAAACAGATCG GGAATGTGATCAACTGGACCAGGGACTCTATTACCGAGGTTTGGTCCTACAACGCAGAGTTACTCGTAGCCATGGAAAACCAACACACCA TTGACCTCGCCGATAGTGAGATGGATAAACTGTATGAACGCGTCAAGCGGCAGTTGCGAGAAAATGCAGAAGAGGATGGCACAGGGTGT TTTGAGATTTTCCATAAGTGCGACGATGACTGTATGGCCTCCATCCGCAATAACACCTACGATCACTCAAAGTACCGGGAAGAGGCCATG CAGAATCGAATTCACATTGACCCAGTTAAACTATCTTCAGGGTACAAAGATGTGATTCTATGGTTTAGTTTTGGTGCATCGTGCTTTATCCT CCTGGCTATCGTTATGGGCCTCGTCTTTATCTGTGTCAAGAACGGGAATATGAGATGTACTATTTGCATC
2849	ATGAACACCCAGATTCTAGTATTCGCCCTGATTGCAATAATCCCAACAAATGCAGATAAAATCTGTCTGGGACACCACGCGGTCTCAAAC GGGACTAAGGTTAATACCCTCACGGAAAGAGGAGTGGAGGTGGTGAACGCTACGGAGACCGTCGAGCGTACAAACATACCCCGGATCT GCTCAAAAGGCAAGAGGACCGTGGATTTAGGACAGTGCGGATTGTTGGGAACAATCACCGGCCCTCCTCAGTGTGACCAATTTCTCGAG TTTAGTGCCGATCTGATTATCGAACGGCGCGAGGGATCAGACGTGTGCTACCCGGGGAAATTCGTAAACGACGAGGCTCTGCGCCAGAT TCTCCGGGAGTCCGGGGGGATTGATAAGGAGGCCATGGGGTTCACATATTCTGGCATTCGGACCAATGGCGCCACTTCTGCCTGCAGA CGCTCCGGCTCGAGCTTTTATGCCGAAATGAAATGGCTCCTTAGCAACACCGATAATGCCGCATTCCCCCAAATGACCAAAAGTTACAAG AATACACGTAAGTCGCCTGCACTTATCGTGTGGGGCATACACCACTCAGTGAGCACAGCTGAGCAAACCAAGTTATATGGTTCGGGGAG TAAGCTGGTGACCGTAGGGTCTTCAAATTATCAGCAGAGTTTCGTCCCTAGTCCAGGCGCCAGGCCTCAGGTTAACGGTCTGTCTGGAC GCATCGATTTTCATTGGTTGATGCTTAATCCGAACGATACAGTGACTTTCTCTTTTAATGGCGCCTTCATCGCCCCCGATCGCGCTAGCTT CCTCCGGGGTAAAAGCATGGGTATCCAGAGCGGCGTGCAGGTTGACGCCAACTGCGAGGGAGACTGCTATCACAGCGGCGGTACCATC ATCTCCAACCTTCCCTTTCAGAACATAGACTCTCGGGCTGTCGGAAAATGTCCTCGATACGTGAAGCAGAGATCACTGCTGCTTGCTACT GGAATGAAGAACGTCCCCGAGATACCGAAAGGGAGAGGTCTCTTTGGTGCTATAGCAGGGTTTATTGAAAACGGGTGGGAAGGCTTGAT CGACGGTTGGTATGGCTTCAGACATCAGAATGCTCAGGGCGAGGGCACAGCAGCAGATTATAAGTCGACACAGAGTGCAATCGATCAGA TTACTGGCAAACTGAATAGGCTTATTGAGAAGACCAATCAGCAATTTGAGCTTATCGACAACGAGTTTAACGAAGTTGAAAAGCAAATTGG GAACGTCATCAACTGGACCCGGGACTCAATCACCGAAGTATGGAGCTATAATGCCGAGTTGCTGGTTGCAATGGAGAACCAGCACACTA TCGATCTTGCGGATTCCGAAATGGATAAACTTTACGAAAGAGTGAAGAGACAGCTCAGAGAAAACGCCGAGGAGGACGGTACGGGTTGT TTTGAGATATTCCACAAGTGCGATGATGATTGCATGGCTAGCATCAGCAACAACACCTATGATCACAGCAAATACCGTGAGGACGCAATG CAGAACAGGATTCAGATTGACCCTGTGAAACTGTCTAGCGGCTACAAGGACGTGATCTTGTGGTTTTCTTTCGGTGCCAGTTGTTTTATCT TACTCGCCATTGTCATGGGGCTCGTTTTCATTTGTGTAAAAAATGGAAACATGAGGTGTACGATTTGTATA
2850	ATGAACACACAAATTCTCGTGTTTGCCCTAATCGCCATTATCCCGACAAACGCAGATAAGATCTGTCTGGGGCACCACGCAGTCAGCAAT GGCACCAAAGTGAATACTCTTACCGAGAGAGGAGTGGAAGTCGTGAATGCCACGGAGACCGTCGAGAGGACTAATATCCCAAGGATTTG TAGCAAAGGTAAGAAAACGGTTGACCTGGGCCAATGCGGCCTGCTGGGCACAATCACAGGTCCCCCCCAGTGCGATCAGTTCCTGGAG TTCTCAGCGGATCTGATAATTGAGCCGCCGAGAGGGGTCAGATGTATGTTACCCAGGTAAATTCGTCAACGAGGAGGCACTGCGGCAGAT CCTGCGAGAAAGTGGAGGGATAGAAAAGGAAGCAATGGGTTTTACCTACTCCGGAATCCGAGCCAATGGGGCCACTTCCGCGTGTAGG AGAAGCGGAAGTTCTTTCTACGCTGAAATGAAGTGGCTCCTGAGTAACACAGATAACGCGGCTTTCCCTCAAATGACCAAGTCATACAAG AATACGCGCAAGTCACCAGCCCTCATTGTGTGGGGAATCCATCATTCCGTCTCGACTGCGGAACAGACAAAACTCTACGGATCCGGAAA CAAACTGGTGACCGTAGGTTCCAGTAATTACCAGCAAAGTTTTGTGCCCTCACCCGGAGCCAGGCCTCAGGTCAATGGGTTAAGTGGGC GCATTGACTTCCACTGGTTGATGCTGAATCCCAACGACACAGTGACTTTCTCTTTCAACGGGGCTTTCATAGCTCCCGATCGGGCCTCTTT CCTGCGGGGTAAGTCTATGGGTATTCAAAGCGGCGTCCAGGTGGATGCTAATTGTGAAGGGGACTGTTACCACTCGGGAGGCACAATCA TTAGTAACCTGCCTTTCCAGAATATTGACTCGAGGGCAGTAGGCAAGTGTCCCCGCTATGTGAAGCAAAGGTCACTCCTTTTGGCAACCG GGATGAAGAATGTGCCCGAGATACCAAAGGGTAGAGGGCTATTTGGCGCAATTGCAGGATTTATCGAGAATGGTTGGGAGGGCCTGATT GACGGCTGGTACGGGTTTCGGCATCAGAATGCCCAGGGGGAAGGGACTGCCGCAGACTACAAAAGCACACAGTCGGCTATTGATCAGA TAACGGGAAAACTGAACAGGTTGATTGAGAAGACCAACCAACAATTCGAACTTATAGACAACGAGTTCAACGAGGTCGAGAAGCAAATCG GCAACGTGATCAATTGGACCCGGGACTCGATCACCGAGGTGTGGTCATATAACGCTGAGCTGTTAGTAGCTATGGAGAATCAGCACACG ATCCACCTTGCAGATAGCGAGATGGATAAACTGTATGAACGGGTCAAGCGGCAGCTGACAGAAAATGCCGAAGAGGACGGTACGGGCT GCTTTGAAATTTTTCATAAGTGCGACGATGACTGCATGGCCAGTATTAGAAACAACACCTACGACCATTCCAAATATCGGGAAGAAGCGAT GCAGAACAGAATTCAGATAGATCCAGTTAAGCTGTCCTCTGGATATAAGGATGTGATACTGTGGTTTTCTTTTGGCGCCAGTTGCTTTATA TTGCTCGCAATTGTGATGGGGCTGGTGTTCATTTGCGTCAAAAACGGGAACATGAGGTGCACCATTTGCATT

2851	ATGAATACACAGATTCTGGTATTTGCTCTGATCGCGATTATTCCGACGAATGCAGATAAGATCTGTTTAGGCCACCACGCTGTTAGCAATG GGACGAAAGTGAATACTCTAACAGAACGTGGGGTGGAAGTCGTCAATGCTACGGAGACTGTCGAGCGCACCAATATTCCAAGGATCTGT TCCAAGGGTAAGAAGACTGTGGACCTCGGGCAATGCGGTCTCCTAGGCACCATTACCGGTCCTCCTCAGTGCGATCAGTTTCTGGAATT CTCTGCAGATCTTATTATAGAGCGGCGCGAGGGCAGTGATGTGTGCTACCCAGGTAAGTTCGTCAATGAAGAGGCTCTGCGTCAAATTTT GAGAGAGTCAGGCGGCATTGATAAGGAGGCCATGGGCTTTACCTACAGCCGAATCCGCACCAACCGAGCAACCAGCGCCTGTCGCAGA AGCGGTAGCAGTTTCTACGCCGAAATGAAATGGCTACTTTCTAATACCGATAATGCTGCGTTTCCACAGATGACAAAGAGTTACAAGAACA CCCGTAAGAGTCCTGCCCTCATCGTGTGGGGGATCCATCACAGTGTGTCAACCGCAGAGCAAACCAAATTGTATGGTTCCGGAAACAAA CTGGTTACCGTGGGCTCTAGCAATTACCAGCAGAGTTTCGTTCCGTCCCCCGGCGCTAGACCCCAAGTCAATGGTCAGAGCGGAAGAAT CGATTTCCACTGGCTGATGCTGAATCCCAATGACACTGTGACTTTCTCCTTTAATGGGGCCTTCATTGCTCCCGACCGCGCTTCATTCTTG CGCGGCAAATCCATGGGAATACAGAGCGGTGTGCAGGTGGATGCCAATTGCGAAGGTGATTGTTATCACAGTGGAGGAACTATTATCTC TAATCTGCCGTTTCAGAACATTGACAGCAGAGCCGTTGGCAAGTGTCCACGATACGTCAAACAGCGGTCCCTCCTGTTGGCCACAGGCA TGAAGAACGTGCCTGAGATCCCTAAGGGGCGCGGACTGTTTGGAGCCATCGCGGGATTTATTGAGAACGGATGGGAAGGCTTGATAGAT GGCTGGTATGGCTTTAGGCACCAGAATGCACAGGGGGAGGGCACCGCCGCTGATTACAAGAGCACGCAGTCTGCGATCGACCAAATAA CCGGTAAGTTAAATCGCCTGATCGAGAAAACCAATCAGCAGTTCGAGCTGATCGACAACGAGTTTAATGAAGTCGAAAAACAGATCGGCA ACGTGATCAATTGGACCAGGGACAGTATCACAGAGGTATGGAGCTACAATGCCGAGCTGCTGGTGGCCATGGAGAATCAGCATACAATT GACCTAGCTGATAGTGAGATGGACAAACTGTATGAGAGGGTCAAGAGACAATTGCGGGAAAACGCGGAGGAAGATGGGACAGGCTGCT TCGAGATCTTCCATAAGTGCGACGACGACTGTATGGCCAGTATCAGAAATAACACGTACGACCACTCGAAGTATAGGGAGGAGGCAATG CAGAACAGAATTCAGATCGACCCTGTGAAGCTTAGTAGTGGATACAAGGACGTGATTCTCTGCTTCTCTTTCGGGGCCTCATGTTTCATTT TGCTCGCTATAGTTATGGGCCTAGTGTTCATCTGCGTGAAGAATGGGAATATGCGGTGCACGATTTGCATC
2852	ATGAATACTCAGATCTTAGTGTTCGCATTGATCGCCATCATCCCAACCAACGCCGATAAAATTTGCCTTGGACATCATGCTGTGAGCAACG GCACAAAGGTCAATACACTTACAGAAAGAGGGGTGGAAGTGGTGAACGCCACAGAAACCGTGGAACGGACGAACATTCCCCGAATTTGT TCTAAGGGAAAGAGAACAGTAGACCTCGGACAGTGTGGCCTCCTGGGCACCATAACCGGCCCTCCACAGTGTGACCAGTTTCTGGAATT CAGCGCCGGATCTGATTATTGAGAGGCGGGAGGGCTCCGATGTCTGCTACCCTGGCAAGTTTGTGAACGAGGAGGCCCTGAGCCAGATC CTACGGGAGTCCGGAGGGATCGATAAAGAGGCAATGGGTTTTACCTACAGCGGCATCCGGACCAATGGAGCCACTTCTGCATGCCGGC GCTCCGGGAGTTCATTTTATGCTGAGATGAAATGGTTACTTTCTAACACCGATAACGCAGCCTTCCCACAGATCACCAAGAGCTATAAAAA TACGCGAAAATCTCCAGCCCTCATAGTCTGGGGTATCCATCACTCTGTTAGCACAGCGGAGCAGACCAAGTTATACGGGTCAGGAAATAA ACTTGTGACAGTGGGCTCCTCTAACTATCAGCAGTCTTTTGTGCCCAGCCCTGGGGCCAGGCCCCAAGTCAATGGCCTTTCTGGTAGGA TTGATTTCCACTGGTTGATGTTGAACCCCAATGATACCGTCACATTCAGCTTTAATGGTGCTTTTATCGCCCCTGACCGAGCTAGTTTCCT AAGAGGAAAATCTATGGGTATCCAGTCTGGCGTTCAGGTGGACGCCAATTGTGAGGGCGACTGTTATCATTCCGGGGGAACCATCATCT CTAACCTTCCCTTCCAAAATATCGACTCCAGGGCAGTTGGCAAATGTCCACGGTATGTGAAGCAGAGATCACTCCTGTTAGCGACTGGGA TGAAAAACGTGCCAGAAATCCCTAAAGGTCGTGGATTGTTCGGAGCGATCGCGGGATTTATCGAAAACGGGTGGGAGGGTCTTATAGAC GGCTGGTATGGGTTTAGACACCAGAACGCACAGGGTGAGGGCACCGCGGCAGACTACAAATCTACACAATCTGCAATAGATCAGATCAC CGGCAAGCTGAACCGACTTATTGAAAAGACCAATCAGCAGTTTGAGTTAATCGATAACGAATTCAACGAGGTGGAGAAGCAAATAGGGAA TGTTATTAACTGGACACGGGACTCAATCACAGAAGTATGGTCGTATAATGCAGAGCTGCTAGTCGCAATGGAGAATCAGCATACGATTGA CCTCGCCGACAGTGAGATGGACAAGTTGTATGAAAGGGTCAAAAGACAGCTCCGGGAAAATGCCGAGGAAGACGGCACAGGCTGTTTT GAAATATTCCATAAGTGCGACGATGACTGCATGGCAAGTATTACGAACAACACGTATGACCATTCAAAGTACAGGGAGGAGGCCATGCAG AACCGCATCCAGATTGATCCAGTTAAGCTTAGCAGTGGCTATAAGGATGTCATCCTGTGGTTTAGTTTTGGCGCCTCCTGTTTTATCCTGC TGGCCATCGTTATGGGGCTTGTGTTCATATGCGTGAAGAATGGCAATATGCGCTGCACCATTTGTATT
2853	ATGAATACACAAATTCTGGTATTCGCCCTCATAGCTATAATTCCCACCAACGCCGATAAGATTTGTTTAGGTCACCACGCCGTGTCAAACG GGACAAAGGTTAATACACTTACAGAGCGGGGCGTGGAAGTCGTGAACGCTACTGAGACAGTCGAAAGGACTAACATCCCCAGAATTTGT AGCAAAGGAAAAAAGACCGTGGATTTGGGGCAGTGTGGCTTACTAGGGACAATCACCGGTCCACCCCAGTGTGACCAGTTCCTCGAGTT CTCAGCGGACCTAATTATCGAGCGTCGAGAGGGCAGCGATGTGTGCTATCCCGGGAAATTCGTGAATGAGGAGGCCTTACGGCAGATCT TGCGTGAAAGTGGGGGCATAGATAAAGAGGCGATGGGGTTCACGTATTCTGGCATTCGCACCAACGGAGCTACCTCAGCCTGTCGTCGT TCAGGATCCAGCTTCTACGCAGAAATGAAGTGGTTACTCTCTAACACAGATAACGCAGCCTTCCCGCAAATGACGAAGAGCTACAAAAAC ACTCGCAAATCCCCCGCCCTCATCGTCTGGGGTATTCACCACAGCGTTAGCACCGCTGAACAGACTAAATTATACGGCAGCGGAAATAAG CTTGTCACCGTCGGGAGCTCCAACTATCAGCAGAGCTTTGTCCCATCTCCTGGGGCCCGGCCCCAGGTGAACGGACAGTCCGGGCGCA TTGATTTCCACTGGTTGATGCTGAACCCAAACGACACTGTTACGTTCAGCTTTAACGGCGCCTTCATTGCACCTGACCGAGCTAGCTTTTT GCGGGGCAAATCGATGGGAATACAGTCAGGGGTCCAGGTAGACGCCAACTGCGAAGGCGATTGCTACCATTCCGGCGGGACAATCATC TCCAACCTGCCCTTCCAGAACATTGACAGCCGGGCAGTCGGGAAGTGTCCCAGATATGTGAAACAACGCTCTCTGCTGCTTGCGACCGG CATGAAAAACGTTCCAGAAATCCCGAAGGGCGAGGTTTGTTTGGCGCTATAGCTGGGTTCATTGAGAACGGCTGGGAAGGCCTCATCG ACGGCTGGTATGGATTTAGGCATCAAAATGCACAGGGCGAAGGTACCGCTGCCGACTATAAGTCAACTCAGAGCGCAATAGACCAAATT ACCGGAAAGCTCAATCGCCTGATTGAGAAAACCAATCAGCAGTTTGAACTGATAGACAACGAATTTAATGAAGTGGAGAAGCAGATCGGG AATGTGATCAACTGGACTCGTGACAGTATTACTGAGGTATGGAGTTATAACGCTGAACTTCTGGTGGCTATGGAGAACCAACACACTATT GATCTCGCCGACTCCGAGATGGATAAACTCTACGAAAGAGTTAAAAGACAGCTTAGGGAAAACGCCGAGCAGGATGGTACTGGATGTTT CGAGATCTTTCATAAATGCGATGACGACTGCATGGCTAGCATCCGGAATAACACCTACGATCATTCTAAGTATCGTGAAGAAGCCATGCA AAATCGTATTCAGATTGATCCGGTAAAATTGAGCTCAGGCTATAAGGACGTCATACTGTGGTTCTCCTTCGGCGCTTCCTGCTTTATCTTA TTAGCAATTGTGATGGGATTAGTTTTCATCTGCGTGAAAAACGGAAACATGCGCTGTACCATTTGCATA

2854	ATGAACACTCAAATTTTAGTGTTCGCCTTGATAGCAATCATCCCCACAAATGCGGATAAAATCTGTCTCGGCCATCATGCCGTGTCTAATG GTACCAAAGTAAATACCCTTACCGAGCGTGGGGTTGAGGTCGTTAATGCCACCGAGACTGTGGAGAGAACCAATATCCCCCGCATCTGT TCCAAGGGAAAAAAGACCGTTGACCTCGGTCAGTGTGGTCTGCTTGGTACAATAACCGGACCGCCCCAGTGCGACCAATTCCTGGAATT CAGTGCGGATCTCATAATCGAAAGGCGAGAAGGCTCCGATGTTTGTTACCCCGGAAAGTTCGTCAACGAGGAGGCCCTGCGACAGATAT TGCGGGAAAGCGGAGGCATCGATAAGGAGGCTATGGGCTTCACATATACTGGAATCCGCACGAATGGGGCCACCTCAGCCTGTCGGCG CAGCGGGTCTAGTTTTTATGCTGAAATGAAGTGGCTTCTTTCGAATACTGACAATGCTGCCTTCCCCCAGATGACCAAGAGTTATAAAAAT ACCAGAAAGAGCCCTGCCCTGATAGTCTGGGGAATCCATCACAGCGTAAGCACTGCCGAACAGACTAAGCTCTATGGCAGCGGGAATAA GCTCGTGACCGTGGGCAGTTCCAATTACCAGCAGAGCTTTGTGCCATCCCCGGGGGCCCGTCCACAGGTTAACGGGCAATCTGGCCGA ATTGATTTTCATTGGCTGATGCTGAATCCAAACGATACTGTGACTTTCTCTTTTAACGGAGCCTTCATTGCACCCGATCGGGCAAGCTTCC TGCGGGGAAAATCTATGGGAATTCAGAGTGGGGTGCAGGTGGACGCAAACTGTGAAGGCGATTGCTACCACTCTGGCGGGACAATCATT AGCAATCTGCCTTTCCAGAACATAGACAGCCGCGCAGTTGGAAAATGCCCGCGTTATGTCAAGCAAAGGAGTTTACTGCTCGCTACAGG CATGAAGAATGTTCCGGAGATCCCGAAGGGAAGAGGTTTGTTCGGCGCCATAGCAGGATTCATTGAAAATGGCTGGGAAGGGTTAATCG ACGGGTGGTATGGTTTTCGACACCAGAACGCACAAGGGGAGGGGACAGCAGCAGATTACAAGAGCACACAGTCAGCAATTGACCAGATT ACTGGAAAGCTAAACCGGCTCATCGAGAAGACCAACCAGCAGTTCGAATTGATTGACAACGAATTTAATGAGGTGGAAAAACAGATCGGT AACGTCATAAACTGGACTAGAGATTCCATTACTGAGGTCTGGTCATATAATGCCGAACTGCTCGTGGCGATGGAGAATCAGCATACCATA GATCTCGCCGACTCCGAGATGGACAAACTTTATGAACGGGTGAAGCGCCAGCTGCGGGAAAATGCCGAAGAGGACGGTACAGGGTGTT TCGAGATTTTCCATAAATGTGATGACGACTGCATGGCAAGCATACGCAATAACACATATGATCACTCTAAGTATAGGGAAGAGGCCATGC AGAACCGAATTCAAATCGACCCAGTAAAGCTTTCTAGCGGCTATAAAGATGTGATTCTGTGGTTTAGCTTCGGGGCCTCTTGTTTTATCCT GTTAGCCATCGTCATGGGGCTAGTTTTCATCTGCGTGAAAAATGGCAACATGCGTTGCACTATATGCATC
2855	ATGAATACGCAGATCCTCGTGTTCGCACTGATAGCGATCATCCCGACAAATGCTGACAAAATCTGTTTGGCCCATCATGCCGTTAGCAAT GGTACCAAAGTGAACACTCTCACTGAGCGCGGCGTGGAAGTGGTGAACGCCACTGAAACAGTAGAAAGAACAAACATTCCCAGAATCTG CAGTAAGGGTAAGAAAACTGTCGACTTGGGCCAGTGTGGCCTATTAGGAACAATCACCGGACCTCCCCAATGCGATCAGTTCCTAGAATT TAGCGCTGATCTGATAATTGAAAGAAGAAGGTTCGGATGTCTGTTACCCGGGTAAATTCGTGAACGAAGAAGCCTTAAGACAGATTCT CAGAGAAAGTGGCGGAATTGATAAAGAAGCCATGGGGTTCACTTATTCTGGCATACGGACAAATGGGGCCACTAGCGCATGTAGACGAT CAGGCAGCTCTTTCTACGCCGAGATGAAGTGGCTGTTATCCAACACAGACAATGCAGCCTTTCCTCAGATGACCAAATCCTACAAAAATA CAAGGAAGAGCCCCGCACTAATCGTATGGGGCATCCATCACTCTGTGTCGACGGCAGAACAAACCAAGCTCTATGGTAGTGGGAACAAG TTGGTGACGGTGGGGTCCTCTAATTACCAGCAGTCGTTCGTGCCCTCTCCAGGTGCCCGGCCCCAGGTGAATGGTCAGTCCGGCCGAA TCGACTTTCACTGGCTTATGCTGAATCCCAATGATACCGTAACATTTAGCTTTAATGGAGCTTTCATTGCTCCTGACCGTGCCAGCTTCTT GAGAGGAAAATCTATGGGCATACAGTCTGGTGTCCAGGTGGATGCTAACTGCGAAGGTGACTGTTACCACTCAGGAGGTACTATAATTAG CAATCTCCCCTTCCAGAACATCGACAGCCGCGCTGTAGGAAAGTGCCCTCGCTACGTGAAGCAACGATCATTACTGCTGGCCACTGGCA TGAAGAACGTACCCGAGATCCCAAAGGGAAGGGGTTTGTTTGGCGCCATTGCGGGCTTTATTGAGAACGGATGGGAGGGTCTGATCGAT GGATGGTATGGATTCCGTCATCAGAATGCCCAGGGCGAGGGGACCGCTGCTGATTACAAATCCACACAATCTGCAATAGATCAAATCACA GGAAAACTGAACCGGCTTATTGAAAAAACCAATCAGCAGTTCGAGTTGATCGACAATGAGTTCAATGAAGTGGAAAAGCAGATCGGCAAC GTCATCAACTGGACTCGGGACTCTATCACAGAGGTCTGGTCCTACAATGCAGAGTTACTAGTTGCCATGGAGAACCAGCACACGATTGAT TTGGCCGATAGCGAGATGGATAAACTCTACGAGCGTGTGAAGAGGCAGCTGCGCGAAAATGCCGAAGAGGATGGAACAGGTTGCTTCG AGATCTTCCACAAATGCGATGACGACTGCATGGCCTCCATCAGGAATAATACTTATGATCACAGCAAGTACCGGGAGGAAGCCATGCAAA ACAGAATTCAGATTGATCCCGTAAAGTTAAGCTCCGGCTACAAAGATGTGATTCTGTGGTTCTCTTTCGGAGCTTCATGCTTCATCCTCCT CGCTATAGTGATGGGTCTCGTATTCATTTGCGTTAAGAACGGCAATATGAGATGCACAATCTGTATT
2856	ATGAATACTCAAATCCTGGTGTTCGCACTCATTGCCATCATACCAACCAACGCAGACAAGATATGCTTGGGCCATCACGCCGTGTCTAAT GGTACGAAAGTGAATACTCTGACTGAGCGCGGGGTCGAAGTAGTAAACGCGACTGAAACAGTGGAACGAACTAATATTCCCAGAATTTG CAGTAAAGGGAAAAAGACCGTCGATCTCGGCCAATGCGGACTGCTGGGGACAATCACAGGGCCACCCAATGCGATCAGTTCCTGGAG TTTAGCGCGGATCTCATCATCGAGCGAAGAGAAGGGTCCGACGTCTGTTACCCGGGTAAATTTGTGAATGAAGAGGCCCTGCGTCAGAT TCTGCGGGAAAGCGGCGGAATTGACAAGGAGGCAATGGGGTTCACGTACAGTGGTATCAGAACCAACGGAGCTACAAGCGCATGCAGA CGGTCCGGGTCCTCGTTTTACGCGGAGATGAAGTGGCTGTTATCCAACACTGACAACGCTGCTTTCCCACAGATGACCAAAAGCTATAAG AACACAAGAAAAAGTCCAGCCCTGATTGTGTGGGGGATCCACCATTCGGTATCAACTGCTGAGCAAACAAAGTTATATGGAAGTGGAAAC AAGCTTGTCACCGTAGGTTCATCCAACTATCAACAGAGTTTCGTGCCCTCCCCCGGAGCCCGCCCCCAGGTGAATGGACAGTCGGGAAG AATAGACTTTCATTGGTTGATGCTGAATCCCAATGACACCGTGACCTTCAGCTTTAATGGGGCCTTCATTGCCCCGGACAGGGCTAGCTT TCTCCGAGGAAAGAGCATGGGCATTCAATCTGGAGTTCAAGTGGATGCCAACTGTGAGGGCGATTGCTACCATAGCGGAGGGACAATCA TCAGCAACCTCCCCTTCCAAAACATTGATAGTCGTGCTGTAGGAAAATGCCCTCGGTACGTTAAACAAAGATCGCTGCTACTGGCTACAG GAATGAAGAACGTGCCCGAGATCCCAAAAGGACGAGGCCTATTTGGGGCCATCGCGGGTTTTATCGAAAATGGATGGGAGGGCCTCATC GACGGGTGGTATGGCTTTCGTCATCAGAATGCTCAAGGCGAAGGAACAGCCGCTGATTACAAATCAACCCAATCTGCCATCGACCAGAT CACCGGAAAGCTCAACCGCCTGATTGAAAAGACTAATCAGCAGTTCGAGCTGATCGACAATGAGTTCAATGAAGTGGAAAAGCAGATTGG TAACGTGATTAACTGGACTAGAGATTCCATTACCGAAGTGTGGTCTTATAACGCCGAGCTACTGGTCGCCATGGAGAACCAGCATACCAT TGATTTAGCTGATTCTGAGATGGACAAGCTGTACGAAAGGGTGAAACGGCAGCTCCGAGAAAATGCCGAAGAAGACGGAACCGGGTGTT TCGAGATCTTCCACAAGTGCGACGACGATTGCATGGCTAGCATTAGAAACAACACTTATGACCACAGCAAGTATCGCGAAGAGGCGATG CAGAATCGAATCCAGATCGATCCTGTCAAGCTGAGCAGCGGGTACAAGGACGTGATCTTATGGTTTTCCTTCGGGGCTAGCTGTTTTATC CTGCTGGCTATCGTGATGGGTCTGGTTTTTATATGTGTTAAAAACGGCAACATGCGGTGCACAATCTGCATT

2857	ATGAATACCCAGATTTTAGTGTTTGCACTTATCGCTATCATACCAACTAATGCTGACAAGATCTGCCTGGGCCACCATGCAGTGAGCAACG GGACCAAAGTGAACACCCTAACCGAGCGGGGCGTGGAGGTGGTAAATGCCACTGAAACCGTCGAAAGGACGAACATTCCGCGGATTTG CAGTAAAGGCAAACGAACCGTGGACCTGGGCCAGTGTGGCCTCCTGGGGACCATCACGGGACCACCCCAATGTGACCAGTTCCTGGAG TTCTCCGCTGACCTTATCATCGAAAGGCGAGAAGGGTCTGATGTCTGTTACCCTGGAAAATTCGTAAACGAGGAGGCTCTCCGACAAATT CTGCGGGAATCCGGCGGCATCGACAAGGAGGCGATGGGTTTTACCTATTCCGGGATACGCACTAACGGAGCTACATCAGCATGTAGGC GCTCTGGGTCTAGTTTCTATGCCGAAATGAAATGGTTGCTGTCAAACACTGACAACGCTGCTTTCCCTCAGATGACTAAGTCCTATAAGAA TACCCGGAAGTCCCCAGCACTAATAGTCTGGGGGATTCACCACTCCGTATCTACTGCGGAGCAGACGAAACTTTACGGGAGCGGGAACA AACTGGTCACTGTTGGGTCATCAAATTATCAGCAATCATTCGTACCCTCTCCCGGCGCTCGGCCCCAGGTCAATGGCCTGTCCGGCAGA ATTGACTTCCACTGGCTTATGCTAAATCCTAACGACACCGTGACCTTCTCCTTCAACGGCGCCTTCATCGCGCCTGACCGAGCTAGCTTT CTGCGTGGGAAAAGTATGGGAATTCAGAGCGGTGTGCAGGTGGATGCAAATTGTGAAGGCGACTGCTATCATAGCGGGGGTACCATTAT CTCTAATCTGCCTTTCCAGAACATCGACAGTAGAGCCGTGGGCAAATGCCCCAGATATGTGAAACAACGTAGCCTTCTGCTGGCAACAGG AATGAAAAATGTGCCCGAGATTCCCAAAGGGCGTGGGCTGTTTGGGGCCATCGCCGGCTTCATCGAAAACGGTTGGGAGGGATTAATCG ACGGGTGGTATGGCTTCAGGCACCAAAACGCTCAAGGGGAGGGCACGGCGGCGGATTACAAATCTACCCAGTCCGCCATTGATCAAATT ACTGGCAAATTAAACAGAATCATCGAAAAGACTAACCAGCAGTTCGAACTGATTGATAATGAGTTCAACGAGGTTGAGAAGCAAATCGGTA ACGTTATCAACTGGACGCGTGACAGTATCACCGAGGTCTGGTCATATAATGCTGAGCTGCTCGTGGCTATGGAAAACCAGCACACTATCG ATCTGGCGGACTCCGAGATGGACAAGTTGTATGAACGGGTGAAAAGACAGCTTCGGGAGAACGCCGAGGAGGACGGGACCGGATGCTT CGAGATCTTTCACAAGTGCGATGATGATTGCATGGCTTCAATTCGCAATAATACATATGACCATTCTAAGTACAGGGAGGAGGCTATGCA GAACCGGATCCAGATAGATCCCGTCAAACTTAGTAGCGGGTACAAGGATGTCATCCTGTGGTTTTCCTTCGGGGCCTCATGCTTTATTCT TCTGGCCATCGTGATGGGACTCGTGTTCATCTGCGTTAAGAATGGCAACATGCGATGTACAATCTGCATC
2858	ATGAACACCCAGATTCTCGTGTTCGCTTTGATCGCGATTATACCTACTAACGCAGATAAAATATGTCTGGGTCATCACGCTGTCAGTAACG GGACAAAGGTGAACACGTTGACCGAGAGGGGCGTCGAGGTGGTTAACGCCACTGAAACAGTGGAACGGACCAATATACCAAGGATCTG TAGCAAGGGTAAGAAGACTGTCGACCTCGGTCAGTGTGGGTTGCTTGGGACCATCACCGGGCCTCCACAGTGTGATCAGTTTCTGGAGT TCTCGGCTGACCTGATAATCGAGAGACGGGAGGGCAGTGACGTGTGCTACCCAGGGAAGTTTGTGAACGAAGAAGCATTGCGACAGATT CTTCGGGAATCAGGTGGAATCGACAAGGAGGCCATGGGATTTACATACTCTGGGATCAGGACCAACGGTGCTACCTCCGCTTGTAGGAG AAGTGGATCAAGCTTTTATGCTGAAATGAAATGGCTTCTGTCCAATACGGACAACGCAGCTTTCCCTCAGATGACAAAGTCCTACAAAAAC ACAAGAAAGTCACCCGCTCTGATTGTTTGGGGAATCCACCACTCAGTGTCCACGGCGGAGCAGACGAAACTCTATGGCTCTGGGAATAA GCTGGTGACAGTCGGGAGCTCAAATTATCAGCAATCCTTCGTTCCATCCCCCGGCGCCCGACCTCAGGTCAACGGCCAATCGGGCAGG ATTGACTTCCATTGGCTAATGCTCAATCCCAATGACACCGTCACATTCTCTTTTAATGGTGCCTTCATCGCCCCAGATCGAGCCTCTTTTCT CAGAGGTAAGTCGATGGGAATACAGTCAGGGGTTCAGGTGGACGCTAATTGCGAAGGGGACTGCTATCATTCTGGGGGCACAATTATTT CAAATCTGCCATTCCAGAACATTGACTCTCGAGCCGTCGGAAAATGCCCTCGATATGTAAAGCAGCGAAGTCTGCTGCTGGCAACCGGC ATGAAAAACGTACCCGAGATCCCAAAGGGCAGAGGGCTTTTCGGTGCGATCGCCGGGTTTATCGAAAATGGCTGGGAGGGCCTGATCG ATGGGTGGTACGGTTTTCGGCACCAGAACGCTCAGGGTGAGGGCACTGCAGCGGATTATAAGAGTACCCAGTCAGCCATTGACCAGATC ACAGGTAAGTTGAACCGCCTCATTGAAAAGACCAACCAGCAGTTTGAGCTGATCGATAATGAGTTTAATGAGGTGGAGAAGCAGATCGGG AATGTTATTAATTGGACGCGCGATTCAATCACAGAGGTATGGTCATACAACGCTGAGCTGCTGGTGGCAATGGAGAATCAACACACGATC GACCTCGCTGACTCAGAAATGGATAAACTGTACGAAAGGGTCAAAAGGCAGCTGCGCGAGAACGCCGAAGAAGACGGAACTGGTTGTTT CGAAATTTTCCACAAGTGTGACGACCACTGTATGGCATCCATCAGAAACAACACATACGACCATAGCAAATACCGGGAGGAGGCAATCCA GAATCGAATCCAGATTGATCCCGTGAAGTTGTCTAGCGGATACAAAGATGTGATCTTATGGTTTTCATTCGGGGCTAGCTGTTTCATCCTC CTGGCCATAGTTATGGGGCTCGTTTTCATCTGCGTAAAGAATGGGAATATGAGATGTACTATCTGCATT
2859	ATGAATACGCAGATTCTGGTGTTCGCTCTCATCGCTATTATTCCTACCAATGCCGATAAGATCTGTCTGGGGCACCACGCCGTCAGCAAC GGCACTAAAGTGAACACCTTGACAGAAAGGGGGGTGGAAGTGGTGAACGCAACGGAAACAGTGGAGAGGACAAACATACCGCGGATTT GCTCGAAAGGAAAACGGACTGTAGACCTGGGCCAATGCGGCCTGCTCGGAACCATCACCGGGCCACCCCAATGTGACCAGTTCCTAGA ATTTTCCGCCGACCTTATAATTGAGCGGAGGGAGGGATCCGATGTGTGTTATCCTGGCAAGTTCGTGAACGAGGAGGCCCTGCGGCAGA TATTGCGTGAATCTGGGGGCATCGACAAGGAGGCGATGGGATTCACATACTCCGGCATCCGCACCAACGGCGCGACCAGCGCATGCCG TCGCTCCGGGTCCTCATTCTACGCTGAAATGAAATGGCTCCTCTCAAATACTGATAATGCTGCTTTTCCTCAAATGACCAAAAGTTACAAA AACACCAGGAAAAGCCCAGCTCTGATTGTCTGGGGCATTCACCATTCCGTATCTACGGCAGAACAGACTAAACTATATGGCAGTGGCAGC AAGCTCGTGACCGTCGGCTCCTCCAACTATCAGCAATCTTTCGTTCCATCACCAGGCGCCCGGCCCCAGGTCAATGGGCTCTCCGGCCG TATCGATTTCCACTGGCTGATGCTCAATCCTAACGACACCGTTACCTTTTCTTTTAACGGCGCCTTTATTGCTCCAGACAGAGCGTCTTTC CTGCGGGGTAAAAGCATGGGCATTCAGTCTGGTGTCCAAGTGGACGCAAACTGTGAAGGGGATTGCTATCATAGTGGCGGCACTATCAT TAGTAACCTGCCTTTCCAGAACATTGATTCGCGGGCAGTTGGCAAATGCCCCAGGTATGTCAAACAGAGATCACTCTTGTTAGCTACCGG AATGAAGAACGTACCGGAGATTCCTAAGGGTAGGGGACTGTTCGGGGCTATTGCCGGGTTCATCGAGAATGGGTGGGAAGGTCTCATC GACGGGTGGTATGGCTTTCGTCATCAGAACGCCCAGGGCGAAGGTACAGCAGCCGATTATAAGTCTACCCAGTCCGCTATCGACCAAAT AACCGGGAAGTTGAACCGGTTAATCGAGAAAACCAACCAACAGTTCGAATTGATCGACAATGAATTTAATGAGGTGGAAAAACAAATAGG GAACGTGATAAACTGGACAAGGGACTCCATTACAGAAGTCTGGAGCTATAACGCCGAACTGCTGGTAGCTATGGAGAATCAGCATACCAT TGACCTGGCAGACAGTGAGATGGACAAACTTTACGAACGGGTTAAAAGACAGTTGCGAGAGAATGCCGAAGAAGATGGAACTGGCTGCT TTGAGATTTTTCACAAATGCGATGATGACTGTATGGCCAGCATTAGAAACAACACATACGACCACAGTAAGTACAGGGAAGAAGCCATGC AAAACCGGATCCAGATTGACCCCGTCAAGCTGAGCTCGGGTTATAAGGATGTAATCTTATGGTTCAGTTTTGGCGCCTCCTGCTTCATTCT TCTAGCAATCGTGATGGGATTGGTCTTCATTTGCGTGAAAAATGGCAATATGAGGTGCACCATCTGCATT

2860	ATGAATACACAGATCCTAGTCTTCGCCTTAATTGCAATAATACCTACCAACGCAGACAAAATTTGTCTTGGGCACCACGCCGTGTCCAACG GGACCAAAGTTAATACCCTCACCGAACGTGGGGTGGAGGTGGTAAACGCTACGGAGACAGTAGAAAGGACCAACATACCTCGCATCTGT TCCAAAGGGAAAAGAACCGTCGACCTGGGGCAGTGCGGGCTGCTAGGAACTATTACTGGCCCACCACAGTGCGATCAGTTTCTGGAATT TAGTGCTGATCTGATCATCGAAAGGCGTGAGGGTAGCGATGTCTGTTATCCCGGGAAGTTCGTGAATGAAGAGGCGCTGAGACAAATTC TGAGACAAAGCGGGGGGATCGATAAGGAAGCAATGGGGTTTACTTATTCTGGCATTAGAACCAATGGGGCCACATCCGCCTGCCGCAG GTCTGGAAGCTCCTTTTATGCTGAAATGAAATGGCTGCTGAGTAATACTGACAACGCCGCTTTTCCTCAGATGACAAAATCTTATAAAAAC ACGAGAAAATCTCCTGCCCTGATTGTGTGGGGCATTCACCATTCGGTTTCAACAGCGGAGCAAACTAAACTATATGGAAGCGGCAGCAAG CTGGTCACAGTGGGAAGCTCTAACTACCAACAGTCTTTTGTTCCCAGTCCCGGCGCCCGACCTCAGGTCAACGGACTGAGTGGCAGAAT TGATTTCCACTGGCTGATGCTTAATCCGAACGATACCGTCACCTTTTCTTTCAATGGTGCTTTTATTGCCCCTGACAGGGCAAGCTTCTTG CGCGGGAAGTCTATGGGAATCCAATCCGGCGTACAAGTGGACGCAAACTGTGAAGGGGACTGTTATCACTCCGGAGGCACAATTATTAG TAACTTGCCTTTCCAGAATATCGATTCCAGAGCCGTGGGTAAATGCCCTCGATACGTAAAACAGCGCTCTCTGCTTCTGGCTACCGGAAT GAAGAATGTGCCTGAGATCCCTAAGGGCAGAGGCCTGTTCGGGGCCATCGCTGGTTTTATTGAGAATGGGTGGGAAGGCTTAATAGACG GGTGGTACGGTTTTCGTCACCAAAATGCACAGGGCGAAGGCACAGCAGCCGACTACAAAAGTACGCAAAGCGCCATAGATCAGATAACA GGAAAACTTAATCGCCTGATCGAAAAAACGAACCAACAGTTCGAACTGATTGACAATGAATTTAACGAAGTCGAAAAGCAGATAGGCAAT GTCATCAATTGGACGAGAGACAGTATTACAGAAGTTTGGAGCTATAATGCAGAGCTGTTAGTGGCGATGGAAAATCAGCACACCATTGAT TTGGCGGACAGCGAAATGGACAAGCTGTACGAACGAGTGAAGCGACAGCTACGCGAGAATGCAGAAGAAGATGGGACTGGCTGCTTCG AAATCTTTCACAAGTGCGACGACGATTGCATGGCCAGCATTCGCAATAACACATACGATCACTCCAAGTATCGAGAGGAGGCGATGCAAA ACCGAATCCAAATTGACCCTGTGAAATTATCTAGTGGGTATAAAGACGTGATACTGTGGTTTTCATTCGGAGCTTCATGTTTCATTCTGTTG GCAATTGTCATGGGACTGGTGTTTATATGTGTAAAAAGCAGAAATATGCGATGCACCATTTGCATC
2861	ATGAATACCCAGATCCTGGTATTCGCATTAATTGCTATCATTCCTACAAATGCTGACAAAATCTGTCTCGGCCATCACGCCGTGAGTAATG GAACTAAAGTGAATACTCTCACTGAGCGCGGGGTAGAGGTAGTGAATGCCACTGAGACCGTCGAGCGAACTAACATCCCCCGCATCTGT AGCAAAGGCAAGAAGACGGTGGACTTAGGACAGTGTGGCCTCCTGGGGACCATCACAGGACCTCCTCAGTGCGACCAGTTTCTGGAATT TAGTGCTGACCTTATCATTGAACGGCGCGAGGGCAGCGATGTCTGCTACCCTGGCAAGTTTGTCAATGAGGAAGCCTTACGACAGATCC TCAGAGAATCAGGGGGGATTGACAAGGAGGCAATGGGGTTCACCTATAGCGGAATCCGGACTAATGGCGCAACAAGTGCATGTAGACG GAGTGGGAGTAGTTTTTACGCCGAAATGAAGTGGCTGTTAAGCAACACGGACAACGCTGCTTTTCCACAGATGACTAAGTCTTACAAAAA CACCAGGAAGTCACCTGCTCTGATCGTGTGGGGAATTCACCATAGCGTGAGCACAGCTGAGCAGACCAAGCTTTACGGAAGTGGCAACA AACTCGTGACAGTGGGCAGCTCCAACTACCAGCAGTCTTTCGTGCCCTCCCCCGGCGCAAGACCTCAGGTCAACGGGCAGTCGGGGCG TATTGACTTCCACTGGCTCATGCTGAACCCCAATGATACTGTGACCTTTTCGTTCAATGGGGCGTTTATCGCCCCTGATCGGGCAAGCTT CCTGAGGGGTAAATCAATGGGAATACAGTCCGGCGTCCAGGTGGATGCCAACTGTGAAGGCGACTGCTATCATTCCGGGGGTACCATTA TCAGCAACCTTCCCTTTCAGAACATTGATTCTCGGGCTGTGGGAAAATGTCCCAGGTACGTCAAGCAGCGGTCACTCCTGCTGGCAACTG GGATGAAGAACGTCCCTGAAATCCCGAAAGGGCGTGGGCTGTTTGGTGCTATCGCTGGATTCATCGAAAACGGCTGGGAAGGGCTTATT GATGGTTGGTACGGGTTTCGCCACCAGAACGCGCAGGGCGAGGGCACCGCTGCAGACTATAAATCTACTCAGTCTGCAATTGACCAGAT CACCGGCAAACTGAACCGCCTGATTGAGAAGACCAACCAACAGTTTGAGCTCATAGATAACGAATTTAACGAGGTGGAAAAGCAAATCGG AAACGTTATCAACTGGACTAGGGACTCTATCACTGAAGTGTGGTCCTACAATGCAGAGCTGCTCGTTGCCATGGAAAACCAGCACACCAT TGACCTAGCTGACTCCGAGATGGATAAACTGTATGAGCGGGTAAAACGGCAGCTGAGAGAGAACGCCGAAGAGGACGGGACAGGATGT TTTGAGATATTTCACAAATGCGACGACGACTGTATGGCAAGCATCCGGAACAACACCTACCATCATTCTAAATATCCGGACCAAGCCATG CAGAATCGCATTCAGATTGATCCCGTCAAACTGAGTTCTGGCTATAAGGACGTAATTCTGTGGTTCTCCTTCGGAGCTAGTTGCTTCATTC TGCTTGCAATAGTAATGGGACTCGTTTTTATTTGTGTCAAGAACGGCAACATGAGGTGTACCATTTGCATA
2862	ATGAATACACAGATTCTGGTCTTTGCTCTCATCGCTATAATCCCGACCAACGCTGACAAAATCTGTCTGGGTCACCACGCCGTATCTAATG GAACTAAGGTGAATACATTGACCGAGCGCGGCGTTGAGGTCGTTAATGCTACAGAAACTGTTGAGCGGACAAATATCCCCCGGATATGC TCAAAGGGAAAAAAGACTGTCGACTTAGGACAATGCGGGTTACTAGGCACAATAACAGGGCCGCCCCAGTGCGACCAGTTTCTCGAGTT CAGTGCCGATCTGATTATCGAACGCCGGGAGGGAAGCGACGTCTGTTATCCCGGCAAATTCGTGAATGAAGAGGCCCTCCGCCAGATCC TGAGAGAATCCGGAGGCATCGACAAGGAGGCCATGGGATTCACTTACTCTGGAATCAGGACCAACGGGGCTACCTCCGCGTGCCGACG GTCCGGGTCCTCTTTCTACGCAGAGATGAAGTGGCTACTATCTAATACCGATAACGCAGCTTTCCCCCAAATGACAAAATCATATAAGAAT ACAAGGAAATCTCCGGCATTGATCGTCTGGGGCATTCATCACTCTGTCAGTACCGCTGAACAAACAAAGCTGTATGGCAGCGGGAACAA GCTGGTTACAGTGGGGTCTAGTAACTATCAACAGTCCTTCGTGCCATCTCCGGGAGCGCGGCCACAGGTGAATGGGCAGTCTGGCCGG ATTGACTTTCACTGGTTAATGTTAAATCCAAACGACACGGTGACTTTTTCATTTAACGGTGCCTTTATTGCTCCTGACAGAGCCTCATTTCT CAGAGGCAAATCAATGGGTATCCAATCAGGCGTGCAGGTGGACGCTAACTGCGAGGGCGATTGCTACCACTCTGGGGGCACAATCATAT CCAACCTGCCCTTCCAGAATATTGACTCAAGAGCCGTCGGGAAATGTCCACGGTACGTGAAACAGAGGAGTCTACTGCTAGCTACAGGA ATGAAGAATGTGCCCGAGATTCCGAAAGGGAGAGGGTTGTTTGGAGCCATCGCAGGATTCATTGAAAATGGATGGGAAGGTCTCATAGA TGGTTGGTATGGCTTTCGACACCAAAACGCGCAGGGCGAGGGGACTGCAGCTGACTACAAAAGTACGCAATCTGCCATTGATCAGATTA CAGGAAAGCTCAACCGACTGATCGAAAAAACAAATCAGCAGTTTGAGCTCATAGACAATGAGTTCAATGAAGTGGAAAAGCAGATCGGCA ACGTTATCAACTGGACGCGCGACTCAATCACAGAAGTGTGGTCTTACAATGCAGAATTGCTTGTCGCCATGGAGAATCAGCACACTATCG ACCTGCCCGACAGCGAAATGGATAAGCTCTACCAGAGGGTCAAGCGCCAACTTCGAGAGAATGCTGAGGAGGACGGAACCGGTTGCTT TGAGATATTTCACAAGTGTGATGACGATTGCATGGCATCAATCAGAAATAATACTTACGACCATTCGAAATACAGGGAAGAGGCAATGCAG AACAGAATTCAGATCGACCCTGTGAAGCTTTCTAGTGGGTATAAAGACGTGATCCTCTGGTTCAGTTTTGGAGCATCTTGTTTCATTCTGC TAGCCATTGTTATGGGACTTGTCTTCATTTGCGTGAAGAACGGGAATATGCGTTGTACGATCTGCATT

2863	ATGAACACTCAGATACTCGTCTTCGCCCTCATAGCCATCATTCCCACGAATGCCGACAAAATCTGTCTGGGTCACCACGCGGTAAGTAAC GGGACTAAGGTCAACACACTGACAGAAAGAGGCGTCGAAGTCGTGAACGCCACAGAGACGGTGGAGAGGACAAACATACCCAGAATTT GCTCGAAGGGCAAGAAGACCGTAGATCTGGGCCAGTGCGGGCTGTTGGGCACCATCACAGGCCCCCCCCAGTGCGATCAGTTCCTGGA ATTCTCTGCCGACCTTATCATTGAAAGGAGAGAGGGGAGCGATGTTTGTTACCCCGGGAAGTTCGTTAACGAGGAGGCCCTTAGACAGA TCCTTCGGGAATCCGGTGGGATTGACAAGGAAGCCATGGGATTCACCTACTCTGGTATACGGACTAACGCAGCCACCTCAGCCTGCAGA AGAAGCGGTAGTTCTTTTTATGCCGAAATGAAATGGCTGTTGTCAAACACCGACAATGCTGCCTTCCCACAGATGACCAAATCATACAAGA ATACCCGCAAGTCTCCCGCCCTGATTGTGTGGGGCATCCATCACTCGGTGTCCACCGCCGAACAGACTAAGCTCTACGGCTCAGGGAAT AAGCTGGTGACTGTTGGCAGCTCAAACTACCAGCAGTCCTTCGTGCCTAGCCCTGGTGCCCGTCCCCAGGTGAACGGCCAGAGCGGTC GGATTGACTTCCATTGGCTTATGTTGAACCCAAACGATACCGTGACCTTCTCCTTCAATGGTGCCTTTATAGCACCCGATCGGGCGTCCTT TCTGCGCGGTAAATCTATGGGGATTCAAAGCGGCGTCCAGGTGGATGCTAATTGTGAAGGAGATTGCTATCACAGCGGGGGAACTATAA TTTCTAACCTGCCATTTCAAAATATTGATAGCCGGGCAGTGGGAAAGTGTCCTCGCTATGTTAAACAAAGATCTCTTTTGCTGGCTACCGG AATGAAGAACGTGCCAGAGATCCCCAAAGGAAGGGCACTGTTCGGAGCAATCGCCGGGTTTATTGAAAATGGATGGGAGGGCCTCATTG ACGGTTGGTACGGGTTCCGCCACCAGAACGCTCAAGGAGAGGGGACGGCAGCGGATTATAAGAGCACTCAGTCCGCAATCGACCAGAT CACAGGAAAGCTGAATCGCCTCATTGAGAAAACAAACCAGCAGTTCGAGTTGATTGATAACGAATTCAACGAGGTGGAAAAACAGATCGG CAATGTGATAAACTGGACCCGCGACTCCATTACTGAGGTCTGGTCTTACAACGCAGAACTGCTTGTCGCAATGGAGAATCAACACACCAT AGATCTCGCTGATTCTGAAATGGATAAGCTGTACGAAAGGGTGAAAAGACAGCTGAGAGAGAATGCTGAAGAAGATGGCACCGGTTGCT TCGAGATCTTTCACAAATGTGACGATGACTGCATGGCCTCCATTAGGAATAACACATATGATCATTCCAAGTACAGGGAAGAGGCCATGC AGAACCGCATTCAGATTGATCCGGTTAAGCTGAGTTCCGGGTACAAAGATGTCATACTATGGTTTTCTTTCGGCGCGAGTTGCTTCATTTT GCTTGCCATAGTGATGGGTTTAGTTTTCATCTGCGTGAAGAACGGAAACATGCGCTGCACTATTTGCATA
2864	ATGAATACCCAGATTCTTGTTTTTGCTCTTATAGCCATAATACCCACGAACGCAGACAAAATCTGCTTGGGCCACCACGCAGTATCTAACG GCACCAAGGTGAATACTCTCACCGAACGTGGGGTGGAGGTCGTGAATGCTACAGAAACAGTGGAGCGCACAAACATTCCTCGTATTTGT TCAAAAGGGAAGAAGACAGTGGATCTGGGCCAGTGCGGCCTATTAGGAACAATCACAGGGCCCCCTCAATGTGACCAGTTTCTGGAGTT TTCCGCGGACCTCATTATCGAGCGACGGGAAGGGAGCGACGTATGCTACCCCGCGAAAATTCGTCAATGAGGAGGCCCTGCGTCAGATT CTTAGGGAGTCAGGCGGCATCGACAAGGAAGCTATGGGATTTACATATTCTGGAATTCGCACTAACGGCGCCACCTCCGCTTGTCGACG GAGCGGATCGTCCTTCTATGCCGAAATGAAGTGGCTGCTGTCGAATACGGACAACGCTGCTTTTCCACAAATGACTAAGAGCTATAAGAA TACCAGAAAGTCTCCTGCGTTAATTGTATGGGGAATACATCATAGTGTGTCTACCGCCGAGCAGACAAAACTGTACGGTTCCGGCAATAA GCTAGTCACTGTGGGCTCTTCCAATTACCAGCAGTCGTTTGTCCCCTCGCCAGGGGCCCGCCCGCAAGTAAATGGACAATCTGGGCGTA TCGATTTCCATTGGCTTATGCTGAACCCTAATGACACCGTGACATTTTCATTCAATGGGGCTTTCATTGCCCCCGACAGGGCTAGCTTCCT TCGCGGCAAGAGCATGGGGATACAATCCGGGGTTCAAGTGGATGCCAACTGCGAGGGCGACTGTTACCACTCAGGCGGAACCATCATT TCAAATCTGCCCTTTCAGAACATTGACAGCCGCGCCGTGGGCAAGTGTCCACGATATGTGAAGCAGAGATCACTGCTCCTTGCAACCGG TATGAAAAATGTTCCTGAGATACCGAAGGGCCGGGGGCTCTTCGGTGCTATCGCTGGATTCATTGAAAACGGGTGGGAAGGACTGATCG ACGGTTGGTATGGATTTAGGCACCAGAACGCACAGGGCGAAGGGACCGCCGCTGATTATAAAAGCACGCAGAGCGCTATAGATCAGATC ACCGGAAAGCTGAACCGCCTGATCGAGAAAACGAACCAACAGTTCGAACTGATTGATAACGAGTTTAACGAAGTGGAAAAACAGATTGGT AACGTGATCAACTGGACTAGAGACTCCATCACCGAAGTGTGGTCATATAACGCAGAGTTACTCGTCGCGATGGAAAACCAACACACCATC GACCTGGCAGACAGTGAGATGGATAAACTCTACGAACGGGTCAAGAGACAACTACGTGAGAATGCCGAGGAAGATGGCACAGGTTGTTT TGAAATTTTTCACAAGTGTGACGACGACTGCATGGCATCAATAAGAAACAATACTTACGACCATAGTAAGTACCGGGAGGAAGCTATGCA GAATAGAATCCAAATAGATCCAGTGAAATTGAGTAGCGGATATAAGGATGTGATTCTATGGTTTAGCTTCGGCGCTAGTTGTTTTATCCTTT TGGCTATAGTGATGGGGCTCGTTTTTATCTGCGTGAAGAACGGAAACATGAGATGTACGATCTGCATT
2865	ATGAATACCCAAATTCTCGTTTTTGCCCTCATCGCCATTATCCCTACTAACGCCGATAAGATCTGTCTGGGGCATCATGCCGTCTCCAACG GGACCAAGGTAAACACTCTGACCGAGCGCGGTGTCGAAGTAGTCAATGCTACAGAGACAGTTGAACGCACTAATATTCCCAGGATCTGT AGTAAAGGAAAGCGTACCGTGGACCTAGGACAATGTGGCTTGCTCGCCACTATCACCGGCCCTCCACAGTGCGACCAATTTCTGGAATT CTCCGCTGATCTGATTATCGAGAGAAGAGAAGGCAGTGACGTGTGCTACCCGGGAAAGTTTGTCAACGAAGAGGCCTTAAGGCAGATCC TGCGCGAGAGCGGGGGTATAGATAAAGAAGCCATGGGCTTCACTTACAGTGGAATTAGGACAAACGGCGCCACGTCGGCCTGCAGAAG GTCTGGGTCCTCATTCTACGCCGAGATGAAGTGGCTTCTGTCGAATACCGACAATGCAGCATTTCCACAAATGACTAAATCCTATAAGAAC ACCCGGAAGTCACCAGCCCTCATAGTGTGGGGGATCCATCACTCCGTGAGCACTGCAGAGCAAACAAAGCTGTATGGGTCCGGGAATAA ACTGGTAACCGTTGGCAGTAGCAATTATCAGCAGTCATTTGTTCCCTCCCCAGGTGCCCGGCCTCAGGTGAATGGTCTGTCTGGTCGCAT CGACTTTCACTGGCTGATGCTGAATCCAAATGATACAGTCACATTTTCGTTTAATGGAGCCTTCATCGCGCCCGACAGGGCTAGCTTCCTA CGCGGCAAGAGCATGGGCATTCAGAGCGGCGTCCAGGTTGACGCCAATTGCGAGGGAGATTGTTATCATAGCGGAGGCACAATCATCT CAAACCTCCCCTTTCAGAATATAGACTCCAGAGCCGTTGGGAAGTGCCCTAGATACGTGAAACAGCGATCCCTGCTCCTAGCTACAGGAA TGAAGAATGTGCCAGAAATCCCCAAGGGTCGGGGGCTCTTTGGAGCCATCGCCGGTTTTATCGAAAATGGTTGGGAAGGCCTCATTAAT GGCTGGTACGGCTTTCGACACCAGAATGCCCAGGGTGAAGGAACAGCGGCAGACTATAAGTCCACACAGAGTGCTATAGACCAGATCAC CGGCAAGCTTAACAGACTGATTGAAAAGACCAATCAGCAGTTCGAGTTGATCGACAACGAATTCAACGAGGTTGAGAAACAAATAGGAAA CGTGATAAATTGGACCCGCGATTCTATAACAGAAGTGTGGTCATATAATGCGGAACTGCTCGTGGCAATGGAGAACCAACATACAATAGA CCTTGCAGACTCCGAAATGGATAAGCTGTACGAGCGCGTCAAGCGACAGCTCAGAGAAAATGCAGAGGAGGATGGAACAGGCTGTTTTG AAATTTTCCACAAGTGTGACGACGATTGCATGGCCTCCATCAGAAATAACACATATGACCATTCAAAGTATCGCGAAGAAGCTATGCAGAA TCGAATCCAAATAGATCCTGTGAAGTTGTCGTCCGGCTACAAAGATGTTATCCTGTGGTTCAGTTTCGGGGCCTCATGCTTCATCCTGCT GGCCATAGTGATGGGTCTCGTGTTCATCTGCGTGAAGAACGGCAACATGCGCTGCACTATCTGCATT

2866	ATGAATACCCAGATTTTGGTGTTCGCACTTATCGCAATAATCCCGACTAACGCTGATAAAATTTGTTTAGGACATCATGCCGTTTCTAACG GAACTAAGGTGAACACACTTACCGAGAGAGGTGTGGAAGTGGTGAACGCTACCGAAACTGTGGAGCGGACTAACATCCCCCGGATATGC TCAAAGGGCAAGAGAACTGTGGATTTAGGTCAATGTGGTCTACTGGGCACAATTACCGGCCCACCCCAATGCGACCAGTTCCTGGAGTT TAGCGCCGACCTGATTATTGAGAGACGTGAGGGCTCAGACGTATGTTATCCTGGGAAATTTGTCAACGAAGAAGCCTTGCGGCAGATCC TACGTGAGAGCGGGGGCATTGACAAAGAGGCCATGGGGTTTACATACTCAGGCATCCGAACTAACGGGGCAACAAGTGCATGTAGACG ATCTGGGTCTAGCTTCTATGCCGAGATGAAATGGCTCCTCTCGAACACCGATAACGCCGCTTTTCCGCAGATGACTAAATCTTATAAGAAT ACAAGGAAGTCCCCCGCTTTAATTGTCTGGGGCATCCACCACTCTGTGAGCACCGCCGAACAGACAAAATTGTACGGCTCGGGGTCCAA GCTTGTCACTGTGGGCTCATCCAATTACCAGCAAAGCTTTGTGCCCTCCCCCGGGGCAAGACCTCAGGTGAACGGGTTGAGCGGAAGG ATTGATTTCCATTGGCTGATGTTGAACCCTAACGACACAGTCACCTTCTCGTTTAACGGGGCCTTCATTGCCCCAGATAGGGCCTCTTTTC TGCGCGGAAAGTCGATGGGGATCCAGTCAGGGGTGCAGGTGGACGCCAACTGTGAAGGCGACTGTTATCACAGCGGAGGAACGATAAT TTCCAATCTACCCTTTCAGAATATAGATTCAAGAGCCGTTGGGAAATGTCCCAGGTACGTCAAGCAGAGGTCGCTCCTCTTGGCTACCGG CATGAAGAATGTTCCTGAAATCCCGAAGGGCAGAGGACTCTTTGGCGCCATAGCCGGCTTTATTGAAAATGGATGGGAAGGATTAATTGA CGGCTGGTACGGCTTCCGGCATCAGAATGCACAGGGCGAGGGCACGGCGGCTGACTACAAGTCTACTCAGAGTGCCATAGACCAGATT ACCGGGAAGTTAAATCGTCTGATCGAGAAGACCAACCAACAGTTTGAACTCATTGACAATGAGTTCAACGAAGTGGAGAAACAGATCGGC AACGTAATCAATTGGACTCGCGATTCCATTACCGAGGTGTGGTCTTATAACGCAGAACTCTTGGTGGCCATGGAAAACCAACACACGATT GATCTCGCAGACAGCGAGATGGACAAGTTGTATGAGAGAGTTAAGCGCCAGCTCAGGGAGAATGCCGAAGAGGACGGAACCGGTTGCT TTGAGATCTTTCACAAATGTGATGATGACTGCATGGCGAGCATTCGTAATAATACTTATGACCATAGTAAATACCGGGAGGAAGCCATGCA GAATCGCATCCAAATTGACCCCGTAAAGCTAAGTTCAGGTTATAAGCATGTTATTCTTTGGTTTTCATTCGGAGCTTCATGCTTCATCCTTC TTGCAATTGTAATGGGACTGGTTTTCATTTGTGTAAAGAACGGAAATATGCGATGTACGATTTGTATT
2867	ATGAATACTCAAATCCTTGTCTTTGCGCTGATCGCTATCATTCCAACAAATGCAGATAAAATCTGTCTAGGACATCATGCTGTGTCAAACG GTACTAAAGTGAATACACTGACGGAGAGGGGAGTCGAGGTGGTGAACGCAACAGAGACAGTGGAACGCACAAATATCCCCCGCATCTGT TCTAAGGGGAAAAAAACCGTGGACTTGGGGCAGTGCGGTTTACTGGGGACCATTACTGGTCCGCCCCAGTGCGACCAGTTCCTAGAGTT TTCCGCAGACCTGATTATAGAGAGACGCGAAGGAAGCGACGTATGTTACCCCGGGAAGTTTGTCAATGAAGAAGCCCTCAGACAGATTC TTCGGGAGTCCGGAGGCATCGATAAAGAGGCTATGGGCTTCACCTACTCCGGCATTCGTACAAATGGTGCTACGTCTGCCTGTCGACGC AGTGGGTCATCCTTTTACGCCGAAATGAAGTGGCTATTGAGCAATACCGACAACGCAGCTTTCCCTCAGATGACTAAGAGCTATAAAAAC ACCCGCAAGAGCCCAGCTCTGATCGTATGGGGAATTCACCACTCAGTTTCGACTGCAGAGCAGACTAAACTGTATGGCTCCGGGAATAA GCTGGTTACCGTGGGTAGCTCTAACTACCAGCAGTCATTCGTTCCTTCACCCGGCGCACGCCCCCAAGTGAATGGGCAATCTGGGCGCA TAGACTTCCACTGGCTTATGCTAAACCCAAACGACACTGTGACCTTTTCATTCAACGGCGCATTCATTGCACCTGACCGCGCGTCATTTCT CCGGGGCAAGAGCATGGGGATCCAGAGCGGCGTACAGGTGGACGCGAATTGCGAGGGGGATTGCTACCATTCAGGAGGAACCATTATT TCTAACCTCCCGTTCCAAAATATAGACTCTCGTGCTGTGGGAAAGTGCCCGAGATACGTCAAACAGCGTTCTCTACTCCTGGCTACGGGG ATGAAGAACGTGCCCGAGATTCCGAAAGGGCGCGGACTGTTTGGCGCCATCGCGGGATTTATTGAGAACGGCTGGGAGGGGCTTATTG ATGGTTGGTATGGCTTCCGGCACCAGAACGCTCAGGGGGAGGGGACTGCAGCTGATTACAAGTCAACACAATCAGCAATCGATCAGATT ACCGGAAAGCTGAACAGACTCATCGAGAAGACGAATCAACAGTTTGAATTGATCGATAACGAATTCAACGAAGTCGAGAAACAAATCGGA AACGTCATTAACTGGACTAGAGATTCCATCACCGAGGTGTGGTCTTATAACGCCGAGCTCCTGGTGGCGATGGAGAACCAGCACACAATT GATCTCGCTGACAGCGAGATGGACAAGTTGTACGAGAGGGTAAAACGACAGCTCCGGGAAAATGCCGAGGAAGATGGGACAGGGTGCT TCGACATCTTTCACAAGTGTGACGACCACTGTATGGCCTCAATTCGAAATAACACCTATGATCACTCCAAATACCGCGAGGAAGCTATGC AGAACAGAATCCAGATTGACCCTGTGAAGCTGAGCAGTGGGTATAAAGACGTCATCTTATGGTTCAGCTTCGGGGCTTCTTGCTTCATAC TGCTTGCCATCGTTATGGGCCTGGTGTTTATCTGTGTCAAGAATGGTAACATGAGGTGCACTATCTGTATA
2868	ATGAACACCCAAATTCTTGTATTTGCTCTGATCGCCATTATTCCAACCAATGCCGACAAGATCTGTTTGGGGCACCATGCCGTTTCTAATG GCACAAAAGTCAATACATTAACTGAGCGCGGCGTAGAAGTCGTGAACGCTACCGAAACAGTTGAGAGAACCAACATCCCTCGAATATGTT CCAAGGGGAAGAAAACAGTGGATCTCGGGCAGTGCGGCTTGCTGGGAACAATAACTGGCCCGCCTCAGTGTGATCAGTTCCTGGAATTT AGTGCTGACCTGATAATCGAGCGAAGAGAGGGGAGCGATGTGTGTTATCCCGGGAAATTTGTAAACGAGGAGGCCCTGCGGCAGATACT TAGAGAGAGTGGGGGAATCGACAAAGAAGCGATGGGGTTTACCTACTCTGGAATCAGAACTAACGGGGCTACAAGCGCCTGCCGGCGC AGCGGATCGTCCTTCTACGCAGAGATGAAGTGGCTGCTCAGCAACACTGACAATGCCGCTTTTCCTCAAATGACCAAGTCTTACAAGAAC ACCCGTAAAAGTCCCGCGCTCATAGTATGGGGTATCCACCATTCTGTCTCAACAGCAGAACAGACCAAACTGTACGGCAGCGGCAACAA ACTGGTGACAGTGGGATCTAGCAATTACCAGCAGAGTTTTGTCCCATCCCCTGGCGCGCGGCCCCAGGTGAATGGGCAAAGTGGACGG ATAGACTTCCACTGGTTGATGCTGAACCCAAATGATACCGTGACATTTTCTTTCAACGGTGCATTTATAGCCCCTGACCGTGCAAGCTTCT TGAGAGGTAAGAGCATGGGAATTCAGTCAGGTGTTCAGGTGGATGCCAATTGCGAGGGCGATTGCTACCACAGCGGCGGCACTATTATT TCTAATCTTCCTTTCCAGAATATAGACTCCCGTGCTGTGGGTAAATGCCCTAGGTACGTAAAGCAAAGGAGCTTACTGTTGGCAACAGGC ATGAAGAATGTGCCAGAAATTCCCAAGGGGCGGGGCCTGTTTGGCGCCATTGCAGGCTTTATTGAGAATGGCTGGGAGGGCCTAATTGA TGGTTGGTACGGCTTCAGGCACCAGAACGCACAGGGGGAAGGCACAGCTGCGGATTACAAAAGTACCCAGAGCGCCATAGATCAAATAA CCGGCAAGTTAAACCGATTGATTGAGAAAACAAACCAGCAATTTGAGCTGATAGACAATGAATTCAACGAAGTCGAAAAGCAGATTGGCA ACGTGATCAATTGGACCCGGGATAGCATTACTGAGGTATGGAGCTATAATGCAGAGTTGCTGGTGGCCATGGAAAACCAGCATACCATC GATCTGGCCGACTCAGAAATGGATAAGCTCTACGAGCGGGTGAAGACGCAGCTGAGAGAAAATGCCGAAGAAGACGGAACCGGCTGCT TTGAGATTTTTCACAAGTGCGACGATGATTGCATGGCCAGTATTAGAAACAACACCTATGATCACTCCAAGTACCGCGAAGAAGCCATGC AGAACCGCATCCAAATCGATCCGGTCAAACTCTCCTCTGGCTATAAGGATGTAATCCTGTGGTTCTCTTTTGGAGCGTCCTGCTTCATATT ATTGGCTATTGTGATGGGCTTGGTGTTTATCTGTGTTAAGAACGGAAACATGCGGTGTACAATCTGCATT

2869	ATGAATACACAGATACTCGTGTTCGCCCTTATTGCTATCATCCCTACAAATGCTGACAAGATATGTCTGGGCCACCACGCCGTGAGCAAT GGCACCAAGGTTAATACCCTAACCGAACGGGGAGTTGAGGTGGTTAACGCTACCGAGACCGTAGAGCGCACTAATATTCCACGCATTTG CTCGAAAGGCAAGAAAACAGTGGACTTAGGCCAATGTGGACTCCTCGGAACAATAACCGGCCCACCACAGTGCGATCAGTTCCTCGAGT TTTCAGCCGACCTAATTATTGAGCGACGGGAAGGTAGCGATGTGTGTTATCCAGGGAAGTTCGTCAATGAAGAGGCCTTGCGGCAGATA CTGCGCGAAAGTGGGGGCATAGACAAGGAGGCTATGGGGTTTACATACAGTGGAATTCGCACCAATGGGGCCACTAGCGCCTGTAGGA GATCAGGCTCCTCTTTCTACGCTGAGATGAAATGGTTACTGTCAAATACTGATAACGCCGCCTTTCCTCAGATGACAAAATCCTATAAGAA CACTAGGAAGTCACCTGCTCTAATCGTTTGGGGAATCCATCACTCCGTGAGCACCGCTGAGCAGACCAAGCTTTATGGCTCAGGAAACAA GCTCGTGACGGTGGGCTCCAGTAACTATCAACAAAGCTTCGTGCCTTCTCCAGGAGCAAGGCCCCAGGTGAATGGACAGTCCGGGAGA ATCGACTTTCACTGGCTCATGCTGAATCCAAACGATACCGTTACCTTTTCATTCAATGGGGCCTTCATCGCCCCAGACAGGGCCAGCTTC CTCCGCGGCAAAAGCATGGGCATACAAAGCGGCGTGCAGGTGGATGCTAATTGCGAGGGTGACTGCTATCATTCAGGCGGCACCATAAT CAGTAACCTGCCCTTTCAAAATATAGATTCGCGGGCAGTAGGTAAATGCCCTCGTTATGTGAAACAGCGGTCCTTGCTGCTTGCGACTGG AATGAAAAACGTGCCCGAGATTCCCAAGGGCCGTGGATTATTTGGCGCTATCGCCGGGTTCATCGAGAACGGCTGGGAGGGTCTCATTG ATGGATGGTACGGTTTCAGACACCAGAACGCCCAGGGGGAGGGGACCGCAGCCGACTACAAATCAACCCAGTCCGCCATTGATCAGAT CACAGGGAAGCTAAATCGCCTGATTGAGAAGACAAATCAGCAGTTCGAGCTGATCGACAACGAATTCAATGAGGTAGAGAAACAGATTGG GAACGTTATTAATTGGACCAGAGATTCCATAACAGAGGTATGGTCTTATAACGCAGAACTGCTCGTTGCGATGGAGAACCAGCATACAAT CGACCTGGCAGACAGTGAAATGGACAAGCTCTATGAGAGGGTCAAGAGGCAATTGAGAGAGAATGCCGAAGAAGACGGGACCGGTTGC TTCGAGATCTTCCACAAATGTGATGATGACTGCATGGCAAGCATTAGAAACAACACCTACGATCACTCCAAATACCGAGAGGAGGCTATG CAAAATCGGATACAGATCGATCCTGTCAAGCTGTCTTCTGGTTACAAAGATGTTATTCTGTGGTTCAGTTTTGGGGCCTCATGTTTCATCTT ACTGGCAATCGTGATGGGACTCGTTTTCATCTGTGTTAAAAATGGCAACATGAGGTGTACCATTTGTATC
2870	ATGAACACACAAATTTTAGTGTTTGCTCTGATTGCCATTATCCCCACCAACGCTGATAAGATTTGTCTGGGGCACCACGCAGTGTCTAATG GCACCAAAGTCAACACACTTACAGAGAGAGGAGTGGAGGTGGTTAACGCCACTGAAACCGTCGAGAGAACCAATATTCCTCGGATCTGC TCCAAGGGCAAGAAAACAGTTGATCTAGGACAGTGTGGGTTGCTAGGGACAATCACCGGACCACCTCAGTGTGATCAGTTCTTAGAGTTC TCCGCTGATCTAATTATTGAGCGAAGAGAAGGGTCCGATGTGTGCTACCCCGGTAAATTTGTCAATGAAGAAGCCTTGAGGCAGATTCTG AGAGAAAGCGGCGGCATCGATAAGGAGGCAATGGGTTTCACTTACTCTGGAATCCGGACCAACGGAGCCACTAGCGCTTGTCGAAGGA GTGGATCGTCCTTCTACGCCGAGATGAAATGGCTCTTGTCAAATACAGATAATGCCGCATTTCCCCAGATGACTAAGAGCTACAAGAATA CACGTAAGAGCCCGGCTCTGATCGTCTGGGGTATTCATCATAGTGTGTCTACTGCAGAGCAGACCAAACTATACGGTAGTGGCAACAAAC TTGTCACCGTCGGCAGTTCTAATTACCAGCAATCCTTTGTCCCTTCCCCCGGCGCTCGGCCTCAGGTGAATGGACAGAGCGGCAGGATT GACTTCCACTGGCTTATGCTGAACCCTAATGATACTGTTACCTTCTCGTTTAATGGGGCGTTTATTGCTCCCGATAGGGCCAGTTTTCTTC GGGGCAAGTCCATGGGGATCCAGTCAGGTGTGCAGGTCGATGCTAATTGTGAGGGCGACTGCTATCATTCTGGCGGCACGATCATCTCA AACTTGCCCTTCCAAAATATAGATAGCCGAGCAGTGGGAAAATGCCCTCGTTACGTCAAGCAGAGGAGCCTCCTGTTAGCCACCGGGAT GAAAAATGTTCCAGAAATACCCAAGGGGAGGGGCTTATTTGGCGCTATTGCAGGATTTATCGAGAACGGTTGGGAGGGACTCATTGACG GATGGTATGGCTTCCGCCACCAGAACGCCCAGGGGGAGGGTACCGCAGCTGATTACAAGTCCACACAGAGTGCCATAGATCAAATCACC GGTAAGCTGAATAGGCTGATCGAAAAAACCAATCAGCAGTTTGAGCTCATCGATAATGAGTTCAACGAGGTGGAGAAACAGATTGGTAAT GTCATCAACTGGACAAGGGACTCGATTACCGAAGTGTGGAGCTATAACGCAGAGCTCTTGGTTGCCATGGAGAATCAGCACACCATCGA TTTAGCAGATAGCGAAATGGACAAACTATACGAAAGGGTGAAACGGCAACTTAGGGAGAACGCCGAAGAGGATGGGACCGGGTGCTTC GAAATTTTTCACAAGTGTGATGACGACTGCATGGCCTCTATCCGGAATAATACCTACGATCACAGCAAATACAGGGAGGAAGCCATGCAG AACCGCATCCAGATCGATCCAGTCAAGCTGTCTTCTGGCTATAAGGACGTGATCCTGTGGTTCTCTTTCGGCGCTAGTTGTTTTATCCTGC TGGCCATCGTGATGGGCCTTGTTTTCATCTGTGTGAAAAACGGGAATATGAGGTGCACAATCTGCATT
2871	ATGAACACACAGATCCTCGTCTTTGCACTCATCGCAATCATCCCCACCAACGCTGATAAAATTTGTCTCGGACATCATGCCGTGAGCAATG GGACCAAAGTTAACACCCTAACGGAGAGAGGTGTCGAGGTGGTGAACGCGACAGAAACCGTAGAACGAACCAATATTCCAAGGATCTGC AGTAAGGGAAAAAGGACCGTGGATTTGGGGCAGTGTGGATTGCTGGGGACTATCACAGGACCACCACAGTGCGACCAATTCCTTGAATT TTCCGCGGATCTGATTATTGAGAGGAGGGAGGGATCTGACGTCTGTTACCCAGGCAAGTTCGTTAATGAGGAAGCGCTGAGACAGATTC TGCGAGAGAGCGGAGGCATAGACAAAGAGGCTATGGGTTTTACCTATTCTGGGATACGCACAAACGGAGCTACAAGCGCATGCCGGAG AAGTGGAAGCTCATTTTATGCGGAAATGAAATGGCTCCTCTCCAACACCGACAATGCAGCATTCCCTCAGATGACCAAATCATATAAAAAC ACCAGAAAAAGTCCGGCTCTCATAGTGTGGGGTATACATCATTCCGTCTCGACCGCAGAGCAAACTAAGCTCTACGGCTCTGGAAACAAA CTCGTGACTGTCGGCAGTTCTAATTACCAGCAGTCTTTTGTACCTAGCCCCGGGGCTAGGCCACAGGTGAATGGCTTATCTGGGCGCAT AGACTTCCACTGGCTTATGCTGAACCCAAACGATACAGTGACCTTCTCCTTTAACGGCGCCTTCATTGCCCCCGACAGAGCTAGTTTCCT GCGCGGAAAATCAATGGGAATCCAGAGTGGGGTGCAGGTGGATGCAAACTGCGAAGGCGACTGCTATCACTCAGGCGGCACAATAATT AGTAACCTACCATTCCAGAACATTGATTCACGCGCCGTCGGGAAGTGTCCGAGATACGTGAAACAGCGGTCTCTCCTGTTAGCAACAGG CATGAAGAATGTGCCTGAGATTCCCAAAGGCCGGGGGCTCTTCGGAGCCATTGCTGGTTTCATCGAGAACGGGTGGGAGGGTCTGATC GATGGCTGGTATGGTTTCCGGCATCAGAACGCGCAGGGGGAGGGCACAGCCGCAGATTACAAGAGTACTCAGAGTGCCATCGATCAGA TTACAGGGAAGCTGAATAGAATAATCGAAAAAACTAATCAGCAGTTTGAACTGATAGACAACGAATTCAACGAGGTCGAGAAGCAGATCG GGAATGTGATCAACTGGACCCGCGACTCAATCACAGAGGTTTGGAGTTATAATGCCGAGCTCCTCGTTGCCATGGAGAATCAACATACAA TTCACCTTGCTGACTCTGAGATGGATAAACTGTATGAACGAGTTAAGCGACAGTTGAGGGAGAATGCGGAAGAAGACGGGACCGCTGC TTCGAAATATTCCATAAGTGCGACGATGACTGTATGGCATCAATCAGGAACAACACCTATGACCACTCCAAGTATCGAGAGGAAGCGATG CAGAACAGGATCCAGATCGATCCAGTCAAGTTGTCCTCTGGCTATAAGGACGTGATTCTGTGGTTTTCTTTCGGCGCAAGCTGTTTCATTT TATTAGCTATCGTCATGGGTCTAGTGTTCATCTGTGTTAAAAATGGCAATATGCGATGTACAATCTGTATT

2872	ATGAACACCCAAATCCTTGTTTTCGCTCTGATTGCAATTATCCCAACGAATGCTGACAAAATTTGCCTGGGCCACCACGCTGTGTCCAACG GAACTAAGGTAAATACTCTGACTGAACGCGGCGTGGAGGTGGTGAACGCCACCGAAACAGTGGAGAGAACTAATATTCCCCGCATCTGT TCTAAGGGCAAAAAGACAGTGGATCTCGGTCAGTGCGGGCTGTTGGGAACAATTACCGGACCTCCCCAATGCGATCAGTTCCTAGAGTT TTCCGCCGACCTGATAATCGAAAGACGTGAAGGCTCTGACGTGTGTTACCCCGGAAAGTTTGTGAATGAAGAAGCACTGCGGCAGATCT TACGCCAGAGTGGCGGCATTGACAAGGAGGCAATGGGATTTACCTACTCCGGAATCAGGACAAATGGCGCAACCAGTGCTTGTAGGCG CTCCGGAAGCTCCTTTTATGCCGAGATGAAATGGCTTCTGTCCAACACGGACAACGCCGCTTTCCCTCAAATGACCAAATCCTACAAAAAT ACCCGTAAGTCACCGGCGCTGATTGTATGGGGGATACACCACAGTGTTTCTACCGCCGAACAAACCAAGCTGTACGGCAGCGGAAATAA ACTGGTAACTGTCGGATCATCCAATTACCAGCAGTCCTTCGTCCCTAGCCCTGGGGCGCGCCCGCAGGTCAATGGGCAGTCTGGAAGAA TTGACTTTCATTGGCTGATGTTAAATCCAAACGACACAGTTACATTCAGTTTCAACGGGGCCTTTATTGCACCCGATAGAGCATCCTTTCT GCGAGGCAAAAGTATGGGGATTCAATCTGGGGTGCAGGTTGATGCGAATTGCGAGGGAGATTGCTATCACTCTGGCGGAACCATCATCA GCAACCTCCCATTCCAAAATATTGACTCACGCGCTGTGGGAAAGTGCCCTAGGTATGTGAAGCAGCGGTCCTTGCTTTTAGCTACCGGAA TGAAAAACGTGCCAGAAATCCCTAAGGGGCGCGGATTATTCGGCGCCATTGCTGGCTTCATTGAGAATGGATGGGAGGGCCTAATCGAC GGTTGGTATGGTTTTCGTCATCAGAACGCACAAGGGGAAGGGACTGCTGCCGACTACAAGTCTACCCAGTCCGCTATAGACCAGATTAC CGGTAAGCTAAATCGGCTTATAGAGAAGACTAACCAACAGTTCGAGCTGATCGACAACGAATTTAACGAAGTGGAGAAGCAGATTGGGAA CGTCATCAACTGGACGCGAGACTCTATTACTGAAGTCTGGTCTTACAATGCTGAGTTGCTCGTGGCTATGGAAAACCAGCATACTATAGA CCTGGCCGACTCAGAGATGGATAAGTTATATGAACGCGTGAAGCGGCAGCTCCGAGAAAATGCCGAAGAGGACGGAACAGGGTGTTTT GAGATCTTCCACAAATGTGACGATGACTGCATGGCATCTATACGGAACAATACCTATGACCATTCCAAGTACCGTGAGGAAGCCATGCAA AATAGAATACAAATCGATCCAGTCAAACTCTCTAGCGGCTACAAGGACGTTATCTTGTGGTTCTCCTTCGGAGCGAGCTGTTTTATTTTGC TCGCGATTGTGATGGGATTGGTGTTTATTTGCGTTAAAAACGGAAACATGCGGTGCACCATCTGTATC
2873	ATGAATACCCAGATCCTGGTCTTCGCTCTGATCGCAATCATTCCTACCAACGCTGACAAAATTTGCCTGGGCCACCACGCAGTTAGTAAT GGAACAAAAGTGAATACTCTTACAGAGCGGGGTGTCGAGGTAGTAAACGCTACGGAAACCGTAGAGCGCACAAATATCCCCAGAATCTG TAGCAAAGGTAAAAGGACAGTAGACTTGGGCCAGTGTGGACTGCTCGGAACAATTACAGGCCCCCCTCAGTGTGATCAGTTCCTGGAAT TCTCAGCCGACCTTATCATCGAACGAAGGCAGGGGAGCGATGTCTGCTACCCAGGCAAGTTTGTGAATGAAGAGGCCCTTCGTCAAATA CTCCGAGAATCGGGGGGAATCGATAAAGAAGCAATGGGATTTACCTATAGCGGAATCAGAACTAACGGCGCAACTTCCGCCTGCAGGCG GTCTGGCAGCTCCTTTTATGCCGAGATGAAGTGGCTATTGAGCAATACTGACAACGCAGCCTTCCCTCAAATGACCAAGTCATACAAAAA TACACGCAAGTCTCCAGCTCTGATCGTGTGGGGCATTCACCATAGCGTCAGCACAGCCGAGCAAACAAAGCTCTACGGCAGCGGCAACA AACTCGTGACCGTCGGCAGTTCTAACTACCAGCAGTCCTTTGTGCCCAGCCCAGGGGCCCGTCCACAGGTTAACGGACTGTCTGGCCG GATCGATTTCCACTGGCTGATGTTAAATCCCAACGACACAGTCACGTTTAGCTTTAACGGAGCCTTTATCGCCCCCGACAGGGCTAGCTT TCTTAGGGGCAAGAGCATGGGAATTCAATCAGGGGTTCAAGTGGACGCAAACTGCGAGGGCGATTGCTATCACTCCGGTGGAACTATTA TAAGTAATCTACCATTCCAAAACATCGATTCAAGGGCGGTGGGTAAGTGTCCTCGCTACGTTAAGCAGCGCAGCCTGCTACTCGCGACG GGAATGAAAAATGTCCCGGAGATACCTAAAGGCAGAGGCCTCTTCGGCGCCATCGCAGGTTTCATCGAAAACGGCTGGGAAGGTCTGAT CGACGGTTGGTATGGATTCCGTCACCAAAACGCACAGGGAGAGGGTACAGCCGCAGACTATAAGTCTACCCAGTCAGCTATCGACCAAA TTACTGGAAAACTGAATCGTATCATTGAAAAGACCAATCAACAATTCGAGCTGATCGACAACGAGTTTAATGAAGTTGAGAAACAGATCGG AAATGTGATTAACTGGACACGTGACAGCATTACTGAGGTCTGGAGCTACAACGCGGAACTTCTCGTAGCGATGGAGAACCAGCACACTAT TGACCTGGCTGATAGTGAAATGGATAAACTATATGAAAGGGTCAAAAGACAACTGCGGGAAAATGCCGAGGAAGATGGTACTGGATGTTT TGAAATCTTTCACAAGTGCGATGACGACTGTATGGCTAGCATCAGGAACAATACATACGACCATTCGAAGTATCGCGAAGAGGCCATGCA GAACCGAATACAGATTGATCCGGTTAAGCTGTCTTCTGGATATAAAGACGTCATCCTGTGGTTTAGTTTCGGGGCGAGTTGTTTCATTCTA CTGGCTATTGTCATGGGCCTGGTTTTTATCTGCGTGAAAAATGGCAACATGCGGTGCACGATCTGCATC
2874	ATGAACACACAGATCTTGGTGTTCGCCCTCATCGCGATAATACCCACTAATGCCGACAAAATTTGTTTGGGTCATCACGCCGTGAGCAAT GGCACTAAGGTGAACACGCTTACAGAACGGGGCGTAGAAGTTGTGAACGCCACAGAGACAGTGGAGAGGACAAATATCCCGCGGATCT GCTCAAAGGGGAAAAAAACCGTTGACCTGGGACAGTGCGGCCTGCTGGGGACAATAACAGGCCCCCCCCAGTGTGACCAATTTCTGGA GTTCTCTGCCGACCTGATTATCGAAAGACGGGAGGGCAGTGATGTCTGCTACCCCGGCAAGTTCGTAAATGAAGAGGCTCTGCGGCAGA TCCTACGGGAGTCCGGAGGAATCGATAAGGAGGCAATGGGGTTTACGTATAGCGGCATTAGAACCAACGGAGCAACCAGTGCTTGCAG GAGGAGCGGCAGCTCATTCTACGCGGAAATGAAGTGGCTGCTGTCAAATACAGATAACGCCGCTTTCCCACAGATGACTAAATCCTATAA GAACACCCGCAAGAGTCCTGCTCTCATCGTTTGGGGTATCCACCACTCTGTGTCAACTGCCGAGCAGACCAAACTTTACGGATCTGGGAA TAAGCTTGTGACAGTTGGGAGCAGCAACTATCAGCAGTCATTTGTGCCCTCTCCGGGTGCCAGGCCACAGGTTAATGGACAGTCGGGGC GTATCGACTTCCATTGGCTTATGCTCAATCCTAATGACACTGTTACCTTTTCTTTCAACGGCGCGTTCATTGCCCCCGACAGAGCGAGCTT TCTTAGGGGGAAATCGATGGGTATTCAGAGCGGGGTCCAGGTCGATGCGAACTGCGAGGGCGATTGCTACCACTCAGGGGGAACAATT ATCTCTAATTTACCCTTTCAGAACATCGACTCAAGGGCAGTGGGGAAGTGTCCGCGATACGTTAAACAACGGAGTTTGCTCCTGGCCACA GGCATGAAAAACGTCCCAGAAATCCCCAAAGGCAGAGGGTTGTTTGGGGCGATCGCCGGCTTTATCGAGAACGGATGGGAAGGCTTAAT TGACGGCTGGTACGGCTTCCGCCACCAGAACGCTCAGGGGGAAGGTACCGCCGCTGATTATAAAAGTACGCAAAGCGCTATTGATCAGA TTACGGGCAAATTGAATCGGCTGATTGAGAAAACAAATCAGCAGTTTGAACTGATCGACAACGAGTTTAATGAGGTGGAGAAGCAGATAG GAAATGTAATTAACTGGACTCGGGATTCGATCACGGAGGTTTGGTCCTATAACGCAGAGCTACTGGTCGCCATGGAGAATCAGCACACTA TTCACCTGGCAGATTCTGAGATGGACAAGCTGTACGAGAGGGTAAAAAGACAACTGCGGGAGAACGCTGAAGAGGATGGAACAGGGTG TTTTGAGATCTTTCACAAGTGCGACGATGATTGCATGGCATCTATCCGCAACAATACTTACGACCACTCTAAGTACAGAGAAGAGGCGAT GCAGAATCGAATTCAGATTGACCCGGTGAAGCTGAGCAGTGGATATAAAGATGTAATCCTCTGGTTTAGCTTCGGAGCCTCATGTTTTATC CTCCTTGCTATCGTTATGGGCCTCGTCTTTATATGTGTCAAAAACGGCAATATGCGCTGCACAATCTGCATT

2875	ATGAATACCCAAATTTTGGTTTTTGCTCTGATTGCCATTATTCCCACTAATGCCGACAAGATCTGCCTCGGCCATCATGCCGTGTCTAATG GCACAAAGGTGAATACCTTGACAGAAAGGGGTGTGGAAGTAGTGAATGCCACAGAAACTGTAGAGCGGACCAATATTCCCAGGATTTGC TCTAAGGGAAAGAAGACCGTTGATCTGGGCCAATGCGGCTTGCTGGGCACCATCACTGGGCCACCACAGTGCGACCAGTTCCTGGAATT TAGCGCCGATCTGATAATAGAGAGACGGGAAGGTAGCGACGTCTGCTATCCTGGGAAATTCGTCAACGAGGAAGCTCTGCGGCAGATCC TACGGGACAGTGGCGGGATAGACAAGCAGGCAATGGGCTTTACGTACTCACGAATTCGAACCAATGGGGCGACGTCCGCCTGCCGGAG ATCGGGCTCTTCTTTTTACGCAGAAATGAAGTGGCTGCTGTCGAATACAGATAATGCGGCTTTCCCCCAGATGACCAAGTCATATAAAAAT ACGAGAAAATCGCCCGCACTTATTGTGTGGGGCATCCACCACTCCGTGTCAACTGCCGAGCAAACCAAGCTATATGGCAGCGGCAATAA GCTGGTGACAGTAGGAAGTTCTAATTACCAGCAGTCTTTCGTGCCGAGCCCTGGAGCCAGACCACAGGTGAATGGCCAGTCGGGCAGG ATAGATTTCCACTGGTTAATGCTTAACCCAAACGACACGGTGACCTTCAGTTTCAACGGTGCCTTCATCGCCCCCGATCGCGCCTCCTTTT TGCGCGGCAAGTCTATGGGTATTCAGAGTGGAGTGCAGGTGGACGCAAACTGCGAAGGGGATTGTTATCACAGCGGCGGGACCATAAT CAGCAATCTCCCGTTTCAAAATATCGATTCTCGGGCAGTCGGTAAATGTCCCAGGTACGTTAAGCAACGGAGCCTGCTGCTCGCTACAGG CATGAAGAATGTGCCCGAGATCCCCAAGGGAAGAGGCCTGTTCGGGGCCATCGCTGGTTTCATCGAGAACGGATGGGAGGGATTAATC GACGGGTGGTACGGTTTCCGACACCAGAACGCACAGGGTGAGGGCACTGCCGCTGATTACAAATCAACCCAGTCTGCAATCGACCAGAT CACTGGGAAATTGAACCGATTGATCGAAAAGACGAACCAGCAGTTCGAACTCATTGACAACGAGTTTAATGAAGTCGAGAAGCAGATTGG AAATGTGATCAACTGGACAAGGGATTCTATCACAGAGGTCTGGTCATACAACGCTGAACTCCTTGTGGCCATGGAGAACCAGCACACCAT CGATCTGGCTGATTCAGAAATGGATAAACTATATGAAAGGGTGAAGCGGCAACTTAGGGAAAATGCTGAGGAGGACGGCACGGGATGCT TTGAAATCTTCCACAAGTGTGATGATGACTGTATGGCGTCCATCCGGAACAATACATATGACCATTCAAAATATAGAGAAGAGGCAATGCA GAACAGGATACAGATTGACCCTGTCAAGTTGAGTTCTGGCTACAAGGACGTGATTCTGTGGTTCAGCTTCGGAGCCAGTTGTTTCATCCT CCTGGCCATCGTGATGGGCCTAGTGTTCATCTGCGTGAAAAATGGCAATATGCGCTGCACAATATGTATA
2876	ATGAATACCCAAATCCTCGTTTTCGCACTGATTGCTATTATACCCACCAACGCTGACAAGATTTGTCTGGGTCACCACGCAGTTTCTAACG GCACAAAGGTGAATACTCTCACAGAGCGCGGGGTTGAGGTCGTCAACGCTACTGAGACCGTCGAACGAACCAATATCCCCAGGATCTGT AGCAAGGGAAAGAAAACAGTCGATTTAGGCCAGTGTGGCCTCTTAGGCACCATCACAGGCCCACCCCAATGCGACCAGTTCCTGGAGTT CTCCGCAGATTTAATCATTGAGCGGCGACAGGGCAGTGATGTTTGCTACCCAGGTAAGTTCGTCAATGAAGAAGCCTTGCGCCAGATCC TGCGGGAAAGCGGCGGCATTGACAAAGAGGCTATGGGATTTACTTATTCTGGCATACGAACTAACGGGGCAACGTCTGCTTGTCGACGC TCAGGCTCATCCTTCTACGCAGAAATGAAATGGTTGCTCAGTAATACAGATAATGCCGCATTTCCCCAAATGACCAAGAGCTACAAGAATA CACGGAAGAGCCCAGCCCTGATTGTGTGGGGTATCCACCATTCTGTAAGTACAGCAGAGCAGACAAAACTGTATGGGTCTGGCAACAAA CTGGTGACGGTGGGCAGTAGCAACTACCAACAGAGTTTCGTGCCGTCACCTGGGGCAAGACCCCAGGTTAACGGGCAGTCCGGCCGCA TAGACTTTCACTGGCTCATGCTGAATCCGAACGATACAGTGACTTTTTCATTCAATGGTGCATTTATCGCCCCCGATCGCGCTTCGTTCCT CCGCGGCAAATCTATGGGAATTCAATCCGGGGTACAGGTCGACGCCAACTGCGAGGGCGACTGTTACCATTCCGGCGGGACTATTATAA GTAATCTACCTTTTCAGAACATAGATTCCAGAGCCGTGGGAAAATGCCCTCGCTACGTGAAACAGAGGTCCCTGCTTCTGGCTACTGGAA TGAAGAATGTTCCAGAGATACCTAAGGGACGGGGCCTTTTCGGGGCAATCGCCGGATTCATCGAAAATGGGTGGGAAGGGCTGATCGAT GGCTGGTACGGCTTTCGCCATCAGAACGCACAGGGAGAGGGTACAGCGGCTGATTACAAATCGACTCAAAGTGCTATAGATCAGATTAC CGGCAAACTGAACAGGCTGATCGAGAAAACGAATCAACAGTTCGAACTGATTGATAACGAGTTTAACGAGGTGGAAAAACAAATCGGAAA CGTTATCAATTGGACGAGGGACAGCATTACCGAAGTGTGGTCGTATAATGCAGAGCTGTTAGTCGCCATGGAGAATCAACACACCATCGA TCTTGCCGACAGCGAGATGGACAAGCTTTATGAGCGCGTGAAGAGACAACTGCGAGAAAATGCCGAGGAAGATGGAACCGGATGTTTTG AGATATTCCACAAGTGCGACGATGATTGTATGGCGAGCATCCGCAACAATACCTATGATCACTCCAAATACAGGGAAGAGGCTATGCAAA ATAGAATCCAAATTGATCCCGTCAAACTGTCAAGTGGGTACAAGGATGTGATATTATGGTTTTCCTTTGGCGCCTCATGCTTTATACTGCT GGCTATCGTGATGGGCTTGGTCTTTATTTGCGTGAAGAATGGCAACATGCGGTGTACAATATGTATT
2877	ATGAATACACAAATTCTGGTGTTCGCACTCATTGCTATCATACCCACTAATGCCGATAAGATCTGTCTGGGCCACCACGCTGTTAGCAACG GGACCAAAGTTAACACCCTTACCGAGAGGGGCGTGGAGGTGGTCAACGCAACCGAAACCGTCGAACGGACAAACATCCCGAGGATCTG CAGTAAAGGAAAGAGAACAGTCGACCTCGGGCAATGTGGACTGCTGGGGACTATTACAGGGCCCCCCCAATGTGATCAGTTCCTGGAGT TTTCTGCGGATTTAATCATTGAGAGGCGGGAGGGGAGTGATGTTTGTTACCCAGGTAAGTTTGTTAATGAAGAAGCGCTGCGCCAGATAC TTAGAGAGAGTGGGGGAATCGACAAGGAGGCTATGGGTTTTACCTACAGCGGCATCAGAACAAATGGCGCCACCAGCGCATGCCGCAG GAGCGGTTCCTCATTCTACGCCGAAATGAAGTGGCTCCTATCCAATACAGATAACGCCGCCTTCCCCCAGATGACAAAGTCCTACAAAAA CACCCGGAAGAGCCCTGCCCTAATTGTGTGGGGTATCCACCACTCCGTGTCTACGGCCGAGCAGACCAAGCTGTACGGAAGCGGCAAC AAACTAGTGACCGTAGGATCTAGTAACTATCAGCAGAGTTTCGTACCTTCACCAGGAGCGCGGCCTCAGGTTAATGGTCTGAGCGGGAG GATCGATTTCCATTGGCTTATGCTGAACCCCAATGACACTGTTACGTTTTCTTTCAACGGAGCATTCATCGCTCCAGACCGCGCGAGCTTT CTCCGAGGAAAATCAATGGGGATACAGTCTGGGGTCCAAGTGGACGCTAACTGTGAGGGGGACTGTTATCACTCAGGTGGAACCATTAT TTCTAACTTGCCATTCCAGAACATAGATTCGCGAGCCGTGGGCAAATGTCCCCGTTACGTGAAACAGCGTTCCCTCCTGCTAGCCACCGG GATGAAGAACGTACCCGAAATTCCTAAAGGGAGAGGCCTCTTTGGCGCTATTGCGGGGTTTATTGAAAACGGATGGGAGGGCCTAATTG ACGGCTGGTATGGGTTCAGACATCAGAATGCACAGGGTGAGGGGACAGCTGCTGACTACAAGTCCACGCAGTCTGCAATAGATCAAATA ACCGGCAAACTCAACCGCTTAATCGAGAAAACGAACCAGCAGTTCGAGCTTATAGACAATGAGTTCAATGAGGTTGAAAAACAGATCGGC AACGTGATTAATTGGACTCGGGACTCTATCACCGAGGTCTGGAGCTATAACGCAGAGTTGCTGGTCGCCATGGAGAATCAGCACACAATA GATCTTGCTGACTCGGAAATGGACAAACTTTACGAAAGGGTCAAGCGGCAGCTGAGGGAGAACGCTGAAGAAGATGGGACGGGTGCT TCGAAATTTTTCATAAGTGCGATGATGATTGTATGGCGTCCATTAGAAATAACACCTACGACCATAGTAAATACCGGGAGGAAGCCATGCA GAACAGGATCCAGATCGACCCAGTAAAACTGTCAAGTGGGTATAAGGATGTTATATTGTGGTTCTCGTTCGGCGCCAGTTGCTTCATCCT CCTCGCTATTGTCATGGGCCTTGTGTTTATTTGCGTGAAGAATGGAAATATGCGCTGCACCATTTGCATT

2878	ATGAATACCCAGATACTGGTGTTTGCCTTAATAGCAATTATCCCAACAAATGCCGACAAGATCTGCCTTGGACACCACGCCGTGTCCAAT GGTACTAAAGTGAACACCTTGACCGAGCGGGGGGTGGAGGTGGTAAATGCCACGGAGACTGTCGAACGTACAAATATCCCCCGCATCT GTTCTAAAGGCAAAAAAACTGTCGACTTGGGCCAGTGCGGACTACTGGGGACGATTACTGGTCCGCCTCAGTGCGACCAATTCCTCGAA TTTAGTGCAGATCTCATTATTGAAAGACGAGAGGGGAGTGACGTTTGCTATCCAGGCAAGTTCGTTAACGAGGAAGCCCTTAGGCAAATT TTCAGGGCAGTCGGGGGGAATTGATAAAGAAGCAATGGGATTTACTTACAGCGGCATTCGGACCAACGGTGCTACCTCAGCATGCAGGCG CTCTGGCTCCTCCTTCTACGCTGAGATGAAATGGCTACTGTCCAACACCGACAATGCAGCATTTCCTCAAATGACTAAGAGCTATAAGAAC ACAAGAAAATCCCCCGCCCTCATAGTCTGGGGGATCCACCACTCAGTGTCCACCGCAGAGCAGACGAAATTATATGGATCCGGAAATAA ACTGGTGACAGTGGGGAGTAGTAACTACCAACAGAGTTTTGTTCCCAGCCCAGGCGCCAGACCTCAGGTAAATGGACAGTCCGGAAGGA TCGATTTTCATTGGCTCATGCTAAACCCCAACGATACCGTCACATTCTCATTCAACGGAGCATTTATAGCTCCCGACCGGGCTTCATTCCT TCGGGGTAAATCAATGGGCATTCAGTCTGGGGTACAAGTGGACGCCAATTGTGAGGGCGACTGCTATCACTCTGGCGGAACAATCATAT CTAATCTCCCATTCCAGAACATCGATTCCCGCGCAGTCGGCAAGTGTCCAAGGTACGTGAAACAGCGCTCCCTGTTGTTGGCAACTGGAA TGAAAAATGTACCAGAGATTCCTAAGGGACGTGGACTGTTCGGAGCTATTGCAGGGTTCATCGAAAATGGGTGGGAGGGCCTAATCGAC GGTTGGTATGGGTTCAGACATCAGAACGCCCAGGGGGAGGGTACTGCCGCCGATTACAAGTCCACCCAGTCCGCAATTGACCAGATCA CGGGTAAATTAAACAGACTCATTGAAAAAACTAATCAGCAGTTCGAGCTTATTGATAATGAATTTAATGAAGTGGAAAAGCAGATAGGGAA TGTGATTAACTGGACCCGCGACTCTATCACAGAAGTCTGGTCATACAATGCTGAACTGCTTGTCGCCATGGAAAATCAGCATACCATAGA CCTTGCAGATTCAGAAATGGACAAGCTGTACGAAAGAGTCAAAAGACAGCTGAGGGAGAATGCCGAGGAGGATGGTACAGGGTGCTTCG AAATCTTCCACAAGTGCGACGATGACTGTATGGCGAGCATCCGCAATAATACGTATGACCATTCTAAATACAGAGAAGAGGCGATGCAGA ATCGGATCCAAATCGACCCGGTAAAGCTCAGTTCGGGCTACAAAGACGTAATACTATGGTTCTCCTTTGGAGCTTCGTGCTTCATTCTCCT AGCTATCGTCATGGGGTTAGTCTTCATCTGCGTGAAGAATGGTAATATGCGCTGTACAATTTGTATA
2879	ATGAACACACAGATCCTAGTGTTTGCCCTCATCGCCATTATTCCAACAAACGCTGATAAGATTTGCCTAGGACATCACGCAGTGTCCAACG GAACTAAGGTGAACACCTTAACTGAGAGGGGAGTAGAGGTGGTGAATGCCACAGAGACTGTTGAAAGAACCAATATTCCCCGTATCTGCT CAAAGGGGAAGAGAACGGTAGACCTGGGGCAGTGTGGCTTGCTGGGCACCATCACCGGCCCACCCCAGTGCGATCAGTTTTTGGAATT CTCTGCCGATTTAATAATAGAACGCAGAGAAGGTTCCGACGTGTGCTACCCTGGAAAATTCGTCAATGAGGAAGCACTGAGGCAGATCTT ACGCGAGTCCGGAGGCATCGACAAAGAGGCAATGGGATTCACTTACTCCGGAATCCGGACTAACGGGGCGACGAGCGCCTGCAGGCGT AGCGGCTCCTCTTTTTACGCGGAAATGAAGTGGCTTCTATCCAACACTGACAATGCCGCTTTTCCACAAATGACCAAGTCTTATAAAAACA CACGCAAGTCGCCCGCCTTAATAGTGTGGGGCATTCATCATTCCGTGTCAACAGCAGAGCAGACTAAGTTGTATGGCAGTGGTTCTAAGT TGGTGACAGTAGGATCCTCCAACTACCAGCAAAGTTTCGTGCCTAGCCCAGGCGCACGTCCCCAGGTGAACGGCCTTTCAGGGAGGATC GATTTCCATTGGCTGATGCTCAACCCCAACGACACTGTTACTTTCTCCTTCAACGGAGCATTTATCGCACCAGATCGGGCCTCCTTCTTGC GGGGAAAATCTATGGGAATCCAGAGCGGAGTGCAGGTGGATGCCAATTGTGAAGGGGATTGCTACCACTCTGGGGGAACAATTATCTCC AATTTGCCCTTCCAGAATATCGATTCCCGAGCTGTCGGAAAATGTCCGAGGTATGTCAAGCAGCGTAGTCTGCTCCTTGCCACGGGCATG AAGAACGTCCCAGAGATTCCCAAGGGCCGGGGCTTATTCGGGGCTATCGCCGGGTTCATCGAGAACGGATGGGAGGGGCTAATAGACG GCTGGTATGGCTTCAGGCATCAGAATGCACAAGGTGAAGGAACGGCGGCTGATTATAAATCCACCCAGAGTGCCATCGATCAGATTACA GGTAAGCTTAACCGGCTAATCGAAAAGACGAATCAGCAGTTTGAACTCATCGATAACGAGTTCAATGAGGTGGAGAAGCAGATCGGAAAT GTCATCAACTGGACTCGAGACTCGATCACTGAGGTGTGGTCCTACAATGCTGAGCTTCTGGTCGCTATGGAAAATCAACACACCATCGAT CTGGCAGACTCAGAAATGGACAAGTTGTACGAGCGGGTGAAGAGACAACTAAGGGAAAACGCCGAGGAAGATGGCACTGGATGCTTTG AAATCTTTCACAAGTGCGACGATGACTGCATGGCTTCAATCAGAAATAACACCTACCATCACAGCAAGTACAGACAAGAAGCAATGCAGA ACCGTATTCAAATCGACCCCGTCAAGCTGTCCAGCGGATACAAGGATGTGATCCTGTGGTTCTCCTTCGGGGCAAGCTGTTTCATCCTGC TGGCTATAGTGATGGGCCTGGTCTTCATTTGTGTCAAGTCCAGGAATATGCGGTGTACTATCTGTATT
2880	ATGAACACCCAGATACTGGTCTTTGCTCTCATAGCGATCATACCAACCAACGCTGACAAAATCTGTCTGGGGCACCACGCGGTGTCTAAC GGCACTAAGGTAAACACCCTCACTGAACGCGGTGTGGAGGTGGTGAACGCCACCGAGACCGTTGAGCGTACAAATATACCCAGGATCTG TTCCAAGGGTAAAAGGACCGTCGACCTTGGCCAGTGTGGCCTGCTGGGCACCATTACGGGCCCTCCCCAATGCGATCAATTTCTTGAGT TCAGCGCCGATTTGATCATAGAGCGACGTGAGGGGTCAGACGTGTGTTACCCTGGGAAATTTGTGAAGGAAGAAGCACTGCGACAGATC CTTCGAGAGTCTGGAGGGATTGATAAGGAAGCTATGGGCTTCACCTATAGCGGAATCAGGACTAACGGCGCTACTAGTGCGTGCAGACG ATCTGGAAGCAGTTTTTACGCCGAAATGAAGTGGCTGCTGTCTAATACTGATAATGCCGCCTTCCCACAAATGACCAAGTCCTACAAGAA CACACGGAAGTCCCCAGCTCTGATCGTGTGGGGCATCCATCACTCGGTCTCCACAGCCGAGCAAACAAAGCTTTACGGGTCAGGAAACA AATTGGTAACCGTGGGCTCTAGTAACTATCAACAGAGCTTCGTCCCGTCCCCCGGTGCACGCCCACAGGTGAACGGTTTAAGTGGGCGC ATCGACTTCCATTGGCTCATGCTGAACCCTAATGACACAGTGACCTTTTCCTTTAACGGGGCATTCATCGCCCCTGATAGGGCATCCTTCC TCCGGGGCAAGAGTATGGGGATACAATCGGGGGTGCAAGTCGACGCAAATTGCGAGGGTGATTGTTACCACTCTGGGGGTACCATCATT TCCAACCTGCCCTTTCAGAACATAGATTCGAGGGCTGTCGGGAAGTGCCCTCGATATGTGAAGCAGCGGAGTCTCCTGCTGGCAACAGG CATGAAGAACGTCCCCGAGATCCCCAAAGGACGAGGACTCTTCGGCGCAATCGCCGGTTTTATAGAGAACGGATGGGAAGGCCTGATC GACGGTTGGTATGGCTTTCGTCACCAGAACGCGCAGGGTGAGGGCACTGCCGCCGACTATAAGAGCACACAGTCCGCCATCGACCAAA TTACTGGAAAGCTTAATAGACTAATCGAAAAAACAAACCAACAGTTTGAGCTGATTGACAATGAGTTTAATGAGGTGGAGAAGCAGATAGG TAATGTGATAAATTGGACCCGGGATTCTATTACCGAAGTGTGGTCATACAATGCGGAACTGCTGGTAGCAATGGAAAATCAGCACACAAT CGATCTGGCCGATAGTGAGATGGACAAGTTGTACGAGAGGGTTAACAGGCAGCTTCGGGAAAATGCAGAAGAGGATCGTACCGGGTGT TTTGAAATCTTCCACAAGTGCGATGACGACTGCATGGCTTCTATCAGAAACAATACATACGACCACAGCAAATATCGGGAGGAAGCCATG CAGAACCGAATCCAGATCGATCCCGTCAAGCTTAGCTCGGGCTATAAAGATGTCATTCTGTGGTTTTCCTTTGGAGCCTCCTGTTTCATCC TCCTAGCCATTGTGATGGGTCTTGTGTTTATCTGCGTGAAGAACGGCAATATGAGGTGCACTATCTGCATT

2881	ATGAATACCCAAATTCTTGTGTTCGCCCTGATCGCTATTATCCCGACGAACGCCGATAAGATCTGCCTAGGACACCACGCTGTGAGTAAT GGAACCAAGGTGAACACACTGACTGAGAGGGGCGTGGAGGTGGTGAACGCGACAGAGACCGTGGAACGAACAAATATTCCGAGGATCT GTTCAAAGGGGAAGAGAACCGTGGATCTGGGCCAGTGCGGACTGCTAGGCACTATTACCGGCCCACCACAGTGTGACCAGTTCCTGGA GTTTAGTGCCGACCTGATTATTGAAAGGCGCGAGGGAAGTGACGTTTGTTACCCCGGGAAATTCGTAAATGAGGAGGCCCTCAGACAAA TACTAAGAGAGTCCGGCGGAATTGACAAAGAGCCGATGGGCTTTACATACAGCGGTATAAGAACGAACGGGGCGACCTCCGCATGTAG GCGTAGTGGCTCAAGTTTCTATGCCGAGATGAAGTGGCTGCTGAGCAATACAGACAATGCCGCCTTTCCCCAGATGACCAAGTCTTACAA GAATACGAGAAAAAGTCCAGCACTCATTGTGTGGGGTATCCATCATTCTGTGAGCACAGCAGAACAAACAAAGCTGTACGGCTCAGGAAA CAAGCTCGTGACTGTGGGCAGCTCTAACTACCAGCAATCATTCGTCCCCTCCCCGGGAGCCAGACCACAGGTTAACGGACTGAGTGGGC GAATTGATTTCCACTGGCTTATGTTAAACCCCAATGATACTGTGACATTTTCTTTCAACGGCGCGTTTATCGCTCCAGATCGTGCCAGTTTT CTGAGGGGGAAGTCCATGGGCATCCAGTCCGGAGTGCAGGTTGACGCCAATTGTGAAGGGGACTGTTACCATTCTGGTGGGACTATTAT TTCGAACCTACCCTTCCAGAATATTGACAGCCGGGCCGTTGGTAAATGCCCTCGGTACGTTAAACAAAGAAGCCTCCTGCTGGCGACCG GAATGAAGAATGTGCCTGAAATTCCGAAAGGACGCGGACTGTTCGGCGCTATCGCAGGGTTTATAGAAAACGGTTGGGAGGGCTTAATT AACGGTTGGTATGGCTTTCGGCACCAGAACGCCCAGGGCGAGGGGACCGCCGCCGACTATAAAAGCACACAGTCTGCGATTGATCAGA TTACCGGGAAGCTCAACAGGCTCATAGAGAAAACAAACCAACAGTTTGAACTGATTGACAATGAATTTAACGAGGTAGAAAAACAGATTG GCAATGTTATCAACTGGACCAGGGATTCAATTACAGAGGTGTGGTCTTATAATGCCGAGCTATTGGTGGCCATGGAAAACCAACATACAA TCGACCTCGCCGATTCAGAGATGGACAAGCTGTATGAGAGGGTCAAAAGACAGTTGCGCGAGAATGCCGAGGAGGACGGGACTGGCTG CTTCGAGATCTTTCACAAATGCGATGACGACTGTATGGCTAGTATCCGCAACAACACGTATGATCACTCTAAGTACCGCGAGGAAGCCAT GCAGAATCGGATCCAAATTGATCCAGTTAAACTGTCTAGCGGCTACAAGCATGTGATTCTGTGGTTTTCTTTCGGAGCATCCTGCTTCATC TTGTTAGCCATTGTGATGGGCCTGGTTTTTATCTGCGTGAAGAACGGCAATATGCGGTGTACTATCTGCATC
2882	ATGAATACGCAGATTCTGGTGTTTGCCTTAATAGCTATTATCCCCACTAATGCCGACAAGATCTGTTTAGGGCACCACGCTGTATCAAATG GCACCAAAGTCAATACGCTCACCGAACGGGGGGTGGAAGTTGTGAACGCGACTGAAACTGTCGAAAGAACCAATATTCCGCGAATTTGC AGTAAGGGAAAGAAGACTGTGGACCTGGGCCAATGTGGCTTGTTGGGCACCATTACCGGTCCACCGCAGTGCGATCAATTTCTTGAATT CTCTGCAGACCTGATAATCGAACGCCGCGAGGGCTCTGACGTGTGTTATCCTGGGAAATTTGTTAATGAGGAAGCCCTCAGGCAGATAC TCAGGGAATCTGGCGGTATTGATAAGGAGGCCATGGGGTTCACGTACTCTGGTATACGCACCAATGGTGCCACATCCGCATGCAGGAGA AGCGGCTCCTCGTTCTATGCAGAAATGAAGTGGTTGCTGTCGAACACAGACAACGCCGCCTTTCCACAGATGACAAAGAGTTACAAGAAC ACAAGGAAGTCACCTGCCCTGATCGTTTGGGGTATTCATCATTCTGTCTCCACAGCTGAGCAGACCAAACTGTACGGGTCAGGAAACAAG CTGGTGACTGTGGGTTCCTCGAATTATCAGCAGAGCTTTGTTCCGTCCCCAGGTGCGAGGCCCCAGGTGAACGGCCAATCCGGGCGCA TCGACTTTCACTGGCTGATGCTTAACCCAAACGATACTGTGACCTTCTCCTTTAACGGAGCGTTTATCGCTCCAGACAGGGCCTCGTTTCT GAGAGGGAAAAGCATGGGAATCCAGTCAGGGGTGCAGGTGGATGCTAATTGTGAAGGTGACTGTTACCACTCAGGAGGCACAATCATTA GCAACCTGCCCTTTCAGAACATTGACTCCAGGGCAGTGGGCAAATGTCCACGGTATGTGAAGCAGAGAAGCCTCCTCCTGGCCACAGGC ATGAAGAACGTGCCAGAGATTCCCAAGGGTAGGGGCCTGTTCGGTGCTATAGCCGGATTTATTGAGAATGGATGGGAAGGGCTGATCGA CGGTTGGTACGGCTTTAGACACCAGAACGCTCAGGGAGAGGGAACTGCCGCAGACTATAAAAGTACTCAAAGCGCGATCGACCAGATAA CAGGTAAGCTCAACCGTCTCATTGAGAAAACAAACCAGCAGTTCGAACTCATCGATAACGAGTTTAACGAAGTCGAGAAACAAATTGGAA ACGTCATTAACTGGACTAGAGATTCAATCACTGAGGTGTGGAGCTACAACGCTGAGCTGCTTGTGGCTATGGAGAATCAGCATACCATAG ATCTGGCTGACTCTGAAATGGATAAACTCTATGAACGAGTCAAGAGACAGCTCAGGGAGAATGCTGAAGAGGATGGCACAGGATGTTTC GAAATCTTCCACAAGTGTGACGACGACTGCATGGCAAGCATTAGGAATAACACATATGATCATTCTAAATATAGGGAACAGGCAATGCAG AATCGGATCCAAATTGATCCAGTCAAGTTATCTTCCGGGTATAAAGACGTGATCCTTTGGTTCAGCTTTGGCGCTTCATGTTTCATCTTGC TTGCCATCGTCATGGGACTCGTCTTTATCTGCGTCAAAAATGGTAACATGAGATGCACAATTTGTATC
2883	ATGAATACCCAGATTTTAGTGTTTGCCCTGATTGCCATAATCCCAACTAACGCCGATAAGATTTGCTTGGGCCACCACGCTGTGAGCAAC GGCACCAAAGTCAACACCCTGACCGAACGCGGCGTGGAGGTCGTCAATGCCACCGAAACCGTGGAGCGAACCAACATTCCACGCATCT GTTCCAAGGGAAAAAGAACGGTTGATCTCGCCCAGTGTGGCCTGCTCGGCACAATCACCGGGCCGCCGCAGTGCGACCAGTTCCTGGA GTTTAGCGCCGACCTAATCATCGAACGCCGGGAGGGTTCTGATGTGTGCTACCCTGGCAAATTTGTGAATGAGGAGGCGCTTAGGCAGA TTCTCAGGGAGTCCGGTGGAATTGACAAGGAAGCCATGGGTTTCACATACAGCGGTATTAGGACTAATGGCGCAACATCAGCTTGTCGA AGATCTGGTTCTAGCTTTTACGCAGAGATGAAATGGCTTCTATCCAATACTGATAATGCAGCTTTTCCCCAAATTACAAAGTCATATAAAAA CACCCGCAAATCGCCCGCTCTCATAGTGTGGGGAATTCACCATTCCGTTAGTACCGCTGAACAGACCAAATTGTATGGGAGCGGTAACAA GCTGGTCACCGTTGGGTCCTCCAATTATCAGCAGTCGTTCGTGCCGAGTCCTGGAGCCCGCCCCCAGGTCAACGGCCTTTCTGGCCGG ATTGACTTTCATTGGCTGATGTTGAACCCTAACGACACAGTTACATTCTCATTTAATGGAGCCTTCATCGCTCCGGATCGAGCCAGCTTCC TCCGTGGTAAGAGTATGGGGATCCAAAGCGGTGTTCAGGTCGATGCCAACTGCGAGGGCGATTGTTATCATTCAGGGGGTACTATAATC AGCAACCTGCCCTTTCAAAACATAGACAGCAGGGCAGTCGGAAAATGTCCTCGATATGTCAAACAACGGTCTCTGTTGCTGGCTACAGGG ATGAAGAACGTGCCGGAGATCCCCAAAGGACGTGGATTATTTGGGGCCATAGCCGGGTTTATTGAGAATGGATGGGAGGGCCTGATTGA TGGATGGTACGGTTTTCGCCATCAGAATGCACAGGGAGAAGGAACAGCTGCCGACTACAAGTCAACGCAGAGTGCTATTGACCAGATAA CTGGCAAGTTAAACCGGCTGATCGAAAAGACTAATCAACAATTTGAGTTGATCGACAATGAGTTTAATGAAGTTGAGAAACAGATCGGCAA CGTCATCAACTGGACGCGCGATAGCATTACAGAAGTCTGGTCTTATAACGCCGAGTTGCTTGTAGCAATGGAGAACCAGCATACCATTGA TCTCGCGGACAGTGAAATGGACAAGCTCTACGAGCGCGTAAAGCGACACCTAAGAGAAAATGCAGAAGAAGATGGGACGGGATGTTTCG AAATCTTCCACAAATGCGATGACGATTGCATGGCCAGCATACGGAATAATACCTACGATCATTCCAAGTATCGGGAAGAGGCCATGCAGA ACCGCATTCAGATTGATCCCGTGAAACTTTCAAGTGGCTACAAGGACGTCATCCTGTGGTTCTCTTTCGGGGCAAGTTGTTTTATTCTGCT CGCTATTGTCATGGGTCTGGTCTTTATCTGTGTCAAAAATGGCAACATGCGGTGCACAATCTGTATT

2884	ATGAACACACAGATTCTAGTGTTTGCACTGATTGCTATCATTCCTACAAATGCAGACAAGATTTGTCTGGGCCATCACGCCGTCTCGAATG GTACAAAAGTGAATACCCTGACTGAAAGAGGGGTCGAAGTGGTTAACGCAACAGAGACCGTAGAGAGAACGAATATTCCAAGAATCTGCT CTAAGGGCAAAAAAACAGTGGATCTGGGTCAGTGTGGACTGTTAGGGACAATCACAGGTCCACCACAATGCGACCAGTTCCTTGAATTCT CGGCGGATCTTATCATCGAGCGCCGAGAAGGTTCCGACGTGTGTTACCCAGGAAAATTCGTCAATGAGGAGGCTTTGAGACAGATCTTG CGCGAAACCGGTGGCATCGACAAAGAGGCTATGGGATTCACTTACTCTGGAATACGAACTAACGGTGCCACGTCCGCTTGCCGCAGGA GTGGATCTAGCTTCTATGCAGAGATGAAGTGGCTATTGAGCAACACTGATAATGCTGCCTTCCCTCAGATGACAAAGAGCTATAAAAACA CTAGAAAGAGCCCAGCATTGATCGTTTGGGGCATTCACCACAGTGTATCCACAGCAGAGCAGACCAAACTGTATGGGAGTGGAAACAAG CTTGTTACTGTAGGGTCCAGCAACTATCAACAGAGCTTTGTTCCGTCACCAGGTGCTCGGCCCCAGGTTAATGGGCAAAGTGGGCGGAT CGACTTCCACTGGCTGATGCTCAATCCTAATGACACGGTCACATTTTCCTTTAACGGGGCCTTCATTGCTCCGGATCGCGCCAGCTTTCT GCGTGGAAAGAGTATGGGTATTCAAAGTGGCGTTCAGGTCGACGCCAACTGCGAGGGCGACTGCTATCATAGCGGTGGAACAATCATCT CTAACCTGCCTTTCCAGAACATCGATAGCAGAGCCGTTGGCAAGTGTCCCAGGTACGTGAAGCAGCGATCACTGTTGTTGGCTACGGGC ATGAAAAACGTCCCAGAGATTCCAAAAGGGAGGGGACTATTCGGCGCCATTGCTGGGTTCATTGAAAATGGCTGGGAGGGTCTGATTGA TGGATGGTACGGGTTTAGGCACCAAAACGCCCAAGGCGAAGGGACAGCAGCTGACTATAAATCCACTCAATCTGCTATCGACCAGATCA CTGGAAAACTCAACAGACTTATTGAGAAGACTAATCAGCAGTTTGAGCTCATCGACAACGAGTTTAATGAAGTGGAGAAGCAGATTGGGA ACGTTATCAACTGGACGAGGGACTCAATCACGGAAGTATGGTCCTACAATGCCGAATTGCTGGTGGCCATGGAGAACCAGCACACCATT GACCTGGCGGACTCCGAAATGGATAAACTCTACGAACGAGTGAAGAGGCAACTCCGAGAAAATGCTGAGGAAGATGGGACCGGTTGTTT CGAGATCTTTCATAAGTGTGACGATGACTGCATGGCTTCTATTAGAAACAATACTTACGATCACTCAAAATATAGAGAAGAGGCAATGCAG AACCGCATTCAGATCCACCCGGTGAAGCTGAGCAGTGGTTATAAGGATGTCATTTTATGGTTTAGTTTTGGAGCATCTTGTTTTATTTTGC TCGCTATCGTAATGGGCCTCGTGTTTATCTGCGTGAAAAATGGTAACATGCGGTGCACGATATGCATT
2885	ATGAATACGCAGATTTTGGTTTTCGCCTTGATTGCCATTATTCCTACAAACGCCGACAAGATCTGCCTGGGACATCACGCCGTGAGTAATG GAACCAAGGTGAATACACTGACCGAGAGGGGCGTAGAAGTTGTAAATGCTACTGAGACAGTGGAGCGAACAAATATTCCCAGAATTTGCT CCAAGGGCAAGAAGACTGTCGACCTTGGCCAATGTGGGCTGCTCGGCACGATCACGGGGCCACCCCAATGTGATCAGTTTCTGGAGTTT TCTGCCGACCTCATTATAGAACGCCGGGAGGGGAGCGACGTATGCTACCCCGGTAAGTTCGTAAACGAAGAAGCCCTCAGACAGATTTT GAGGGAGTCCGGAGGCATCGATAAGGAGGCAATGGGCTTCACCTACAGCGGCATCAGAACAAACGGTGCCACAAGTGCATGTAGGCGC TCTGGCTCATCATTCTATGCAGAGATGAAGTGGCTGCTTTCCAATACTGATAATGCGGCCTTTCCTCAGATGACTAAATCATATAAGAACA CCAGAAAATCTCCCGCCTTGATCGTCTGGGGGATCCATCACAGCGTCTCTACCGCTGAGCAGACCAAACTCTACGGTTCAGGGAACAAA TTAGTGACCGTTGGCAGCTCCAACTACCAGCAATCATTTGTGCCTTCCCCAGGTGCCAGGCCGCAAGTAAACGGTCAGTCAGGACGAAT TGATTTCCACTGGCTAATGCTGAACCCGAATGACACAGTCACCTTCAGCTTTAACGGGGCTTTCATCGCACCCGACCGCGCCAGTTTCTT ACGGGGAAAATCAATGGGAATACAATCCGGCGTTCAAGTGGACGCGAATTGCGAGGGAGATTGTTACCATAGTGGTGGTACAATTATCTC CAATCTTCCCTTTCAGAATATCGACAGTAGGGCAGTGGGAAAATGTCCCCGATACGTCAAACAGCGGAGCCTGCTCCTCGCCACCGGGA TGAAGAACGTCCCCGAGATTCCAAAAGGTAGGGGTTTGTTCGGAGCTATCGCAGGGTTCATCGAAAACGGCTGGGAAGGCCTGATAGAC GGTTGGTATGGATTCCGACACCAAAACGCTCAAGGCGAGGGCACTGCAGCAGATTACAAATCTACACAAAGCGCTATCGACCAGATTAC CGGAAAACTTAACCGTCTAATCGAAAAGACCAATCAGCAGTTCGAGCTCATCGATAACGAATTCAATGAGGTGGAGAAACAGATCGGAAA CGTAATCAATTGGACACGAGATTCCATCACTGAGGTTTGGTCTTATAACGCCGAACTCTTAGTGGCTATGGAGAATCAGCACACCATCGA CCTGGCTGATTCCGAGATGGACAAGCTGTACGAAAGAGTGAAGAGACAGCTTCGCGAAAACGCAGAGGAGGATGGGACCGGATGTTTC GAAATCTTCCACAAATGTGATGATGATTGCATGCCCAGCATCCCGAACAACACATATGATCATTCGAAATATCGCCGAGGAGGCTATGCAA AATCGGATCCAGATCGATCCCGTCAAGCTAAGCTCTGGGTACAAGGACGTGATTTTATGGTTTTCCTTCGGCGCTTCATGTTTCATCCTTC TGGCCATCGTGATGGGCCTAGTGTTTATCTGCGTCAAGAATGGAAACATGAGATGTACGATCTGCATC
2886	ATGAACACTCAGATTCTGGTATTTGCGCTAATCGCAATCATTCCCACCAACGCAGACAAGATTTGTCTGGGACATCATGCCGTGAGCAAC GGCACAAAAGTGAACACTCTGACTGAAAGGGGAGTGGAGGTTGTAAATGCGACCGAAACTGTGGAAAGAACCAATATACCCCGAATCTG TAGCAAGGGTAAAAAGACCGTAGACTTAGCACAGTGTGGTCTCTTGGGGACAATCACAGGCCCCCCTCAATGTGACCAGTTCTTAGAGTT TAGCGCAGATTTGATTATCGAGCGAAGGGAGGGAAGTGACGTTTGTTACCCTGGGAAGTTCGTGAATGAAGAAGCTCTCCGTCAGATCTT GAGGGAGTCCGGAGGCATAGACAAAGAGGCCATGGGGTTCACTTACTCCGGGATTCGCACAAATGGAGCTACAAGTGCCTGTAGAAGG TCAGGCAGTTCTTTCTATGCTGAGATGAAATGGCTCTTATCAAACACAGACAACGCTGCCTTCCCACAGATGACAAAATCCTACAAAAACA CTCGCAAATCCCCAGCCCTTATCGTGTGGGGGATCCATCACTCCGTGTCCACTGCGGAACAGACTAAGCTGTATGGATCCGGGAACAAA CTGGTAACTGTAGGGAGTAGCAACTACCAGCAGTCCTTCGTCCCATCTCCTGGGGCACGCCCCCAGGTAAACGGACAGTCTGGAAGAAT TGATTTCCACTGGTTGATGCTGAACCCAAATGATACTGTGACCTTCTCTTTCAATGGCGCTTTCATCGCGCCCGATCGCGCTAGCTTCCTA AGGGGAAAATCCATGGGGATTCAGTCTGGCGTCCAGGTCGATGCAAATTGCGAAGGCGACTGTTACCATTCTGGAGGGACCATTATCAG TAACCTGCCATTCCAGAATATTGACTCGAGGGCAGTGGGTAAGTGCCCCAGGTACGTGAAACAAAGATCACTGCTTCTGGCTACAGGAAT GAAGAACGTACCTGAGATCCCTAAAGGGCGGGGATTATTCGGAGCCATCGCAGGCTTCATAGAGAACGGGTGGGAGGGCCTGATTGAT GGGTGGTACGGATTCAGGCACCAAAATGCCCAGGGAGAAGGGACCGCTGCGGATTACAAGAGTACCCAGTCCGCTATAGACCAGATCA CAGGCAAACTTAATAGGCTGATTGAAAAGACCAATCAGCAGTTTGAACTCATCGATAACGAATTTAACGAGGTGGAAAAGCAGATTGGGA ACGTGATTAACTGGACAAGGGATTCTATTACCGAGGTGTGGAGCTACAATGCCGAGCTGTTGGTTGCGATGGAGAATCAGCACACTATC GATCTAGCAGACTCCGAAATGGACAAGTTGTATGAAAGAGTGAAGCGCCAATTAAGGGAAAACGCAGAGGAGGATGGGACCGGATGTTT TGAGATCTTCCATAAATGCGATGACGACTGCATGGCCAGCATTAGAAATAATACCTACGACCATAGTAAGTATAGGGAGGAGGCGATGCA AAACAGAATTCAAATAGACCCAGTGAAACTGAGTTCTGGGTACAAAGACGTTATTCTCTGGTTTTCATTCGGCGCCAGCTGCTTTATACTG CTGGCTATCGTCATGGGCTTGGTCTTCATCTGCGTGAAAAACGGGAATATGCGATGCACAATTTGTATT

2887	ATGAATACCCAGATCCTGGTGTTCGCTCTCATCGCCATCATACCAACAAACGCTGACAAGATCTGCCTAGGTCACCACGCTGTGTCTAAT GGAACCAAAGTGAATACACTGACCGAGCGCGGTGTGGAGGTGGTGAACGCTACAGAGACCGTGGAACGCACTAATATCCCTCGAATATG TTCTAAGGGCAAGAAAACTGTAGATCTGGGGCAGTGCGGCCTTCTGGGAACTATCACGGGCCCTCCACAGTGCGATCAATTTTTGGAGT TTTCTGCCGACCTCATCATCGAACGCCGTGAGGGATCGGACGTGTGTTATCCAGGAAAGTTTGTGAATGAGGAAGCGCTCCGGCAGATC TTAAGAGAAAGCGGTGGTATTGACAAACAAGCAATGGGCTTCACCTATTCAGGAATTAGGACCAACGGAGCAACCAGTGCGTGTAGACG AAGCGGGTCTTCATTCTACGCCGAAATGAAGTGGCTGCTTTCTAACACCGATAACGCAGCATTCCCACAAATGACCAAGAGCTACAAGAA TACAAGAAAATCCCCTGCATTAATTGTATGGGGAATCCATCACTCCGTCTCAACAGCAGAACAGACAAAACTTTATGGCAGTGGCAACAAA CTCGTCACCGTTGGTAGCTCCAACTACCAGCAGTCTTTCGTGCCAAGTCCCGGGGCCCGACCGCAAGTGAACGGGCAGTCTGGCAGAA TCGACTTTCACTGGCTCATGCTGAATCCAAACGACACAGTTACCTTCAGCTTCAATGGGGCATTCATTGCCCCAGACAGAGCCAGCTTTC TTCGAGGCAAGAGCATGGGCATCCAGAGCGGGGTACAGGTCGACGCAAACTGTGAAGGAGATTGCTACCACTCCGGCGGGACGATCAT TTCGAATCTGCCATTCCAGAATATCGACTCTAGGGCTGTGGGGAAATGTCCTAGGTATGTGAAACAGCGGAGCCTCCTTCTGGCAACAG GAATGAAAAACGTACCTGAGATCCCTAAGGGGCGAGGATTGTTTGGTGCCATCGCGGGATTCATTGAAAATGGCTGGGAGGGATTGATT GACGGCTGGTACGGATTCCGGCATCAAAACGCGCAAGGGGAGGGTACGGCGGCAGATTACAAAAGCACTCAGTCCGCGATCGACCAGA TCACCGGCAAACTCAATAGACTTATCGAGAAGACTAACCAACAGTTCGAGCTCATCGACAATGAGTTTAATGAAGTAGAGAAACAGATCG GCAATGTCATAAACTGGACCAGGGACAGCATCACCGAAGTGTGGTCCTATAACGCCGAGCTCCTGGTGGCCATGGAGAATCAGCACACT ATAGACTTAGCTGACTCTGAGATGGACAAGCTCTATGAGAGAGTTAAACGGCAGCTCAGAGAAAACGCAGAAGAAGATGGCACCGGGTG CTTCGAGATCTTTCATAAATGTGATGACGACTGCATGGCCTCGATACGTAACAATACATATGATCATAGTAAGTATCGGGAGGAAGCAATG CAGAACAGGATTCAGATCGACCCCGTTAAATTGTCCAGCGGCTATAAGGACGTAATTCTGTGGTTCTCGTTTGGCGCCTCCTGTTTCATTT TGTTGGCCATTGTGATGGGTCTCGTCTTCATTTGCGTAAAAAATGGCAATATGCGGTGCACTATTTGTATC
2888	ATGAATACTCAGATCCTAGTTTTTGCTCTGATTGCCATAATTCCGACGAATGCGGACAAGATTTGCCTGGGACACCACGCTGTCAGTAAC GGTACCAAAGTAAATACCCTGACGGAGCGAGGTGTCGAGGTGGTGAATGCCACCGAAACTGTGGAGAGAACCAACATCCCTCGGATCTG TAGCAAGGGCAAGAAAACTGTGGACCTCGGTCAGTGTGGGTTGCTCGGGACCATTACAGGCCCCCCACAGTGCGATCAATTTCTGGAGT TCTCTGCCGACTTAATTATTGAAAGACGGGAGGGTAGCGACGTTTGTTATCCCGGCAAATTCGTAAATGAGGAGGCCTTGCGCCAGATTT TGCGCGAGTCCGGCGGCATTGATAAAGAGGCTATGGGATTCACTTATTCTGGCATTAGGACAAACGGCGCGACCAGTGCATGTCGCCGA TCCGGCTCCTCCTTTTACGCTGAGATGAAATGGCTGTTGTCGAACACTGACAACGCTGCATTTCCACAGATGACAAAGTCCTATAAGAAC ACCAGGAAGAGTCCGGCCCTGATCGTGTGGGGAATACATCACTCTGTGTCCACCGCCGAACAGACCAAGCTGTACGGAAGTGGCAATAA GCTGGTTACTGTCGGGAGCAGCAATTACCAGCAGAGTTTTGTGCCGTCACCCGGGGCCCGGCCTCAGGTGAATGGTCAGTCCGGTAGA ATAGACTTCCACTGGCTTATGCTAAACCCAAATGACACCGTTACCTTCTCATTTAACGGTGCCTTTATTGCCCCCGACCGGGCTTCATTCC TGCGCGGGAAATCCATGGGCATCCAATCTGGGGTGCAGGTTGACGCTAATTGTGAGGGAGACTGTTATCATAGCGGAGGCACCATAATC TCCAATCTGCCCTTCCAGAACATTGACAGCAGAGCAGTAGGCAAATGCCCCAGGTATGTCAAGCAACGAAGCCTTCTTCTCGCTACCGGT ATGAAGAATGTGCCCGAAATACCAAAGGGACGGGGGCTCTTTGGCGCGATAGCCGGGTTCATCGAGAATGGATGGGAGGGACTCATTG ACGGTTGGTATGGGTTCCGCCATCAGAACGCGCAGGGGGAAGGCACAGCAGCCGACTATAAGTCCACACAGAGTGCAATTGACCAAATT ACCGGCAAATTAAACAGACTCATTGAGAAGACTAACCAGCAGTTTGAGTTAATTGATAATGAGTTTAACGAAGTGGAGAAACAAATTGGAA ACGTGATCAACTGGACACGAGATAGCATTACGGAGGTATGGTCATATAACGCCGAGCTGCTCGTTGCGATGGAGAATCAACACACAATTG ATTTAGCAGACTCCGAAATGGACAAACTTTACGAACGGGTGAAACGCCAGCTGCGCGAAAACGCTGAAGAAGATGGGACAGGTTGCTTT GAAATTTTTCACAAGTGCGATGATGACTGCATGGCCTCAATCCGTAATAATACATACGATCACAGCAAGTATAGGGAACAAGCCATGCAG AATAGGATTCAAATAGACCCGGTGAAATTATCTTCTGGCTATAAAGATGTTATTCTGTGGTTCTCCTTCGGCGCGTCATGTTTCATTTTATT GGCCATTGTGATGGGCCTGGTTTTCATATGCGTGAAGAACGGAAATATGAGATGCACCATCTGTATT
2889	ATGAACACACAAATCCTGGTGTTTGCCTTAATAGCTATTATCCCAACCAACGCAGACAAGATCTGTCTTGGACACCATGCTGTAAGCAATG GAACCAAAGTGAACACATTGACCGAGCGTGGGGTCGAAGTGGTTAACGCCACCGAGACAGTGGAGAGGACTAATATCCCCAGAATTTGT AGTAAAGGGAAGAGGACAGTTGACTTAGGACAATGCGGACTGTTGCGTACCATTACAGGTCCACCCCAGTGTGATCAGTTTCTGGAATTC AGCGCTGATTTGATCATCGAGCGGAGAGAAGGCAGTGATGTCTGTTACCCTGGGAAATTCGTCAACGAAGAGGCATTGCGCCAGATTTT GAGGGAATCCGGCGGAATCGACAAAGAAGCCATGGGGTTCACATACTCAGGGATCCGGACGAATGGCGCAACGAGCGCTTGTAGGAGG TCCGGGAGTTCTTTCTATGCGGAGATGAAATGGCTATTGAGCAACACAGATAATGCCGCTTTCCCTCAAATGACCAAGTCATACAAAAATA CAAGGAAGAGCCCAGCCCTTATCGTGTGGGGTATCCATCACAGTGTGTCTACCGCAGAACAGACAAAACTCTACGGCAGCGGGAACAAA CTTGTGACCGTCGGCTCTTCAAACTATCAACAGAGTTTCGTTCCGTCTCCAGGCGCAAGGCCACAGGTGAATGGATTAAGTGGGCGGAT CGATTTTCATTGGCTTATGCTCAACCCTAATGATACAGTCACATTCTCTTTTAATGGAGCTTTCATTGCCCCAGACAGGGCGTCCTTCCTG CGAGGCAAATCTATGGGTATACAAAGCGGCGTCCAGGTAGACGCAAATTGTGAGGGTGATTGCTACCACAGCGGGGGAACCATTATCAG CAACTTGCCATTCCAAAACATCGATTCCCGTGCTGTGGGCAAGTGTCCAAGATATGTGAAACAGAGGAGTCTGTTGTTGGCGACGGGAAT GAAAAACGTTCCCGAGATTCCAAAGGGCCGGGGACTTTTTGGGGCCATCGCCGGATTCATAGAGAACGGCTGGGAAGGACTCATTAATG GATGGTATGGGTTCCGGCATCAGAACGCTCAGGGGGAGGGCACCGCTGCTGACTACAAGAGTACCCAGTCTGCCATCGATCAGATTAC CGGTAAGTTGAATAGGCTGATCGAGAAAACTAACCAGCAGTTTGAGCTCATTGATAATGAATTCAACGAGGTAGAAAAACAGATAGGCAA CGTGATTAACTGGACCCGTGACTCCATAACGGAGGTATGGAGTTATAATGCAGAACTCCTCGTGGCTATGGAAAATCAGCACACAATTGA TTTAGCTGACTCAGAAATGGATAAGCTGTACGAGCGGGTCAAACGCCAACTGCGCCGAGAATGCAGAAGAACACGGGACCGGGTGTTTCG AAATCTTTCATAAATGCGACGATGACTGTATGGCCAGCATCCGTAACAACACTTACGACCACAGCAAGTATCGGGAGGAAGCTATGCAAA ATAGGATCCAGATCGACCCTGTCAAATTATCCAGTGGGTACAAAGATGTCATCCTGTGGTTTTCCTTTGGCGCGTCATGCTTTATTTTGTT GGCTATCGTGATGGGCCTCGTTTTTATCTGTGTGAAGAATGGAAACATGCGATGCACCATCTGTATT

2890	ATGAACACTCAGATTCTGGTGTTCGCACTAATCGCAATCATACCTACTAATGCCGACAAAATTTGCCTCGGCCACCATGCCGTGTCAAATG GCACAAAAGTGAATACCCTGACGGAGCGCGGCGTCGAGGTTGTCAATGCCACAGAGACAGTGGAACGGACCAACATTCCACGTATTTGT TCAAAAGGCAAGCGTACCGTTGACCTGGGTCAGTGCGGACTTCTCGGGACAATCACTGGACCACCACAGTGCGATCAGTTCTTGGAGTT TTCAGCAGACCTTATCATCGAGAGGCGGGAAGGTAGCGACGTTTGCTACCCCGGCAAATTTGTTAATGAAGAAGCCTTAAGGCAGATCCT ACGAGAATCAGGGGGCATTGACAAAGAAGCGATGGGATTCACTTACAGTGGAATTAGAACAAACGGTGCCACGTCTGCTTGCAGACGCT CGGGCTCTAGCTTTTATGCTGAAATGAAATGGTTGCTGAGCAATACCGATAATGCTGCCTTCCCACAGATGACAAAAAGTTACAAGAATAC TAGAAAGTCACCCGCACTGATCGTCTGGGGGATTCACCATTCAGTCAGCACCGCCGAGCAGACCAAACTTTATGGTAGCGGTAACAAGT TAGTGACAGTTGGAAGCTCCAATTATCAACAGTCATTTGTGCCATCTCCCGGGGCTCGACCCCAAGTGAACGGGTTAAGTGGTCGGATC GACTTCCACTGGTTGATGCTCAACCCTAACGACACAGTCACATTCTCCTTCAACGGCGCATTCATAGCCCCCGACCGGGCAAGCTTCCTA AGGGGAAAGAGTATGGGAATCCAGTCTGGGGTGCAAGTTGATGCCAACTGCGAAGGGGATTGCTATCATTCCGGCGGGACAATTATAAG CAACCTGCCATTCCAGAACATCGATTCACGGGCAGTCGGGAAGTGCCCCAGGTATGTGAAGCAGCGGTCACTGCTGTTGGCTACTGGCA TGAAAAACGTGCCGGAGATCCCAAAAGGCCGTGGGCTCTTCGGGGCTATCGCCGGGTTTATCGAGAACGGCTGGGAGGGCCTGATTAA TGGGTGGTATGGCTTTAGGCATCAGAACGCGCAAGGAGAGGGAACCGCGGCCGACTATAAGAGCACACAGTCTGCGATCGATCAAATAA CCGGTAAGCTCAATCGGCTGATTGAAAAAACCAACCAGCAGTTTGAGCTAATCGACAATGAATTCAACGAGGTGGAAAAGCAGATTGGTA ACGTCATTAATTGGACAAGAGACAGTATTACTGAGGTGTGGAGTTATAACGCTGAACTGCTGGTTGCCATGGAGAATCAGCACACTATTG ACCTGGCAGATTCTGAGATGGATAAGCTCTATGAGAGAGTGAAACGGCAGCTGAGAGAGAATGCTGAAGAGGATGGAACCGGGTGCTTC GAAATTTTTCATAAGTGTGATGACGATTGTATGGCCTCGATTAGAAATAATACATCACGATCACTCCAAGTACCGAGAGGAGGCCATGCAGA ACAGGATCCAAATTCACCCGGTTAAGCTGAGTAGTGGCTACAAGGATGTTATATTATGGTTTAGCTTCCGAGCCTCCTGCTTCATACTGCT GGCTATTGTGATGGGGTTAGTCTTTATCTGCGTGAAAAATGGGAACATGAGATGTACAATCTGTATA
2891	ATGAACACGCAGATCCTAGTCTTTGCCCTTATTGCCATAATCCCTACCAACGCCGATAAGATCTGCCTCGGTCACCACGCCGTTTCCAAT GGCACAAAGGTGAATACTTTGACAGAACGGGGGGTTGAGGTGGTCAATGCTACCGAAACCGTAGAGCGAACAAATATCCCGAGAATATG TTCCAAAGGCAAGAAGACCGTGGACCTCGGACAGTGTGGATTACTCGGGACAATCACAGGTCCCCCACAGTGCGATCAGTTCTTGGAAT TCTCTGCCGACCTCATTATCGAACCGACAGAAGGATCGGATGTGTGTTATCCTGGCAAGTTCGTCAATGAGGAAGCTCTTAGACAGATCT TACGCGAGAGTGGTGGCATCGACAAGGAAGCAATGGGTTTCACATATAGCGGAATTCGGACTAATGGAGCAACTTCAGCTTGTAGGAGA AGCGGCTCATCTTTTTATGCCGAAATGAAGTGGCTGTTATCTAACACCGATAACGCTGCCTTCCCACAGATGACGAAATCATACAAGAACA CTAGGAAGAGCCCAGCTCTGATTGTGTGGGGTATTCATCATAGTGTCTCCACGGCGGAACAGACTAAGCCTGTATGGGTCCGGCAATAAA CTTGTCACCGTAGGCAGCAGCAACTACCAGCAGAGCTTCGTGCCTAGCCCAGGAGCTAGGCCCCAGGTGAATGGACAGTCTGGGCGAA TAGACTTCCACTGGCTTATGCTCAATCCGAATGACACAGTGACATTCAGCTTCAATGGGGCTTTTATTGCTCCAGACAGAGCCAGCTTCCT GAGAGGCAAATCAATGGGGATCCAAAGCGGAGTACAGGTTGACGCCAACTGTGAAGGCGATTGCTACCACTCAGGGGGCACAATTATCT CTAACCTGCCATTCCAGAATATTGATTCGAGAGCTGTGGGGAAGTGTCCTAGGTATGTAAAACAGCGGAGCCTGCTCCTCGCTACAGGG ATGAAGAATGTCCCGGAAATCCCAAAAGGACGGGGCCTGTTTGGCGCAATTGCCGGCTTCATAGAGAATGGGTGGGAGGGACTGATCG ATGGATGGTACGGGTTTCGCCACCAGAACGCCCAGGGAGAAGGGACAGCCGCTGATTACAAATCCACACAAAGCGCAATCGACCAGAT CACAGGTAAACTGAATAGGCTCATCGAGAAGACCAATCAGCAATTCGAACTTATCGACAATGAGTTTAATGAGGTGGAGAAACAAATCGG AAATGTGATCAATTGGACGCGAGATTCTATTACTGAGGTGTGGTCATACAATGCCGAACTCCTTGTGGCTATGGAAAATCAGCACACCATA GACCTGGCCGACTCCGAGATGGATAAGCTATACGAGCGCGTCAAGCGCCAGCTGAGGGAAAATGCTGAGGAAGACGGAACGGGTTGCT TCGAAATTTTTCACAAGTGCGACCATGACTGCATGGCCTCTATTAGAAATAATACATACGACCACTCTAAGTATAGGGAAGAGGCCATGCA GAATCGCATACAGATCGATCCTGTGAAACTCTCCAGCGGGTATAAGGATGTCATTCTGTGGTTCAGTTTCGGCGCTAGCTGTTTTATACTT TTGGCTATTGTTATGGGTCTAGTGTTTATTTGCGTGAAAAACGGCAATATGCGGTGCACAATATGCATT
2892	ATGAATACCCAGATCCTTGTCTTCGCATTAATAGCAATCATTCCAACCAATGCTGACAAGATCTGTCTGGGTCACCATGCAGTTTCCAATG GCACAAAAGTTAATACACTGACAGAACGCGGGGTGGAAGTAGTCAACGCCACCGAAACCGTAGAGAGGACCAATATCCCCAGGATCTGC TCAAAGGGAAAGCGTACAGTGGACCTGGGACAGTGTGGGCTGCTCGGCACAATTACAGGACCACCACAGTGTGACCAGTTCCTGGAGTT TAGCGCTGACTTAATAATCGAGAGAAGAGAGGGAAGCGACGTCTGTTATCCTGGCAAGTTTGTCAAGGAGGAGGCTCTTAGGCAGATCT TGAGAGAGTCGGGCGGCATAGATAAAGAGGCAATGGGATTCACCTATTCAGGGATTAGGACCAATGGTGCCACAAGTGCCTGCAGAAGA TCAGGCTCTAGTTTTTACGCTGAGATGAAGTGGCTTCTGTCGAACACAGATAACGCCGCGTTCCCTCAGATGACTAAGTCATACAAAAATA CACGCAAAAGTCCTGCCTTAATCGTGTGGGGGATACATCATAGTGTATCCACCGCCGAGCAGACTAAGCTCTACGGTAGCGGTAATAAG CTGGTCACTGTGGGCAGTTCCAACTACCAGCAGAGTTTTGTTCCCAGCCCCGGTGCTCGTCCCCAAGTGAACGGACTTAGCGGTAGGAT TGACTTTCATTGGCTGATGTTGAATCCTAATGACACGGTCACTTTTAGTTTCAACGGCGCTTTCATCGCTCCCGATAGAGCTTCCTTCTTG CGCGGAAAGAGTATGGGGATACAGTCAGGCGTGCAGGTCGACGCGAATTGTGAAGGGGATTGCTACCATAGTGGAGGCACTATTATCTC AAACCTCCCTTTCCAGAACATCGACAGTAGGGCCGTGGGAAAATGCCCTAGGTACGTTAAGCAGAGATCCCTATTGCTGGCCACGGGTA TGAAGAATGTCCCCGAAATTCCAAAAGGAAGAGGACTGTTCGGCGCCATCGCTGGCTTTATCGAGAACGGGTGGGAGGGGCTCATCGA CGGGTGGTACGGGTTCAGACACCAGAATGCCCAAGGGGAGGGCACCGCCGCTGACTACAAATCTACGCAAAGTGCCATTGACCAGATT ACCGGAAAGCTCAACCGGCTGATTGAGAAGACCAACCAGCAATTTGAGCTAATTGACAATGAATTTAACGAGGTCGAGAAGCAAATTGGG AACGTCATAAACTGGACTAGGGATTCAATTACCGAAGTCTGGAGCTATAACGCAGAACTATTGGTCGCTATGGAGAACCAGCACACAATC GATCTCGCTGATTCTGAGATGGATAAGCTGTATGAGAGGGTGAAGAGACAGTTGAGGGAGAATGCTGAGGAGGATGGTACCGGCTGTTT TGAAATCTTTCATAAGTGTGACGATGACTGTATGGCTTCAATCCGGAACAATACGTACGATCACAGTAAATATAGGGAAGAAGCTATGCAG AACAGAATTCAGATTGATCCTGTGAAACTAAGCAGTGGTTATAAAGATGTGATCCTCTGGTTCAGTTTCGGCGCCAGCTGTTTCATCCTGC TCGCCATCGTAATGGGACTGGTCTTCATATGCGTGAAGAATGGAAACATGCGTTGCACCATCTGTATT

2893	ATGAATACACAGATTCTGGTGTTCGCTTTGATAGCGATCATTCCAACAAATGCAGACAAGATCTGCCTCGGTCATCACGCGGTTTCAAATG GCACCAAAGTGAACACTCTCACCGAACGGGGGGTCGAGGTAGTTAACGCGACCGAGACTGTGGAGCGTACTAATATCCCCAGGATCTGT TCAAAGGGAAAGAGAACTGTGGACCTTGGGCAGTGTGGTCTGCTGGGCACCATCACTGGCCCCCCCCAGTGTGACCAGTTCCTGGAGT TCTCAGCCGACCTTATAATTGAGCGACGCGAAGGATCTGATGTATGCTACCCAGGCAAGTTTGTGAAAGAGGAGGCGTTACGCCAGATT CTGAGAGAGTCTGGTGGGATCGACAAAGAAGCGATGGGTTTCACATACAGCGGGATCCGGACCAACGGTGCCACCTCGGCATGTCGTC GTAGCGGCTCATCCTTCTACGCCGAGATGAAATGGCTGCTGTCAAACACCGACAACGCCGCTTTCCCGCAGATGACTAAGAGCTACAAA AACACCAGAAAGTCACCCGCCCTGATTGTGTGGGGAATTCATCACTCTGTGTCTACCGCCGAACAGACCAAATTGTATGGTTCGGGGAAT AAGCTGGTTACGGTAGGATCGTCAAATTACCAACAAAGCTTTGTGCCCTCTCCAGGAGCCAGACCTCAGGTAAACGGCCTCTCAGGAAG GATTGACTTCCACTGGCTGATGCTGAATCCCAATGATACCGTAACATTCTCATTTAATGGCGCGTTCATCGCACCGGATAGGGCATCCTTC TTAAGGGGGAAGAGTATGGGCATCCAGTCCGGGGTTCAAGTGGATGCCAATTGCGAAGGCGATTGTTACCACTCCGGAGGAACGATTAT TAGCAATCTGCCCTTCCAAAACATCGACTCTAGGGCCGTGGGCAAATGCCCTAGATATGTGAAGCAGAGGTCCCTGCTCCTAGCGACTG GCATGAAGAATGTGCCCGAGATACCAAAAGGGAGAGGACTATTCGGTGCCATCGCCGGGTTCATTGAAAACGGATGGGAAGGCCTCATC GATGGGTGGTACGGATTTCGGCACCAGAATGCGCAAGGGGAAGGTACTGCTGCTGATTATAAGAGCACTCAGAGCGCTATTGATCAGAT TACCGGTAAATTGAACCGTCTCATTGAAAAGACGAACCAGCAGTTCGAGCTGATCGATAATGAGTTTAATGAAGTGGAAAAGCAAATCGG GAACGTGATCAACTGGACAAGAGACTCAATCACTGAAGTCTGGTCCTACAATGCCGAGCTTTTGGTTGCCATGGAGAACCAGCATACGAT CGACCTTGCCGATAGCGAGATGGACAAACTATACGAGCGAGTCAAGAGGCAACTCAGAGAAAATGCAGAAGAGGATGGAACCGGCTGTT TCGAGATCTTTCATAAGTGCGACGACGATTGCATGGCTTCCATTAGAAATAACACCTACGACCACTCTAAGTATAGAGAGGAGGCTATGC AAAATCGCATCCAGATTGATCCGGTGAAACTCTCCTCGGGCTACAAAGATGTGATCCTGTGGTTCTCCTTTGGCGCAAGCTGTTTCATTTT ATTAGCTATTGTGATGGGGCTCGTCTTTATCTGCGTGAAGAATGGCAACATGCGGTGTACCATCTGTATA
2894	ATGAACACTCAGATACTCGTCTTTGCGCTCATCGCTATTATTCCTACCAACGCGGATAAAATTTGCCTCGGGCACCACGCAGTGTCCAAC GGAACTAAAGTGAACACGCTGACTGAACGTGGTGTGGAAGTGGTGAACGCAACAGAAACAGTGGAAAGGACTAATATTCCGCGTATATG CAGTAAGGGTAAGAAGACGGTCGACCTGGGGCAGTGCGGGCTTCTCGGTACCATCACGGGTCCTCCTCAGTGTGATCAGTTCCTTGAGT TTTCAGCCGACCTCATCATTGAACGTCGTGAGGGAAGTGATGTTTGTTATCCCGGGAAGTTCGTGAACGAAGAGGCATTGAGACAGATTC TGCGGGAGTCTGGGGGAATCGATAAGGAGGCCATGGGTTTCACATACTCCGGTATCAGGACTAACGGCGCAACCTCCGCCTGCCGGCG AAGCGGCAGCTCTTTCTACGCTGAAATGAAGTGGCTGTTGTCTAACACCGATAACGCCGCTTTCCCTCAGATGACCAAATCCTATAAGAA TACGAGAAAGTCTCCTGCTCTCATCGTATGGGGGATCCACCATAGTGTGTCTACCGCCGAGCAGACCAAACTGTACGGCTCAGGCAACA AATTGGTGACTGTAGGGAGCAGTAATTATCAGCAAAGCTTCGTTCCTTCCCCTGGAGCCAGACCTCAGGTGAATGGACAGTCAGGGCGG ATCGATTTCCATTGGCTGATGCTGAACCCAAACGATACAGTGACATTCTCATTCAATGGCGCGTTCATAGCCCCAGATCGCGCCTCTTTCC TGCGCGGGAAATCAATGGGCATTCAATCTGGGGTACAGGTGGACGCTAACTGTGAGGGGGACTGCTATCATTCCGGGGGCACAATTATT AGTAACCTCCCTTTCCAGAACATTGATTCCAGAGCCGTGGGTAAGTGCCCGCGGTACGTGAAGCAGAGGTCACTGCTGCTAGCCACCGG AATGAAAAATGTGCCTGAAATCCCAAAAGGACGGGGCTTATTCGGCGCTATTGCTGGCTTCATCGAAAATGGGTGGGAGGGGTTGATCG ACGGTTGGTACGGGTTCCGCCACCAGAACGCTCAGGGCGAAGGAACCGCCGCTGACTATAAGTCCACCCAGTCTGCCATCGACCAGAT CACAGGAAAACTGAATAGGCTGATAGAAAAAACCAATCAGCAGTTCGAACTGATTGACAACGAGTTCAACGAGGTCGAGAAACAGATCGG AAATGTTATTAATTGGACCCGAGATAGCATCACTGAAGTCTGGTCCTATAACGCTGAGCTTCTAGTAGCGATGGAAAATCAGCACACGATC GACCTGGCAGATAGCGAAATGGACAAACTCTACGAGAGAGTGAAGCGCCAGCTTCGAGAAAACGCCGAAGAAGATGGAACTGGGTGCT TTCAAATTTTCCACAAGTGTGATGACGATTGCATGGCCTCTATCCGCAATAATACCTACGATCACAGTAAATATCGCGAAGAAGCGATGCA AAATCGTATCCAGATCGATCCTGTAAAGCTCAGCTCCGGATACAAAGACGTGATTCTCTGGTTCAGTTTTGGCGCCAGCTGTTTTATCCTG CTCGCCATAGTCATGGGTTTGGTATTTATCTGCGTCAAGAACGGGAACATGCGTTGCACAATCTGCATA
2895	ATGAATACTCAAATCCTGGTATTCGCCCTAATTGCAATCATTCCAACCAACGCTGACAAGATCTGTTTAGGCCACCACGCCGTTTCGAACG GTACAAAGGTGAATACCCTGACAGAGAGGGGCGTGGAAGTGGTGAACGCGACAGAGACCGTGGAACGGACCAATATCCCCCGCATTTG CTCCAAGGGAAAAAAAACTGTAGACCTCGGCACAGTGCGGCCTGCTCGGAACGATCACCGGACCCCCTCAGTGCGACCAATTTCTGGAAT TTAGTGCAGATCTCATTATTGAGAGGCGGGAAGGAAGTGACGTTTGCTATCCCGGCAAATTTGTGAATGAAGAGGCCCTTCGCCAGATCC TCAGAGAAAGCGGGGGCATAGACAAAGAGGCCATGGGCTTTACCTATTCTGGCATTCGAACCAACGGCGCTACCAGTGCTTGCAGACGG AGCGGCTCGTCCTTCTACGCAGAGATGAAGTGGCTTTTGTCCAATACCGACAACGCTGCTTTCCCTCAGATGACCAAGTCTTACAAGAAC ACCCGAAAGTCTCCAGCGCTGATTGTGTGGGGGATTCATCACTCAGTGAGTACAGCAGAGCAGACTAAGCTGTATGGATCCGGCAATAA ACTCGTGACTGTAGGTTCCTCAAATTATCAACAGAGCTTTGTCCCCTCTCCAGGAGCCAGGCCCCAGGTGAACGGACAATCTGGGCGTAT AGATTTTCACTGGCTGATGTTAAATCCAAACGACACCGTAACCTTCTCGTTTAATGGAGCCTTTATCGCTCCTGACAGAGCGAGCTTTTTA AGAGGAAAAAGTATGGGAATACAGTCCGGAGTTCAGGTGGATGCCAACTGTGAGGGTGATTGCTACCACAGTGGCGGTACAATTATCAG TAATCTACCCTTCCAAAATATCGATTCTCGGGCAGTGGGCAAGTGTCCCCGCTATGTCAAGCAGCGGAGCCTTCTCTTGGCAACAGGTAT GAAAAACGTGCCTGAAATCCCTAAAGGAAGGGGCCTGTTCGGTGCCATTGCAGGATTTATCGAGAATGGGTGGGAAGGGCTTATTGACG GGTGGTATGGATTCCGGCACCAGAATGCCCAGGGGGAAGGCACCGCCGCTGACTACAAATCTACACAGAGTGCCATTGATCAGATCAC GGGCAAACTCAACAGACTTATCGAGAAGACAAATCAGCAATTTGAGCTGATAGACAATGAGTTCAATGAAGTGGAAAAACAGATCGGGAA TGTGATTAACTGGACCAGAGATTCCATCACAGAGGTCTGGTCCTATAACGCCGAACTGCTCGTTGCCATGGAAAACCAGCATACCATCGA TCTTGCCGATTCCGACATGGACAAGCTCTATGAACGCGTCAAGCGACAGCTCAGAGAAAATGCAGAGGAGGATGGGACAGGATGTTTTG AAATTTTCCACAAGTGCGATGACGACTGTATGGCATCCATTCGCAACAACACATACGATCACAGCAAGTATAGAGAGGAGGCCATGCAGA ACAGAATACAGATCGATCCAGTTAAACTGAGTTCTGGCTATAAGGATGTGATTCTCTGGTTCTCTTTCGGAGCCAGCTGCTTTATCCTGCT GGCTATCGTCATGGGCCTGGTGTTCATCTGCGTCAAGAATGGAAACATGCGGTGCACGATTTGCATT

2896	ATGAATACTCAGATCCTTGTGTTCGCCTTAATCGCCATCATCCCTACTAACGCCGACAAGATCTGCCTGGGCCACCACGCTGTTTCCAAT GGCACGAAGGTCAACACTCTCACCGAAAGAGGGGTTGAGGTGGTGAACGCCACCGAAACAGTAGAGAGAACGAACATCCCCCGTATTT GCTCCAAGGGGAAACGCACTGTGGATTTAGGACAGTGCGGCCTGCTGGGGACAATTACCGGACCTCCTCAGTGCGACCAATTCCTGGA ATTTTCAGCTGACCTCATCATAGAGCGGCGAGAGGGGTCCGATGTATGCTACCCTGGAAAGTTTGTGAATGAAGAGGCCCTTCGGCAGA TCCTGCGCCGAGAGCGGGGGGATTGATAAGGAAGCCATGGGTTTTACCTATTCCGGAATTAGAACGAACGGCGCTACATCAGCTTGCCGC CGTTCAGGTTCCAGCTTCTATGCCGAGATGAAGTGGCTACTTTCCAATACCGACAATGCGGCATTCCCTCAGATGACTAAGTCTTACAAG AATACTCGGAAGTCACCAGCCCTTATCGTCTGGGGCATTCACCACAGCGTGTCAACCGCGGAGCAAACCAAACTATATGGGTCGGGGTC TAAACTCGTAACCGTGGGTTCGAGCAATTACCAGCAGAGCTTCGTTCCCAGTCCTGGCGCTCGACCTCAGGTGAATGGGCTGAGTGGGA GGATAGACTTTCATTGGTTAATGTTAAATCCTAACGATACTGTCACCTTCTCCTTCAATGGTGCTTTCATTGCACCAGACCGTGCTAGCTTC CTCCGCGGGAAATCGATGGGCATACAGTCCGGAGTGCAAGTCGACGCTAACTGTGAAGGCGATTGTTACCATTCAGGCGGCACCATAAT CAGCAATCTGCCTTTTCAGAACATTGATAGTCGTGCTGTGGGAAAGTGTCCTAGATACGTGAAGCAGAGGAGTCTTCTTTTGGCCACTGG CATGAAAAACGTACCAGAGATCCCCAAAGGACGGGGCCTGTTTGGCGCTATCGCGGGCTTTATAGAAAATGGCTGGGAGGGATTAATCG ACGGCTGGTACGGTTTTAGGCACCAGAATGCCCAAGGCGAAGGTACAGCAGCCGATTACAAAAGTACACAGTCCGCCATCGATCAGATT ACCGGGAAACTCAATCGTCTCATTGAGAAGACAAACCAACAGTTTGAGCTTATAGACAACGAATTTAATGAAGTGGAGAAACAGATTGGAA ACGTAATCAACTGGACACGGGACTCCATAACTGAGGTCTGGAGCTACAATGCTGAACTCTTGGTGGCCATGGAGAACCAGCATACCATT GACTTGGCAGACTCTGAGATGGATAAGCTGTACGAAAGGGTGAAACGGCAGCTTCGTGAAAATGCCGAGGAGGATGGCACCGGATGTTT TGAGATATTCCACAAGTGTGATGACGACTGCATGGCGAGCATTCGTAACAATACTTACGATCACTCAAAGTACCGGGAGGAAGCCATGCA GAATAGAATCCAGATCGATCCTGTGAAGCTGTCCTCGGGCTACAAGGACGTCATCCTGTGGTTCTCCTTTGGCGCTAGCTGTTTCATTCT GTTGGCGATCGTTATGGGTCTCGTCTTTATTTGCGTTAAGAACGGCAACATGCGTTGCACAATTTGCATT
2897	ATGAATACTCAAATTCTCGTCTTTGCCCTGATTGCTATCATTCCCACCAATGCAGACAAGATTTGCCTCGGTCACCACGCCGTATCAAACG GCACAAAAGTGAACACACTGACCGAAAGAGGTGTCGAGGTCGTTAATGCCACTGAGACCGTTGAGCGGACTAACATCCCTAGAATCTGC AGCAAGGGGAAGAAAACTGTGGATCTTGGGCAGTGCGGACTGCTGGGAACAATAACCGGGCCTCCCCAATGCGACCAATTCCTGGAGT TTTCCGCTGATCTCATTATTGAGCGCAGAGAAGGGAGCGATGTATGCTACCCCGGAAAGTTTGTCAACGAGGAAGCTTTGCGCCAGATTC TGAGGGAATCTGGCGGCATCGACAAAGAGGCTATGGGATTCACATACAGTGGTATAAGGACTAATGGAGCAACCAGCGCTTGTCGGCGG TCGGGGAGCTCCTTTTATGCTGAAATGAAATGGCTGCTTTCTAACACTGATAACGCTGCCTTCCCCCAGATGACTAAATCGTACAAGAACA CACGAAAATCGCCGGCACTGATCGTGTGGGGAATACACCATTCAGTGTCCACTGCCGAACAGACCAAACTGTATGGCAGCGGCAACAAG CTCGTGACCGTTGGTTCCTCGAACTATCAGCAAAGCTTCGTGCCTTCGCCAGGTGCTCGCCCTCAGGTCAATGGACAGTCCGGAAGGAT TGACTTTCACTGGCTTATGTTGAACCCGAACGATACTGTTACATTTAGTTTTAACGGCGCCTTTATTGCACCCGACAGGGCCTCATTTTTG CGAGGCAAGTCGATGGGCATACAATCTGGCGTGCAGGTAGATGCAAACTGTGAAGGCGATTGCTATCACTCCGGAGGCACAATTATTAG TAATCTACCCTTCCAGAATATCGACTCTCGGGCGGTCGGCAAATGTCCCCGATACGTAAAGCAACGCTCCCTGTTGCTGGCCACTGGTAT GAAGAACGTCCCTGAAATCCCCAAAGGCCGGGGATTGTTCGGCGCCATAGCAGGATTCATCGAGAACGGCTGGGAAGGTCTCATAGAT GGTTGGTATGGTTTCCGTCACCAGAACGCCCAGGGGGAAGGCACCGCAGCGGATTACAAGTCGACGCAGTCTGCAATCGACCAGATCA CCGGTAAACTGAACCGCCTGATTGAGAAAACAAACCAACAATTCGAATTAATTGACAATGAGTTTAATGAGGTAGAGAAACAGATTGGCAA CGTGATTAACTGGACAAGGGATTCAATTACCGAAGTTTGGAGCTACAACGCCGAGTTGCTGGTCGCTATGGAAAACCAGCACACAATCGA CCTGGCCGACTCCGAGATGGACAAGTTATATGAACGGGTGAAACGGCAGCTGCGGGAGAATGCCGAGGAGGATGGTACAGGGTGTTTT GAAATCTTTCACAAATGTGATGATGATTGCATGGCGTCAATACCCAACAACACATATGACCACTCAAAATATCCAGAAGAAGCCATGCAGA ACAGAATACAGATTGACCCCGTGAAGCTTTCAAGTGGCTATAAAGATGTCATTCTGTGGTTCTCCTTCGGGGCATCCTGCTTTATCCTGCT CGCCATCGTTATGGGGCTTGTGTTCATTTGTGTCAAGAATGGAAACATGAGGTGCACCATCTGCATC
2898	ATGAATACCCAGATCCTGGTTTTTGCCCTGATCGCTATCATACCAACAAACGCTGACAAGATCTGTCTTGGCCACCACGCTGTCTCTAATG GGACCAAAGTCAATACCCTGACCGAAAGGGGAGTGGAGGTAGTCAACGCTACTGAGACAGTCGAAAGAACTAATATTCCTAGAATTTGCA GTAACGGCAAGAAAACCGTGCGATCTGGGGCAGTGTGGCCTGCTGGGCACCATAACCGGGCCTCCTCAGTGCGATCAGTTCCTGGAGTT TTCCGCCGATCTGATCATTGAACGCAGGGAAGGCAGCGACGTATGCTACCCAGGGAAGTTTGTTAATGAGGAAGCACTCCGCCAGATTC TCCGAGAGTCTGGGGGGTATTGATAAGGAGGCAATGGGTTTTACATACAGCGGCATTAGGACTAATGGGGCTACTTCAGCTTGCCGAAGG AGCGGATCTTCCTTTTACGCTGAGATGAAATGGCTCCTGAGCAATACCGACAATGCCGCCTTTCCTCAGATGACGAAATCTTATAAGAACA CAAGAAAGTCCCCAGCACTGATCGTGTGGGGGATTCACCACTCAGTCTCCACAGCCGAGCAGACCAAGCTGTACGGCTCAGGCAATAAA CTCGTGACGGTTGGCAGCTCCAATTACCAGCAGTCCTTTGTCCCTAGTCCTGGAGCCCGGCCTCAGGTGAATGGGCAGTCCGGTCGAAT AGACTTTCACTGGCTCATGTTAAACCCCAATGACACAGTGACCTTCAGCTTCAATGGGGCTTTCATCGCCCCCGACCGTGCCAGCTTTCT GCGGGGGAAAAGCATGGGGATACAGTCTGGGGTTCAGGTGGATGCCAATTGTGAAGGGGACTGTTACCACTCTGGGGGCACCATCATC AGTAATTTACCATTCCAGAATATTGACTCGAGAGCCGTAGGAAAGTGCCCCAGGTACGTGAAGCAGAGATCTCTGCTGCTTGCGACGGG TATGAAAAATGTCCCAGAGATCCCGAAAGGCCGGGGACTTTTTGGAGCAATCGCCGGCTTCATCGAGAATGGCTGGGAAGGGCTGATCG ACGGCTGGTACGGCTTTCGCCACCAGAACGCCCAGGGCGAGGGTACAGCGGCAGACTATAAGTCCACTCAGAGTGCCATCGATCAGAT CACAGGTAAGCTCAATCGTCTCATTGAGAAGACCAATCAGCAGTTCGAGCTCATCGACAACGAATTCAACGAGGTTGAGAAACAGATCGG CAACGTGATCAACTGGACAAGGGACAGTATCACCGAGGTGTGGAGCTACAACGCTGAGCTGCTGGTGGCTATGGAAAACCAGCACACCA TCGATCTGGCAGATTCCGAAATGGATAAGCTTTATGAGCGGGTGAAGCGTCAACTGAGAGAGAACGCCGAGGACGATGGAACTGGCTG CTTCGAAATCTTTCATAAATGTGATGACGACTGCATGGCCTCAATCCGGAATAACACTTATGACCACAGCAAATACAGAGAAGAGGCGAT GCAGAATCGAATCCAGATAGACCCCGTCAAGCTTTCATCGGGTTATAAGGACGTTATCCTATGGTTTTCTTTTGGCGCCAGCTGCTTTATC CTCCTCGCGATTGTCATGGGACTTGTGTTCATTTGCGTGAAAAACGGCAATATGCGCTGTACAATTTGTATC

2899	ATGAACACACAGATTCTGGTTTTCGCCTTGATTGCCATTATTCCTACTAATGCAGACAAGATCTGTTTGGGCCATCACGCCGTAAGTAACG GTACGAAGGTCAACACCTTGACAGAGCGCGGGGTTGAGGTTGTCAATGCAACAGAAACCGTCGAACGGACAAACATCCCAAGGATCTGC TCAAAAGGCAAGAAAACCGTCGACCTGGGGCAATGTGGCCTTCTCGGAACTATAACCGGACCCCCACAATGCGATCAATTTCTGGAGTT CAGCGCTGACCTAATAATAGAGAGACGGGAGGGGTCAGATGTGTGTTATCCGGGGAAATTCGTGAACGAAGAAGCCCTTAGGCAGATCT TACGGCGAAAGCGGTGGCATCGACAAAGAGGCCATGGGATTTACCTACAGCGGGATCCGGACGAACGGTGCCACTAGTGCATGTCGGAG GTCGGGAAGCTCTTTTTACGCCGAGATGAAGTGGCTTCTGAGCAATACAGATAATGCTGCTTTCCCGCAGATGACAAAGTCTTACAAGAA TACAAGAAAATCCCCAGCTCTCATAGTCTGGGGTATCCATCATTCTGTATCCACGGCCGAGCAAACCAAACTTTACGGCTCTGGGAATAA ACTGGTGACAGTGGGATCATCTAATTATCAGCAGTCTTTCGTGCCATCACCAGGCGCACGACCGCAAGTTAACGGACAGTCTGGTAGAAT CGACTTTCATTGGCTAATGCTGAACCCGAATGATACTGTGACCTTCAGTTTCAATGGGGCTTTCATAGCACCAGACAGGGCATCATTTCTC AGAGGAAAATCAATGGGCATTCAGTCTGGTGTGCAGGTGGATGCAAATTGCGAGGGAGACTGCTATCACAGTGGCGGCACTATCATCTC GAATCTCCCGTTCCAGAATATCGATAGCCGCGCCGTGGGGAAGTGCCCTCGGTACGTTAAACAACGATCTCTGCTGTTAGCCACCGGCA TGAAGAATGTACCCGAGATCCCTAAGGGTCGAGGCCTCTTTGGCGCCATTGCTGGCTTTATCGAGAATGGCTGGGAAGGACTCATAGAC GGCTGGTATGGGTTCCGGCATCAGAACGCTCAGGGAGAAGGGACCGCTGCGGACTATAAGTCAACACAGTCCGCCATTGACCAAATCA CAGGCAAGCTGAACCGGCTGATTGAAAAAACTAATCAACAATTCGAGCTGATAGATAACGAGTTCAACGAAGTCGAAAAGCAGATTGGCA ATGTCATTAATTGGACACGCGATAGCATCACTGAAGTGTGGAGCTACAACGCAGAGCTGCTGGTTGCCATGGAGAACCAGCACACCATC GATCTGGCTGACTCTGAGATGGACAAGCTGTACGAACGGGTGAAGCGTCAGCTCCGTGAGAACGCTGAAGAAGATGGGACAGGTTGCT TCGAAATCTTTCATAAATGCGACGACGATTGTATGGCCTCAATAAGGAACAATACATATGACCACTCTAAGTATAGAGAGGAGGCCATGCA GAATCGTATCCAGATCGATCCAGTGAAGCTGTCTTCAGGGTACAAAGATGTGATCCTCTGGTTTTCATTCGGTGCCAGCTGTTTTATCTTA CTGGCCATCGTCATGGGCCTCGTCTTCATTTGTGTGAAAAACGGTAACATGAGATGCACCATTTGTATT
2900	ATGAATACGCAGATTTTGGTGTTCGCCCTGATCGCCATAATCCCCACTAACGCCGACAAAATCTGTCTGGGGCACCACGCGGTGTCTAAT GGGACTAAGGTAAACACACTGACAGAGAGAGGCGTTGAGGTTGTTAACGCCACGGAGACAGTAGAACGTACAAACATCCCCCGCATCTG CTCCAAGGGCAAAAAGACTGTGGATCTGGGGCAGTGCGGTTTGCTCGGAACCATTACAGGCCCCCCTCAGTGTGACCAATTTCTTGAAT TCTCCGCCGACCTCATCATCGAGCGCCGGGAAGGAAGTGATGTTTGCTATCCCGGGAAATTTGTGAATGAGGAAGCCCTGAGGCAAATT CTCCGGGAGTCAGGGGGTATTGACAAAGAGGCTATGGGGTTTACATACTCCGGTATCCGTACCAATGGAGCAACCTCGGCTTGCCGTAG GTCCGGCAGCTCCTTTTACGCTGAGATGAAATGGCTGCTCTCGAACACCGACAATGCCGCATTTCCCCAGATGACTAAGTCTTACAAGAA TACCCGGAAGTCTCCGGCCCTGATTGTGTGGGGAATCCACCATTCCGTGTCCACCGCCGAACAGACTAAGCTTTACGGCTCGGGCAACA AACTGGTCACCGTTGGCTCTTCTAACTATCAGCAGAGCTTTGTGCCTTCACCAGGGGCTCGACCTCAGGTGAACGGGCAGTCCGGCCGG ATCGATTTCCACTGGCTAATGCTGAACCCAAATGATACTGTTACGTTTTCCTTCAATGGTGCTTTTATAGCACCTGATCGCGCTTCCTTTCT GAGGGGGAAAAGCATGGGCATACAATCTGGGGTTCAGGTTGATGCCAACTGTGAGGGGGATTGCTATCACTCTGGTGGGACCATAATTT CCAACCTCCCCTTTCAGAACATCGACTCACGAGCCGTCGGTAAGTGTCCAAGATACGTGAAGCAACGCTCATTACTCCTGGCTACCGGAA TGAAAAATGTTCCAGAGATCCCGAAGGGGAGGGGTCTGTTCGGTGCCATTGCGGGATTTATTGAAAACGGATGGGAGGGACTAATCGAC GGATGGTATGGGTTTAGGCATCAGAACGCACAGGGCGAGGGAACCGCCGCGGACTATAAAAGCACTCAGTCCGCCATTGATCAGATTAC AGGGAAACTCAATAGACTCATTGAGAAGACGAATCAGCAGTTCGAGCTGATCGACAACGAGTTCAATGAGGTCGAAAAACAGATCGGGA ATGTGATAAATTGGACTAGGGATTCAATAACAGAGGTCTGGAGCTACAACGCAGAGTTACTCGTTGCGATGGAGAACCAACATACTATTG ATCTTGCAGACTCCGAGATGGATAAGCTATACGAGCGGGTCAAGCGACAGTTAAGGGAGAATGCAGAAGAAGATGGAACTGGGTGCTTC GAAATATTTCATAAGTGCGACGACGACTGTATGGCCTCAATCCGCAACAACACATACGACCACTCAAAGTACAGGGAGGAAGCCATGCAC AACCGGATCCAAATTGACCCGGTGAAACTGAGCTCTGGCTACAAGGACGTGATCCTATGGTTTTCTTTTGGGGCCAGTTGCTTCATCCTA CTCGCTATAGTCATGGGTCTGGTGTTCATTTGTGTCAAGAACGGGAACATGCGTTGTACTATCTGCATT
2901	ATGAATACCCAAATCTTGGTGTTCGCGTTGATCGCAATCATCCCCACTAACGCCGACAAAATCTGTTTAGGGCATCACGCTGTTTCCAACG GTACGAAAGTGAATACACTGACTGAACGCGGGGTCGAGGTCGTTAACGCTACAGAGACGGTGGAGAGAACTAATATCCCTAGGATCTGC TCCAAGGGCAAGCGCACTGTGGACCTGGGGCAATGCGGACTGCTAGGGACAATTACTGGCCCGCCCCAGTGCCACCAATTTTTGGAGT TTAGCGCAGACCTGATCATAGAGCGACGCGAGGGTTCAGACGTTTGTTACCCTGGAAAGTTTGTAAATGAGGAAGCCCTTCGCCAGATC CTGCGGGAGAGCGGCGGTATCGATAAAGAGGCTATGGGGTTTACGTACTCTGGAATCAGAACCAATGGGGCCACCTCTGCTTGTAGAAG ATCTGGATCCTCCTTTTATGCAGAAATGAAGTGGTTGTTATCAAATACTGATAACGCGGCATTTCCACAGATGACTAAGAGCTACAAGAAC ACGCGGAAGTCACCCGCTTTAATAGTCTGGGGCATCCACCACTCAGTTAGTACAGCTGAGCAGACTAAGCTGTATGGCAGTGGTAACAA GCTGGTCACCGTGGGTAGCTCTAATTATCAGCAATCTTTCGTACCATCACCCGGAGCTAGACCTCAGGTGAACGGTCTGTCAGGCCGGA TCGACTTCCACTGGTTGATGCTCAATCCCAATGACACTGTTACATTCAGCTTTAATGGCGCCTTCATCGCTCCTGACCGAGCCAGCTTTCT GCGCGGCAAGTCTATGGGGATCCAATCTGGCGTGCAGGTTGATGCGAACTGCGAAGGGGACTGCTACCACAGTGGAGGCACCATCATT TCTAACCTGCCATTCCAGAATATCGACAGTCGCGCCGTGGGGAAGTGTCCTCGATACGTGAAGCAGAGGAGCCTCCTCCTAGCGACTGG AATGAAGAACGTCCCAGAAATCCCTAAGGGCCGGGGGCTCTTTGGAGCGATTGCCGGATTTATCGAAAACGGCTGGGAAGGGCTTATTA ATGGCTGGTACGGCTTCAGGCATCAGAATGCTCAAGGCGAGGGAACTGCCGCCGACTACAAGTCCACTCAATCCGCCATTGATCAGATA ACCGGGAAACTTAACCGCCTGATCGAGAAGACAAACCAGCAGTTTGAGCTGATAGATAATGAGTTCAATGAAGTCGAGAAACAGATAGGA AACGTCATTAATTGGACCCGCGACTCAATTACAGAAGTCTGGAGCTATAACGCAGAGCTGTTGGTAGCGATGGAAAACCAACACACAATT GACCTGGCCGATTCCGAAATGGATAAGCTGTACGAGCGCGTGAAAACACAGTTAAGGGAAAATGCCGAGGAAGATGGTACCGGCTGTTT CGAGATCTTCCACAAGTGCGATGATGATTGCATGGCCTCCATCCGAAATAATACTTACGATCACTCCAAGTACAGAGAGGAGGCAATGCA AAACAGAATTCAGATCGACCCCGTTAAGTTATCAAGCGGATACAAGGACGTCATCCTGTGGTTCTCATTTGGCGCATCGTGTTTTATTCTG CTTGCTATTGTGATGGGTCTCGTTTTCATTTGTGTCAAGAACGGGAATATGAGGTGTACGATCTGCATC

2902	ATGAACACCCAGATTTTAGTCTTTGCACTGATTGCAATTATTCCCACTAATGCTGACAAGATCTGTCTGGGCCACCACGCTGTGTCAAACG GAACTAAGGTGAACACGCTGACTGAGCGAGGTGTCGAAGTGGTGAATGCGACGGAAACCGTGGAGAGAACAAATATCCCCAGAATCTGC AGTAAAGGAAAAAAAACTGTGGATCTGGGCCAATGCGGGCTGCTGGGTACTATCACCGGGCCACCACAATGCGACCAGTTTCTTGAGTT CTCCGCAGATCTCATCATCGAACGCCGGGAAGGCTCCGATGTTTGTTACCCAGGGAAGTTCGTAAATGAAGAAGCCTTAAGGCAAATCCT TAGACAGAGCGGAGGAATTGACAAAGAGGCAATGGGCTTTACCTACTCAGGAATTAGAGCCAACGGCGCTACTTCCGCTTGTCGACGAA GTGGCTCCAGTTTCTATGCCGAAATGAAGTGGCTGTTGTCTAACACGGATAACGCAGCCTTTCCTCAGATGACTAAGTCGTACAAAAACA CCCGGAAATCTCCAGCTCTGATTGTGTGGGGAATCCACCATAGTGTGTCCACCGCCGAGCAAACCAAACTTTATGGGAGCGGAAACAAG TTGGTTACTGTTGGATCCTCGAACTATCAGCAGAGCTTCGTGCCTAGCCCAGGCGCTCGTCCACAGGTGAACGGACTCTCCGGTAGAAT CGATTTTCATTGGCTTATGCTCAATCCTAATGACACTGTGACCTTTAGCTTCAACGGCGCGTTTATCGCCCCTGATCGAGCCTCTTTCCTG CGCGGAAAGTCAATGGGTATCCAGTCGGGCGTTCAGGTGGATGCAAATTGCGAGGGGGACTGCTATCACTCTGGTGGCACCATTATCTC AAACCTGCCATTTCAGAACATCGACTCCCGTGCGGTCGGTAAATGCCCGAGATACGTTAAGCAGCGAAGCCTGCTACTGGCAACAGGCA TGAAGAACGTGCCTGAGATACCTAAGGGACGTGGTCTGTTTGGAGCAATTGCCGGGTTCATTGAGAATGGCTGGGAGGGGCTCATCGAT GGGTGGTATGGCTTTAGACATCAAAATGCTCAGGGAGAAGGTACTGCAGCCGATTACAAGAGTACCCAGAGCGCCATCGATCAGATCAC TGGCAAGTTAAACCGCCTTATCGAAAAGACCAACCAGCAGTTTGAGCTTATCGACAATGAATTTAACGAAGTGGAGAAGCAAATCGGCAA CGTCATCAACTGGACTCGCGATTCTATCACCGAGGTCTGGTCATATAACGCAGAGTTACTCGTTGCGATGGAAAACCAGCACACTATTGA CTTGGCCGACTCTGAGATGGACAAACTGTACGAAAGAGTTAAAAGACAGCTGCGGGAAAACGCTGAGGAAGATGGGACTGGCTGTTTCG AGATCTTTCACAAATGCGATGACGACTGCATGGCTTCGATTCGGAATAATACATACGATCACTCTAAGTACAGAGAAGAGGCTATGCAGA ACAGAATCCAAATCGATCCCGTTAAGCTGTCATCAGGCTACAAGGACGTGATCCTGTGGTTCTCTTTCGGGGCGAGTTGCTTTATCCTTC TTGCCATCGTGATGGGCCTTGTGTTTATTTGTGTTAAAAACGGTAACATGCGCTGTACAATCTGTATA
2903	ATGAACACACAAATTCTGGTCTTCGCACTGATTGCAATCATCCCAACTAACGCTGACAAAATTTGTCTGGGGCACCATGCGGTCAGTAATG GCACCAAGGTCAACACTCTCACTGAACGCGGAGTGGAAGTGGTGAACGCAACCGAGACAGTGGAGCGGACCAACATCCCCAGAATATG CTCAAAGGGTAAAAAGACTGTGGATTTGGGACAGTGCGGTCTTCTTGGGACAATCACCGGCCCTCCCCAGTGTGACCAGTTCCTGGAAT TCTCTGCCGACCTGATTATAGAGCGGCGTGAGGCTCCGATGTGTGTTACCCCGGCAAGTTCGTGAACGAGGAGGCACTGCGACAGAT CTTGAGAGAAAGCGGGGGGATTGACAAAGAGGCTATGGGCTTCACTTATTCGGGCATCAGGACTAATGGCGCAACCTCCGCTTGTAGGA GATCAGGATCCTCTTTCTACGCCGAAATGAAGTGGCTGCTGTCGAATACAGACAATGCCGCTTTCCCCCAAATGACTAAGAGTTATAAAAA CACGCGTAAATCCCCAGCGCTGATTGTCTGGGGAATCCACCACTCGGTGAGCACTGCAGAGCAGACTAAACTGTACGGGAGCGGCAATA AACTTGTTACAGTGGGATCCTCGAATTACCAACAGAGCTTCGTCCCATCACCAGGTGCTCGACCACAGGTGAACGGCCAGTCTGGGCGA ATAGATTTTCATTGGCTCATGCTTAATCCCAATGACACTGTGACTTTTTCTTTCAATGGGGCCTTCATCGCGCCTGATCGGGCCAGTTTCC TGAGAGGCAAGAGTATGGGAATCCAGTCCGGCGTGCAGGTGGACGCTAATTGTGAGGGGGATTGCTATCACTCAGGTGGCACTATAATT TCAAATCTTCCTTTCCAGAACATCGATTCTCGAGCCGTCGGAAAGTGCCCTAGGTACGTGAAACAAAGGTCACTCCTTTTGGCAACCGGG ATGAAGAACGTGCCAGAGATTCCCAAGGGTCGGGGGCTCTTTGGCGCTATTGCTGGATTTATCGAGAATGGGTGGGAGGGTCTGATAGA CGGTTGGTATGGCTTCAGACACCAGAACGCCCAGGGGGAGGGCACCGCCGCCGACTATAAGAGTACCCAGAGCGCAATTGACCAGATT ACCGGAAAATTGAACCGACTGATTGAGAAGACAAATCAGCAGTTTGAGTTGATCGACAACGAATTCAATGAGGTTGAAAAACAGATCGGG AACGTTATAAACTGGACACGAGATAGCATAACAGAGGTCTGGAGTTACAATGCTGAACTGCTGGTTGCTATGGAAAACCAGCATACTATT GACCTTGCGGACAGCGAGATGGACAAACTATACGAAAGAGTGAAGCGCCAGCTGCGTGAGAACGCCGAAGAAGACGGGACTGGATGCT TCGACATATTCCACAAGTGTGATGACCATTGTATGCCAAGCATACGGAACAACACATACGATCATAGCAAATATACGGAGGAGGCGATGC AGAACAGAATACAGATAGACCCTGTGAAGTTATCTTCCGGTTACAAAGACGTTATCCTCTGGTTTAGCTTCGGAGCCAGCTGCTTTATATT ATTAGCCATCGTCATGGGCCTCGTTTTCATCTGCGTGAAGAATGGAAATATGCGCTGCACCATCTGCATC
2904	ATGAATACGCAGATCTTGGTATTCGCACTTATAGCCATCATCCCCACCAACGCCGATAAGATTTGCTTAGGCCACCATGCTGTTTCTAACG GCACAAAAGTGAACACACTTACCGAGAGAGGGGTCGAAGTTGTGAACGCCACGGAGACAGTCGAAAGGACAAACATTCCTCGAATCTGC AGTAAGGGCAAGAAAACAGTGGATCTCGGCCAGTGTGGGCTGCTGGGAACCATCACGGGTCCTCCGCAGTGTGACCAGTTCCTGGAGT TCTCTGCCGATCTCATCATCGAAAGGCGGGAGGGATCCGACGTGTGCTACCCAGGCAAATTCGTGAACGAAGAAGCGCTTCGGCAAATT CTGCGAGAGTCAGGGGGGATAGATAAAGAGGCAATGGGGTTTACTTATTCTGGCATTAGAACCAATGGGGCCACGAGTGCATGTAGACG GTCCGGAAGTAGCTTTTACGCAGAAATGAAGTGGCTGTTGTCTAACACTGACAACGCTGCATTTCCACAGATGACCAAGTCTTATAAAAAC ACGAGAAAATCCCCAGCCCTCATTGTGTGGGGGATCCACCATTCTGTGAGCACCGCCGAGCAAACGAAGCTGTATGGCAGTGGTAACAA GCTAGTCACAGTAGGATCCAGTAATTACCAGCAAAGCTTCGTGCCCAGTCCCGGAGCTCGGCCTCAGGTCAACGGGCAATCTGGGAGAA TCGACTTCCACTGGCTGATGCTGAACCCAAACGACACAGTGACATTTTCTTTTAATGGCGCTTTCATTGCCCCCGACAGGGCTAGTTTTCT GAGAGGCAAGTCCATGGGGATCCAAAGCGGAGTGCAGGTGGACGCTAATTGCGAGGGCGATTGTTACCACAGTGGGGGGACCATTATC AGTAATCTGCCTTTCCAGAACATCGATAGTAGAGCCGTGGGCAAATGTCCACGTTACGTCAAGCAGCGGAGCCTGCTCTTAGCCACGGG CATGAAGAACGTGCCTGAGATCCCAAAGGGCAGAGGGCTGTTTGGCGCTATCGCGGGATTCATTGAAAATGGGTGGGAGGGACTAATC GACGGATGGTACGGTTTTCGACACCAGAACGCACAGGGGGAAGGGACAGCTGCCGATTATAAGTCTACTCAATCTGCTATCGACCAGAT AACTGGGAAACTTAATCGGCTGATTGAGAAAACAAACCAGCAGTTTGAGCTGATAGACAACGAATTCAACGAAGTGGAGAAACAGATCGG TAACGTGATAAACTGGACTAGGGACTCTATCACCGAGGTGTGGTCCTACAACGCGGAACTCCTTGTCGCTATGGAGAATCAGCACACTAT AGACCTGGCAGACTCTGAAATGGATAAACTGTACGAACGGGTTAAGCGGCAATTGCGGCAGAACGCTGAGGAGGATGCCACAGGATGT TTTGAAATTTTTCATAAATGTGACGACGATTGCATGGCTTCCATTCGCAACAATACTTATGATCATTCGAAGTACCGGGAAGAGGCCATGC AAAACCGTATCCAGATCGATCCCGTCAAACTGTCGTCGGGGTACAAGGACGTTATACTTTGGTTCAGCTTTGGAGCCTCTTGTTTCATCTT GCTCGCTATAGTGATGGGCTTGGTGTTTATATGCGTAAAAAATGGCAACATGCGGTGCACAATTTGCATT

2905	ATGAACACTCAGATACTCGTCTTTGCTCTCATTGCAATTATTCCTACCAATGCTGATAAGATTTGCTTGGGGCACCACGCAGTTTCCAATG GGACAAAGGTGAACACCCTAACTGAGCGGGGCGTTGAAGTGGTGAACGCAACCGAGACAGTGGAAAGAACAAATATCCCCCGCATTTGC AGTAAAGGCAAAAAAACCGTCGACTTGGGACAGTGTGGCCTGCTTGGGACTATCACCGGACCCCCACAGTGTGACCAATTTTTGGAGTTT TCAGCTGATCTGATTATCGAAAGACGCGAGGGAAGCGATGTTTGCTATCCAGGAAAATTTGTGAATGAAGAAGCACTCCGACAGATCCTG CGGGAGTCCGCCCGAATAGATAAGCAAGCAATGGGGTTCACTTACTCGGGCATCCGGACTAACCGAGCCACTTCCGCCTGCAGAAGGT CAGGCTCGTCGTTCTATGCCGAGATGAAGTGGCTGCTTTCAAACACTGATAATGCCGCATTCCCACAGATGACAAAGTCATACAAAAACA CCAGGAAATCACCCGCATTGATAGTGTGGGGCATCCATCATTCTGTGTCCACTGCCGAGCAGACTAAACTGTACGGGAGCGGGAATAAG CTTGTTACCGTCGGGTCAAGCAACTATCAGCAAAGTTTTGTACCAAGCCCGGGCGCACGGCCCCAGGTTAATGGTCAGTCCGGTCGCAT TGATTTCCATTGGCTAATGCTAAATCCAAATGACACTGTAACGTTCTCATTCAACGGAGCGTTCATCGCTCCAGACCGTGCCTCTTTTCTG CGCGGGAAAAGCATGGGGATACAGTCCGGGGTGCAGGTAGATGCTAACTGTGAGGGAGACTGTTATCACTCTGGCGGCACGATCATCT CCAACCTTCCCTTTCAGAACATTGACAGCAGAGCCGTCGGGAAATGCCCCCGATATGTTAAGCAGAGGTCACTTCTTCTCGCTACGGGTA TGAAGAATGTCCCAGAAATTCCTAAAGGCCGGGGATTGTTTGGAGCAATCGCCGGCTTCATCGAAAACGGTTGGGAAGGCTTGATTGAT GGCTGGTACGGCTTCCGCCACCAAAATGCTCAGGGAGAAGGAACTGCAGCTGACTATAAATCAACACAGAGCGCCATTGACCAGATCAC AGGGAAGCTGAACAGACTTATCGAGAAAACCAATCAGCAGTTTGAGCTCATAGACAACGAATTTAACGAAGTGGAGAAACAGATTGGCAA CGTTATCAATTGGACCCGGGACTCAATCACTGAAGTCTGGTCCTACAATGCAGAGCTGCTCGTCGCTATGGAAAACCAGCACACTATCGA TTTGGCTGACAGCGAAATGGATAAACTATATGAAAGGGTCAAGCGTCAGCTCCGAGAAAATGCAGAAGAGGATGGAACAGGCTGTTTCG AAATCTTCCATAAGTGTGATGACGATTGCATGGCCAGTATCAGAAACAACACTTACGATCACTCCAAATACAGGGAGGAGGCCATGCAGA ACCGGATCCAGATTGACCCTGTAAAACTGTCGAGCGGCTATAAAGATGTTATCCTGTGGTTCAGCTTCGGGGCCAGCTGCTTCATACTGC TTGCCATTGTGATGGGACTGGTGTTCATTTGCGTTAAAAATGGAAACATGAGATGCACCATCTGTATC
2906	ATGAACACACAAATCCTGGTGTTCGCTCTAATAGCGATCATCCCAACCAATGCCGATAAGATCTGTCTGGGCCACCACGCCGTATCCAAT GGGACTAAGGTTAATACACTGACAGAGCGAGGAGTCGAAGTGGTGAACGCAACCGAGACAGTCGAAAGGACGAATATTCCCAGGATCTG CAGCAAGGGGAAGAAGACTGTTGACCTGGGACAATGTGGCCTACTGGGAACTATAACCGGCCCTCCACAATGCGACCAATTCCTGGAGT TTTCCGCTGACCTGATTATTGAACGGCGCGAAGGGAGCGACGTATGTTATCCCCGGGAAATTCGTGAATGAGGAGGCTCTGCGCCAGATC CTGCGCGAGAGCGGGGGAATCGATAAGGAGGCCATGGGTTTCACCTATTCCGGAATAAGGACTAACGGCGCAACCTCAGCTTGTAGAC GCAGTGGCTCTTCTTTTTATGCTGAAATGAAATGGTTATTATCAAACACGGACAACGCCGCTTTTCCCCAGATGACCAAGAGTTATAAGAA CACCCGGAAATCACCTGCTCTGATAGTATGGGGAATTCATCACTCGGTGTCCACTGCGGAGCAAACCAAACTGTACGGGAGCGGCAATA AGTTGGTTACAGTAGGAAGCTCCAATTATCAGCAAAGTTTCGTCCCCTCCCCTGGCGCCAGGCCTCAAGTGAACGGACAGTCAGGGAGA ATCGACTTCCACTGGCTGATGCTCAATCCAAACGACACCGTAACTTTCTCATTCAATGGGGCATTTATCGCCCCAGACCGGGCCTCGTTC TTACGAGGTAAATCCATGGGGATTCAGAGCGGCGTCCAAGTGGATGCCAACTGCGAAGGCGATTGCTACCACTCTGGTGGGACAATCAT CTCGAACTTGCCGTTCCAAAACATCGATTCCCGAGCAGTGGGGAAATGCCCCAGATACGTAAAACAAAGGTCGCTGCTCCTTGCTACCG GCATGAAAAACGTCCCTGAAATCCCAAAGGGTCGTGGTCTGTTTGGCGCAATTGCTGGGTTTATTGAGAACGGCTGGGAAGGCCTGATA GACGGGTGGTATGGATTTCGTCATCAGAATGCACAGGGCGAGGGGACCGCAGCCGATTACAAGAGTACCCAGTCCGCTATTGACCAGAT AACAGGGAAACTGAACCGGCTGATTGAGAAGACCAACCAGCAGTTCGAGTTGATTGACAATGAGTTCAACGAAGTCGAGAAGCAGATCG GGAATGTAATAAATTGGACAAGAGACAGCATTACAGAGGTGTGGAGCTACAATGCAGAGCTGTTAGTGGCCATGGAGAACCAACACACG ATTGACCTTGCCGATTCCGAGATGGACAAGTTGTCACGAGAGAGTAAAGAGGCAGCTGAGGGAGAATGCTGAAGAAGACGGAACAGGCT GCTTTGAAATCTTCCACAAGTGTGATGATGACTGTATGGCTTCGATACGCAATAATACATACGACCATAGCAAGTATCGGGAGGAAGCCA TGCAGAATAGAATTCAGATTGATCCCGTGAAACTAAGCTCGGGATACAAGGACGTGATACTATGGTTTTCATTTGGCGCCTCCTGTTTCAT CCTCCTGGCTATTGTCATGGGTTTGGTTTTTATTTGTGTTAAAAACGGCAACATGCGGTGTACTATTTGCATC
2907	ATGAACACGCAGATACTGGTTTTCGCACTCATTGCAATCATCCCTACGAACGCCGACAAGATTTGTCTGGGCCACCACGCTGTCTCTAAC GGCACCAAGGTGAACACCCTGACCGAGCGCGGAGTGGAAGTGGTCAATGCGACAGAAACTGTGGAACGGACAAACATTCCACGGATCT GTAGCAAAGGCAAGAAGACAGTGGATCTGGGGCAGTGCGGCCTACTGGGCACCATCACCGGACCTCCTCAATGCGATCAGTTTCTTGAG TTTTCCGCCGATTTGATTATAGAAAGACGGGAGGGATCCGATGTCTGCTACCCAGGCAAGTTTGTCAACGAGGAGGCCCTAAGACAGATT CTGCGGGAATCTGGCGGGATCGACAAGGAGGCAATGGGCTTCACCTACAGCGGGATCAGGACAAACGGAGCCACAAGCGCTTGTAGGC GCTCTGGTTCCTCGTTTTACGCAGAAATGAAGTGGTTACTTTCAAATACGGACAATGCGGCCTTCCCCCAGATGACCAAATCATATAAGAA TACTCGCAAGAGCCCGGCATTGATTGTCTGGGGCATCCATCATTCTGTGTCTACTGCTGAACAGACTAAACTGTATGGCTCTGGCAACAA GCTGGTGACTGTCGGAAGCAGTAACTACCAACAGAGCTTCGTGCCTAGCCCAGGGGCAAGACCCCAGGTCAACGGGCAGAGCGGGCG TATTGACTTTCACTGGCTGATGCTGAACCCAAACGACACTGTGACATTCAGTTTCAATGGGGCCTTTATAGCCCCTGACAGGGCCAGCTT CTTGAGGGGCAAGTCCATGGGAATACAGTCTGGTGTCCAGGTCGATGCCAATTGCGAGGGGGACTGTTATCATAGCGGCGGCACAATCA TCTCAAATCTGCCTTTCCAGAATATCGATAGTCGTGCCGTGGGAAAATGCCCCCGTTATGTAAAGCAAAGATCCTTGCTGCTAGCTACCG GGATGAAGAATGTGCCAGAAATTCCAAAGGGACGAGGCCTGTTCGGCGCTATTGCTGGCTTCATTGAGAACGGGTGGGAAGGCCTGATA GATGGGTGGTACGGGTTCAGGCACCAGAACGCACAGGGCGAGGGTACAGCTGCCGACTATAAATCTACGCAGAGCGCAATTGACCAGA TCACCGGGAAGCTAAACCGTCTGATTGAAAAGACCAATCAACAATTTGAGCTCATAGATAATGAGTTCAACGAAGTTGAAAAGCAGATTGG GAACGTCATCAACTGGACACGGGATAGTATAACGGAAGTGTGGAGTTATAACGCCGAATTACTGGTGGCGATGGAAAACCAGCACACTA TCGATTTGGCCGACTCCGAGATGGATAAGCTGTACGAGCGGGTTAAACGCCAGTTAAGGGAGAATGCTGAGGAGGATGGCACAGGTTG CTTTGAGATTTTCCATAAATGCGATGACGATTGCATGGCCTCTATTCGAAATAACACATACGACCACTCGAAGTACCGTGAGGAGGCAAT GCAAAATCGAATCCAGATTGACCCAGTGAAACTCTCCAGTGGCTATAAGGATGTGATCTTGTGGTTCTCTTTTGGGGCCTCATGTTTTATT CTCCTTGCGATCGTGATGGGATTGGTTTTTATCTGTGTCAAGAATGGAAACATGAGATGCACGATCTGTATC

2908	ATGAATACACAGATCTTAGTTTTTGCACTAATAGCTATCATCCCCACAAACGCCGACAAAATTTGTCTGGGTCATCATGCTGTGTCTAACG GTACCAAGGTTAATACCCTGACCGAACGCGGGGTTGAGGTGGTTAACGCGACTGAAACGGTGGAACGCACGAATATACCCAGGATCTGT TCAAAAGGCAAAAAGACCGTTGATCTGGGTCAGTGCGGCCTTCTGGGGACAATCACCGGCCCCCCCCAGTGCGACCAGTTCCTCGAGTT TAGTGCTGATTTGATCATTGAGCGAAGAGAGGGGTCCGACGTGTGCTATCCAGGGAAGTTTGTTAATGAAGAGGCGTTGCGCCAGATCC TACGCGAATCCGGGGGAATCGATAAGGAAGCAATGGGCTTCACCTATTCTGGCATTAGAACCAACGGAGCTACATCCGCCTGTAGAAGG AGTGGCTCGTCTTTCTATGCAGAGATGAAATGGCTGCTGTCCAACACCGACAATGCCGCTTTCCCCCAGATGACTAAAAGCTACAAGAAT ACCCGAAAAAGTCCAGCTCTGATCGTCTGGGGCATCCACCACTCAGTTAGTACAGCCGAACAGACAAAGTTGTATGGCAGTGGCAACAA ACTAGTCACAGTTGGCTCGTCCAACTATCAGCAGAGTTTTGTGCCCAGTCCCGGGGCACGTCCGCAGGTCAATGGACAGAGCGGCCGG ATTGATTTCCATTGGCTGATGCTGAACCCTAACGATACAGTGACATTTTCCTTCAACGGCGCGTTCATCGCACCCGATCGCGCTTCATTCC TGAGGGGGAAGAGCATGGGGATTCAGTCGGGAGTCCAGGTTGACGCCAACTGTGAAGGAGACTGCTACCATTCAGGGGGGACAATCAT CTCCAACCTACCATTCCAGAACATTGATTCAAGAGCTGTTGGAAAGTGTCCAAGATATGTCAAGCAAAGGTCTCTATTGCTAGCTACCGGT ATGAAGAATGTCCCTGAAATACCGAAGGGGAGGGGTCTGTTCGGTGCCATAGCTGGTTTTATCGAAAATGGGTGGGAGGGTCTTATCGA CGGGTGGTATGGATTTCGCCACCAGAACGCCCAGGGGGAAGGCACTGCAGCTGACTATAAGTCGACTCAGTCTGCCATCGACCAGATC ACCGGTAAATTAAATAGACTGATTGAGAAGACAAATCAGCAATTCGAGCTCATCGATAATGAGTTCAACGAGGTCGAGAAGCAGATTGGA AATGTGATTAACTGGACGCGCGATTCCATTACCGAGGTGTGGAGCTATAACGCGGAGTTGTTAGTGGCGATGGAGAATCAGCATACAATT GATCTAGCGGACAGTGAAATGGACAAGCTGTACGAGAGAGTTAAACGACAGCTGCGCGAGAACGCCGAGGAGGATGGGACAGGGTGTT TTGAGATCTTCCATAAGTGTGACGATGATTGTATGGCTAGCATTAGAAATAACACTTACGATCATAGCAAATACAGGGAGGAGGCCATGCA GAACCGAATTCAAATCGACCCCGTGAAGCTGTCTTCGGGCTACAAAGACGTTATTCTCTGGTTTAGCTTTGGTGCGTCATGCTTCATTCTG CTGGCGATTGTAATGGGGTTGGTTTTCATATGCGTTAAGAACGGCAACATGCGGTGCACTATTTGTATT

Table 18: Codon Optimized Sequences Encoding H10 Hemagglutinin

Seq ID No.	Nucleic Acid Sequence

	5'$^{7Me}G_{ppp}G_{2'OMe}$GGAAAUAAGAGAGAAAAGAAGAGUAAGAAGAAAUAUAAGAGCCACCAUGUACAAAAUCGUCGUCAUUAUCGCAUUGCUGG GCGCUGUCAAGGGACUCGACAAGAUCUGUCUCGGACACCACGCGGUGGCUAACGGAACCAUCGUCAAGACCCUGACUAACGAGCAGGA AGAAGUGACCAACGCCACCGAAACUGUGGAAUCCACCGGGAUCAACAGAUUGUGCAUGAAGGGUCGGAAGCACAAGGAUCUGGGAAAC UGCCACCCGAUUGGAAUGCUCAUCGGCACCCCGGCAUGCGAUCUGCAUCUGACUGGGAUGUGGGAUACCCUUAUCGAGCGGGAAAAC GCGAUCGCAUACUGUUACCCCGGUGCCACCGUGAACGUGGAGGCGCUCAGACAGAAGAUUAUGGAGUCAGGCGGCAUCAACAAGAUUU CCACGGGCUUCACCUACGGGUCCUCCAUUAACUCCGCCGGUACCACUCGGGCCUGCAUGCGGAACGGAGGGAACUCCUUUUACGCCG AGCUCAAGUGGCUUGUGUCAAAAUCCAAGGGACAGAAUUUCCCCCAAACCACCAACACCUAUAGGAACACCGACACCGCCGAACAUCUC AUUAUGUGGGGCAUCCAUCACCCUUCGAGCACACAGGAGAAGAAUGACCUCUACGGCACCCAGUCGCUGAGCAUCUCCGUGGGCUCAU CGACCUAUCGCAACAACUUCGUGCCUGUGGUCGGCGCCCGACCUCAAGUCAACGGACAGUCCGGACGCAUUGACUUCCAUUGGACUCU GGUGCAACCGGGAGACAACAUCACUUUCUCCCACAACGGCGGACUGAUUGCCCCAAGCCGCGUGUCAAAGCUGAUCGGUAGAGGGCUG GGUAUUCAGUCGGAUGCUCCCAUCGAUAACAACUGCGAAUCCAAGUGCUUUUGGAGAGGCGGCUCCAUCAAUACUCGGCUGCCGUUUC AGAACCUGAGCCCGAGGACCGUGCGGCAGUGCCCAAAAUACGUCAAUCGCCGGUCACUGAUGCUGGCGACCGGAAUGAGGAACGUGC CUGAACUCAUCCAGGGACGGGGGCUGUUCGGCGCCAUCGCCGGCUUCCUGGAAAACGGAUGGGAGGGAAUGGUGGACGGUUGGUAC GGCUUCCGCCACCAAAACGCCCAGGGAACUGGACAGGCCGCCGACUACAAGUCCACACAGGCCGCAAUAGACCAGAUCACUGGGAAGC UCAACCGCCUUGUGGAAAAGACUAACACUGAAUUCGAGUCGAUCGAGUCUGAGUUCUCCGAGAUCGAACACCAGAUUGGGAACGUGAU CAACUGGACGAAGGACUCCAUUACCGACAUCUGGACCUACCAGGCGGAACUGCUCGUGGCGAUGGAGAACCAGCACACUAUCGACAUG GCCGACUCCGAGAUGCUGAAUCUGUACGAACGCGUGCGGAAGCAACUGAGACAAAACGCUGAAGAAGAUGGAAAGGGUUGUUUCGAGA UCUACCACGCCUGCGACGACAGCUGCAUGGAAAGCAUUAGGAACAAUACCUACGACCACUCGCAGUACCGGGAGGAGGCCCUUCUGAA CCGGCUGAACAUUAACCCCGUGACCUUGAGCUCCGGCUACAAGGACAUCAUCCUGUGGUUCUCGUUUGGAGCCAGCUGCUUCGUGCU GCUGGCCGUCGUGAUGGGAUUGUUCUUCUUCUGCCUGAAAAACGGAAACAUGCGCUGCACCAUCUGUAUUUGAUAAUAGGCUGGAGCC UCGGUGGCCAUGCUUCUUGCCCCUUGGGCCUCCCCCCAGCCCCUCCUCCCCUUCCUGCACCCGUACCCCCGUGGUCUUUGAAUAAAG UCUGAGUGGGCGGCAAAA

3076	ATGTACAAGATCGTGGTGATCATCGCTCTGCTCGGCGCCGTTAAGGGACTGGATAAGATTTGTTTAGGGCACCATGCAGTGGCAAACGGGA CCATTGTCAAGACACTCACCAATGAACAGGAGGAAGTGACCAACGCGACTGAGACTGTTGAGTCCACCGGCATAAACAGGCTTTGTATGAA AGGCAGAAAGCACAAGGATTTGGGTAATTGTCACCCAATAGGAATGCTAATTGGGACTCCTGCCTGCGATCTCCACCTGACAGGCATGTGG GACACCCTGATTGAACGGGAGAATGCTATCGCATACTGCTACCCAGGTGCAACCGTAAACGTGGAGGCCCTGCGACAGAAGATAATGGAG AGCGGAGGCATCAACAAGATCAGTACGGGATTTACCTACGGGTCTTCTATCAATTCGGCTGGCACAACACGTGCTTGCATGAGAAACGGAG GCAACTCCTTCTACGCTGAACTCAAATGGTTGGTGAGTAAGAGTAAGGGTCAGAATTTCCCACAGACGACCAACACATACAGGAACACCGAT ACGGCCGAACACCTCATCATGTGGGGCATACACCACCCTAGTAGTACACAAGAGAAAAACGACCTGTATGGGACACAATCCCTTTCAATCA GTGTTGGTAGCTCGACCTACCGTAATAATTTCGTCCCTGTGGTGGGTGCTAGACCGCAGGTCAATGGCCAGTCAGGAAGGATCGATTTCCA CTGGACCCTCGTGCAGCCAGGTGATAACATTACTTTTTCTCATAATGGCGGACTAATTGCGCCTAGCAGGGTGTCAAAGCTTATTGGGAGAG GCCTAGGCATACAAAGCGACGCCCCAATCGATAACAATTGCGAGTCGAAGTGTTTCTGGAGGGGCGGATCAATTAACACACGTCTGCCCTT CCAGAATCTGTCACCAAGGACTGTGGGCCAGTGTCCAAAGTATGTTAATCGACGTTCACTCATGCTTGCCACCGGCATGCGGAATGTGCCT GAACTGATTCAGGGCCGCGGACTTTTTGGCGCCATTGCAGGTTTCTTGGAGAACGGGTGGGAAGGTATGGTCGACGGATGGTATGGCTTC AGACACCAGAACGCCCAAGGAACTGGCCAGGCCGCCGATTATAAGAGCACCCAGGCTGCGATCGATCAAATTACCGGTAAACTCAACCGG CTGGTCGAGAAGACTAATACAGAGTTTGAGTCAATAGAATCCGAGTTTTCCGAAATAGAACATCAGATCGGTAATGTAATCAACTGGACCAA GGACAGCATCACCGACATATGGACATACCAGGCCGAACTGCTCGTCGCCATGGAGAATCAACACACGATTGACATGGCCGATTCGGAAATG CTGAATCTGTACGAAAGGGTGCGTAAACAGTTGCGGCAGAATGCAGAAGAAGATGGAAAAGGGTGTTTTGAAATCTACCACGCCTGCGACG ATAGCTGCATGGAATCTATCCGGAATAACACCTATGACCATTCCCAATACCGAGAAGAGGCCCTGCTCAACCGCTTGAACATTAATCCCGTT ACATTGTCATCCGGATACAAGGACATCATCTTGTGGTTCTCTTTCGGGGCTTCCTGTTTTGTGCTTCTAGCAGTTGTGATGGGTCTGTTTTTC TTCTGCTTAAAGAACGGCAATATGAGATGTACTATTTGCATC
2909	ATGTACAAGGTAGTGGTAATTATCGCTCTGCTAGGTGCCGTTCGTGGCTTGGACAAGATTTGTTTGGGGCACCATGCCGTAGCCAACGGAA CAATTGTGAAGACACTGACTAACGAACAAGAGGAAGTAACCAACGCTACAGAAACAGTCGAATCTAAAAGTTTGGGCAAGTTATGTATGAAA GGGCGGAGTTACAACGACCTCGGAAATTGCCACCCCATCGGAATTTTAATCGGCACTCCAGCCTGCGATCTGCACTTAACAGGAACCTGGG ACACCCTCATCGAACGCGAAAATGCCGTTGCATACTGTTACCCAGGTGCGACTGTCAACGAGGAAGCGCTCCGCCAGAAGATCATGGAAAG CGGGGGGATTTCAAAAATCTCTACCGGGTTTACATACGGGACTTCCATCAACAGCGCTGGCACCACAAAGGCCTGCATGCGCAATGGCGGT AACAGCTTCTACGCTGAGCTCAAGTGGCTGGTCAGCAAAAACAAAGGCCAGAACTTCCCCCAAACTACTAATACCTACAGAAACACAGATAC CGCCGAGCACCTCATAATTTGGGGGATCCATCACCCATCTAGCACTCAGGAGAAAAACGATCTCTATGGCACACAGTCTCTCAGCATTTCAG TGGGAAGCTCGACATACCAGAATAACTTTGTTCCCGTGGTGGGCGCAAGACCACAGGTCAACGGCCAGTCCGGGAGAATTGATTTCCATTG GACGCTGCTGCAACCTGGAGACAACATCACCTTTTCTCATAATGGCGGCCTAATTGCGCCTTCCAGAGTTAGCAAGCTGATCGGCCGGGGC CTAGGTATTCAGAGCGAGGCCCCAATTGACAACGGTTGCGAATCAAAGTGCTTCTGGAAAGGCGGATCCATTAACACGAAGTTGCCCTTCC AGAACTTATCACCCAGAACCGTGGGGCAGTGTCCCAAATACGTGAATAAGAGGTCCCTGATGCTCGCTACCGGTATGCGGAATGTGCCTGA GATAATGCATGGAAGAGGACTGTTTGGGGCAATCGCTGGGTTTATCGAAAACGGTTGGGAAGGGATGGTTGATGGATGGTACGGGTTTCG GCATCAGAATGCACAGGGAACTGGTCAGGCCGCCGATTACAAATCAACGCAAGCCGCCATCGACCAGATAACTGGAAAGCTGAATAGGCT CATCGAAAAAACCAACACCGAGTTCGAATCAATTGAGTCTGAGTTCTCTGAGATCGAGCACCAAATCGGGAATATCATCAATTGGACCAAAG ATAGCATTACTGATATCTGGACATATCAGGCTGAGCTGCTGGTCGCCATGGAGAACCAGCATACAATCGACATGGCAGACTCGGAGATGCT TAACCTGTATGAGAGGGTGCGTAAACAATTGCGACAGAATGCCGAAGAGGATGGTAAGGGTTGCTTCGAAATCTACCACGCTTGCGACGAT TCTTGCATGGAAAGTATAAGAAATAATACCTATGATCATTCACAATACCGCGAAGAGGCTTTGCTGAACAGACTCAACATCAATCCGGTGAAG CTTTCCTCTGGGTACAAGGATATAATTCTGTGGTTTTCCTTTGGAGCTTCCTGTCTCATATTATTGGCTGTTGTTATGGGCCTCGTTTTTTTTT GCCTGAAGAACGGTAACATGCGCTGTACAATATGTATC
2910	ATGTACAAAATTATCGTGATCATCGCTCTTCTGGGCGCCGTCAAAGGTCTCGATAAAATTTGCCTTGGGCACCATGCTGTGGCAAACGGAAC TATTGTAAAAACTCTTACGAATGAGCAAGAAGAGGTGACCAATGCCACCGAAACCGTCGAGTCCACCGGGATCAATAGGTTGTGCATGAAG GGGCGTAAACACAAGGACCTTGGCAATTGCCATCCGATCGGCATGTTGATCGGAACACCAGCTTGTGACCTTCACTTGACCGGAACATGGG ACACTCTGATCGAGCGAGAGAACGCAATTGCATATTGCTATCCGGGAGCAACCGTGAATGTGGAGGCCCTGCGGCAGAAGATCATGGAAT CCGGAGGTATCGACAAAATCTCTACCGGATTCACCTACGGATCTTCAATTAACTCCGCTGGCACTACACGCGCTTGCATGCGAAATGGCGG TAACTCATTCTACGCCGAGCTGAAGTGGCTGGTGTCCAAGTCTAAAGGGCAGAATTTTCCGCAGACCACTAACACCTACAGAAACACTGATA CCGCCGAACATCTCATTATGTGGGGAATCCACCACCCATCAAGCACTCAAGAAAAAAACGACTTGTATGGTACACAGTCACTAAGCATTTCC GTGGGCAGCTCTACCTACCGCAACAACTTTGTCCCCGTGGTTGGCGCCCGACCCCAAGTAAATGGTCAGTCCGGCAGGATCGATTTTCATT GGACCCTGGTGCAGCCAGGGGATAACATCACTTTTAGCCATAACGGCGGCTTGATAGCGCCGTCAAGGGTGTCCAAACTGATCGGAAGAG GGCTGGGAATTCAGAGTGACGCACCCATCGATAACAACTGCGAAAGCAAATGTTTCTGGCGGGGGGGTTCGATCAACACCCGATTACCTTT CCAGAACCTCTCCCCCAGGACTGTTGGTCAGTGCCCGAAATATGTTAATCGACGCAGCTTGATGCTCGCCACCGGGATGCGCAATGTTCCC GAGCTCATTCAGGGGCGTGGCCTCTTTGGCGCAATAGCTGGGTTCCTCGAGAATGGTTGGGAAGGCATGGTGGATGGATGGTACGGCTTC CGGCATCAAAACGCACAGGGCACAGGCCAGGCCGCAGATTACAAGTCTACACAGGCCGCCATTGATCAGATTACAGGTAAACTGAACAGG CTAGTGGAAAAGACCAATACCGAGTTCGAATCCATCGAATCTGAGTTCTCAGAGATAGAGCACCAGATCGGGAACGTTATCAACTGGACCAA GGACAGCATTACCGACATTTGGACCTACCAGGCAGAGCTCCTGGTCGCCATGGAAAATCAGCATACCATCGACATGGCCGATTCAGAGATG CTTAACCTTTATGAGCGGGTGAGGAAACAGCTGAGGCAGAATGCAGAAGAAGATGGTAAGGGTTGCTTCGAAATTTACCATGCTTGTGATG ATAGTTGTATGGAGAGTATCCGTAACAATACTTACGATCACAGCCAGTACAGGGAAGAAGCACTCCTCAATCGCCTCAACATCAATCCTGTA ACTCTCTCGTCTGGTTACAAAGATATCATCTTATGGTTCTCTTTTGGGGCCTCATGTTTCGTGCTCCTGGCCGTGGTGATGGGCCTGTTCTTT TTCTGCTTAAAGAATGGAAATATGAGGTGCACAATCTGCATT

2911	ATGTACAAGATAGTCGTTATCATCGCCTTACTCGGAGCAGTTAAAGGTTTAGATAAGATCTGCCTGGGGCACCACGCCGTCGCTAACGGCA CTATTGTGAAGACGCTGACCAATGAACAAGAGGAAGTCACTAACGCAACAGAAACCGTGGAGAGCACTGGCATCAATCGCCTCTGTATGAA AGGGCGCAAGCATAAGGACCTCGGCAATTGCCACCCTATAGGCATGCTGATTGGAACGCCTGCTTGTGACCTTCACCTGACAGGGATGTG GGATACCCTAATTGAACGAGAGAACGCGATCGCATACTGCTATCCTGGTGCCACTGTGAACGTGGAAGCCTTACGCCAAAAAATCATGGAG AGCGGAGGGATCAATAAAATCAGTACAGGTTTCACCTACGGATCCTCCATCAACAGTGCCGGTACTACACGCGCCTGTATGAGAAACGGAG GCAACAGCTTCTATGCTGAGCTGAAATGGTTAGTGAGCAAATCCAAAGGTCAGAATTTTCCCCAGACCACAAATACTTACCGAAATACGGAC ACCGCCGAGCATCTGATAATGTGGGGCATCCACCATCCTAGTTCTACTCAGGAGAAAAACGACCTTTATGGAACGCAGAGCCTGTCTATAA GCGTGGGGTCTTCAACATACAGGAATAACTTTGTCCCTGTAGTGGGCGCCGGTCCTCAGGTGAATGGGCAAAGCGGCCGCATCGATTTTCA CTGGACCCTGGTTCAACCTGGAGACAATATTACGTTCAGTCACAACGGCGGCCTTATTGCCCCTTCCAGAGTCTCTAAGCTCATTGGCAGG GGACTCGGTATCCAATCCGACGCCCCTATTGACAACAACTGCGAGTCCAAATGCTTCTGGAGGGGAGGAAGCATCAACACCCGCCTGCCAT TTCAGAATCTTTCACCTCGAACCGTGGGCCAGTGTCCTAAATACGTCAATCGGAGAAGCTTGATGTTGGCCACTGGGATGCGGAACGTACC CGAGCTTATCCAAGGCAGAGGCCTTTTCGGTGCCATCGCAGGGTTCTTGGAAAATGGCTGGGAAGGCATGGTCGATGGCTGGTATGGCTT TCGCCATCAAAATGCTCAGGGCACGGGCCAGGCCGCTGACTACAAAAGTACACAAGCTGCCATCGATCAGATTACAGGCAAGCTCAATAGG TTGGTGGAGAAGACAAACACCGAGTTTGAGTCAATCGAGAGCGAGTTTAGCGAGATAGAGCATCAGATCGGCAACGTGATTAACTGGACTA AGGACTCTATAACAGACATTTGGACGTACCAGGCAGAGCTCCTTGTAGCAATGGAAAATCAGCACACAATTGACATGGCTGACTCTGAAATG CTCAACCTGTATGAGCGGGTTCGGAAGCAGCTGAGGCAGAATGCAGAGGAAGATGGTAAGGGCTGCTTTGAAATCTACCATGCCTGTGATG ACAGTTGCATGGAGAGTATTCGAAATAACACATACGATCATTCGCAGTACAGAGAGGAAGCCTTGTTGAACCGCCTGAATATCAACCCCGTG ACATTATCCTCAGGCTATAAAGACATCATCCTCTGGTTTTCGTTTGGGGCCTCGTGCTTTGTCTTGCTGGCTGTTGTCATGGGGCTGTTTTTC TTTTGTTTAAAGAACGGGAACATGCGATGCACAATCTGTATT
2912	ATGTACAAGATTGTGCTGGTACTTGCACTTCTGGGTGCCGTTCACGGCCTCGACAAAATCTGCCTAGGCCACCACGCTGTGCCTAACGGGA CCATTGTGAAAACCCTGACTAACGAGAAGGAGGAGGTTACCAACGCAACCGAGACTGTAGAGTCTAAGTCCTTGGATAAACTGTGCATGAA GAACCGAAACTACAAAGACCTGGGCAATTGCCATCCTATCGGGATGGTCGTCGGAACTCCAGCCTGCGATCTCCACCTAACTGGCACCTGG GACACCCTCATCGAGAGGGACAATTCAATTGCATACTGCTATCCTGGGGCTACCGTGAGTGAAGAAGCACTGCGACAAAAAATCATGGAGT CAGGAGGGATCGACAAAATATCGACGGGATTCACATACGGCTCATCTATTAATTCAGCAGGAACCACGAAAGCTTGTATGCGCAACGGAGG TAACTCCTTCTATAGCGAACTCAAGTGGTTAGTGTCTAAGAATAAGGGACAGAACTTTCCACAGACAACAAACACTTACCGGAACACAGACA GCGTAGAACACCTCATAATATGGGGGATCCACCACCCAAGCTCCACACAGGAGAAGAACGATTTATACGGCACTCAGTCCCTGAGCATCAG CGTCGGATCAAGCACCTACCAGAACAACTTCGTCCCCGTGGTGGGAGCTCGCCCACAGGTGAATGGCCAGAGCGGACGCATCGATTTCCA CTGGACTATGGTCCAGCCGGGTGACAACATCACCTTCTCTCACAATGGGGGGCTGATTGCTCCGAACCGAGTGTCCAAGCTGAAAGGGAG GGGGTTAGGAATCCAGAGCGGCGCCTCGGTAGACAACGACTGCGAGAGCAAATGTTTCTGGAAAGGTGGCTCGATTAATACCAAGCTTCC ATTTCAGAATTTATCTCCTAGAACTGTCGGCCAGTGTCCAAAATACGTCAACAAAAAGTCTCTGCTCCTCGCAACAGGCATGCGGAATGTTC CCGAAGTTGCACAGGGCCGAGGCCTGTTTGGTGCCATTGCCGGGTTTATCGAGAACGGATGGGAGGGCATGGTGGATGGATGGTACGGA TTCCGACACCAGAATGCGCAGGGGACGGGCCAAGCAGCGGATTACAAGTCCACACAAGCAGCAATTGACCAAATCACTGGCAAGTTGAAT CGCCTCATAGAGAAAACTAATACAGAGTTCGAATCTATCGAAAGTGAGTTTTCCGAGATCGAACACCAGATCGGGAATGTGATCAACTGGAC GAAGGATTCTATTACTGACATATGGACCTACCAGGCCGAGCTCCTGGTGGCTATGGAAAACCAACATACAATTGACATGGCTGACTCTGAAA TGCTGAATCTGTACGAACGAGTGAGGAAGCAGCTCCGCCAAAACGCAGAGGAAGACGGAAAGGGTTGTTTTGAGATATATCACAAGTGCGA CGACAACTGCATGGAGTCCATCCGAAACAATACTTACGACCATACACAATACAGAGAGGAGGCCCTGCTGAACCGCCTGAATATCAATCCC GTGAAGCTCAGTTCCGGTTACAAGGACGTCATTTTATGGTTCAGCTTTGGAGCGTCCTGCTTCGTGCTTCTGGCCGTCATCATGGGCCTTGT CTTCTTTTGTCTGAAAAACGGTAATATGAGATGCACCATCTGCATT
2913	ATGTATAAAATTGTCGTGATCATCGCCCTGCTCGGGGCCGTCAAGGGATTGGATAAGATATGTCTCGGTCATCATGCAGTGGCTAACGGCA CGATCGTGAAGACCCTGACCAACGAGCAGGAGGAAGTGACTAATGCCACCGAAACCGTTGAGTCTACTGGTATCAATAGGCTGTGCATGAA AGGGACAAAGCACAAGGACCTCGGCAATTGTCACCCAATCGGGATGCTCATTGGGACACCCGCATGTGACCTCCACCTGACAGGCATGTG GGATACACTGATTGAGCGCGAAAATGCCATCGCGTACTGCTACCCTGGCGCCACCGTGAATGTGGAAGCTCTGCGACAGAAAATAATGGAG AGTGGGGGGATTAATAAGATCTCTACAGGGTTTACGTACGGATCAAGTATTAACTCCGCCGGGACTACTCGAGCATGTATGCGCAATGGCG GCAATTCCTTTTACGCCGAGCTGAAGTGGCTTGTGTCGAAGTCAAAAGGACAGAATTTTCCCCAGACAACAAACACATATCGCAATACCGAT ACTGCAGAACACTTGATTATGTGGGGGATTCACCACCCGTCATCTACTCAGGAGAAGAACGATCTGTATGGCACTCAGAGCCTGAGTATCTC AGTAGGCAGCAGCACCTACAGAAATAACTTCGTGCCGGTTGTCGGGGCCCGCCCACAAGTGAATGGCCAGTCAGGGCGGATCGACTTTCA CTGGACGCTGGTTCAGCCTGGAGATAACATTACGTTTAGTCACAATGGCGGGCTCATCGCGCCAAGTCGGGTATCCAAACTGATCGGCCGA GGACTTGGGATACAATCGGACGCCCCCATCGATAACAATTGCGAATCCAAATGTTTTTGGAGAGGAGGATCCATAAATACACGACTGCCCTT TCAGAACCTGTCTCCTCGGACAGTTGGGCAGTGTCCAAAGTACGTAAACCGGAGGTCTCTCATGCTCGCGACAGGAATGCGTAATGTGCCA GAACTGATTCAGGGCCGCGGCCTCTTCGGTGCGATCGCGGGGTTCCTCGAGAATGGTTGGGAAGGCATGGTGGACGGGTGGTATGGTTTT CGTCATCAGAACGCTCAAGGCACTGGGCAGGCCGCCGATTACAAGTCCACGCAGGCGGCAATTGACCAAATCACCGGGAAGTTAAACCGG CTTGTCGAGAAAACCAATACCGAGTTTGAATCCATAGAGTCCGAGTTTTCCGAGATCGAGCACCAAATCGGAAACGTTATAAATTGGACCAA GGACTCTATCACCGACATCTGGACCTACCAGGCCGAACTCCTGGTCGCCATGGAAAATCAGCACACAATAGACATGGCAGATTCAGAGATG CTGAACCTGTACGAGCGGGTGCGTAAGCAGCTCCGGCAGAATGCGGAAGAGGACGCGCAAGGGATGTTTCGAAATATACCACGCTTGTGAT GATTCATGTATGGAGTCGATCAGGAACAATACATACGACCATTCCCAGTACAGGGAAGAGGCCCTCCTGAACAGGCTTAACATCAACCCCG TCACCCTGTCCAGCGGCTACAAGGATATTATCCTTTGGTTTTCATTCGGCGCAAGCTGCTTCGTCTTACTGGCCGTGGTAATGGGACTATTC TTTTTTTGCCTGAAAAACGGCAACATGAGGTGTACCATTTGTATT

2914	ATGTACAAAATCGTGCTCGTTCTGGCCCTGCTCGGCGCCGTGCACGGCCTCGACAAGATTTGCCTGGGACATCACGCTGTGCCCAACGGT ACTATTGTCAAGACACTGACGAACGAAAAAGAAGAGGTTACAAACGCTACCGAAACCGTTGAATCTAAGAGCTTGGATAAACTGTGCATGAA AAATCGCAACTATAAAGACCTCGGTAATTGTCACCCTATAGGGATGGTGGTCGGCACCCCAGCTTGCGACCTGCATTTGACAGGTACTTGG GATACTTTGATCGAACGGGATAACAGTATCGCCTACTGTTATCCGGGAGCCACGGTGTCGGAAGAGGCTCTGCGTCAGAAAATCATGGAGA GCGGCGGGATAGACAAGATCTCCACTGGCTTCACTTATGGCTCCAGCATTAATAGCGCAGGTACAACCAAGGCTTGCATGAGAAATGGGGG GAACTCCTTCTACTCAGAACTGAAATGGCTGGTGTCCAAAAATAAGGGACAGAATTTCCCGCAGACGACAAACACCTATAGAAACACAGACT CTGTAGAGCACCTCATTATTTGGGGTATCCATCACCCAAGCTCCACCCAGGAAAAGAATGACCTGTACGGCACACAGTCCTTGAGCATTTCA GTGGGCAGCTCCACATATCAGAATAACTTTGTGCCAGTCGTTGGCGCACGACCCCAGGTGAATGGGCAGTCAGGTCGCATTGACTTCCACT GGACTATGGTGCAGCCCGGCGATAACATTACCTTCTCCCATAACGGGGGACTGATAGCCCCCAATAGGGTCTCTAAGCTCAAGGGTCGGG GTTTGGGCATCCAGTCCGGAGCATCCGTCGATAACGACTGTGAGTCGAAATGCTTTTGGAAAGGTGGATCCATCAATACAAAGCTACCTTTC CAAAATTTATCCCCTCGAACGGTCGGACAGTGCCCTAAATACGTAAACAAGAAGTCACTGCTCCTCGCTACCGGCATGCGCAATGTACCTGA AGTTGCCCAAGGACGGGGGCTTTTCGGCGCAATCGCTGGCTTCATTGAGAATGGATGGGAGGGCATGGTTGACGGCTGGTACGGCTTTAG GCACCAGAATGCTCAGGGTACCGGTCAGGCAGCAGATTATAAAAGCACCCAGGCTGCCATCGATCAGATAACCGGGAAGTTGAACAGGCT GATCGAGAAGACCAACACTGAGTTTGAGAGCATTGAATCGGAGTTCTCGGAAATCGAACATCAAATCGGCAATGTGATCAATTGGACTAAGG ATTCTATCACCGACATCTGGACGTACCAGGCAGAACTGTTAGTTGCAATGGAAAATCAGCATACTATTGATATGGCAGACAGTGAGATGCTG AACCTGTACGAGCGTGTTAGAAAGCAGCTCAGACAGAACGCCGAGGAGGACGGGAAAGGATGCTTTGAAATCTATCACAAATGTGATGATA ATTGCATGGAGTCAATCAGGAATAATACATACGATCACACGCAATACAGAGAGGAGGCTCTCCTGAACCGCCTTAATATCAACCCCGTGAAA CTCAGCTCCGGATATAAAGACGTCATCCTCTGGTTTTCTTTTGGAGCAAGCTGCTTCGTGCTGCTCGCCGTTATCATGGGGTTGGTGTTCTT CTGCCTGAAGAATGGTAACATGAGATGCACCATTTGTATA
2915	ATGTACAAGATTGTGCTCGTTCTGGCACTATTGGGCGCAGTGCATGGTCTGCATAAAATCTGTCTGGGTCATCACGCAGTGCCCAACGGTA CTATCGTCAAAACACTAACGAACGAGAAGGAAGAAGTAACTAATGCAACGGAGACAGTGGAGTCAAAAAGTCTGGACAAGTTATGTATGAAG AACAGAAACTATAAGGATCTCGGTAATTGCCACCCCATCGGTATGGTGGTTGGGACCCCAGCCTGCGACCTTCACCTGACGGGAACCTGG GACACCCTCATCGAGCGGGACAACTCCATTGCCTATTGCTATCCTGGGGCCACCGTCTCCGAGGAGGCACTGAGGCAGAAAATTATGGAAT CCGGCGGGATTGACAAGATTTCTACCGGCTTTACCTATGGCTCCTCAATCAACAGCGCTGGGACAACTAAAGCCTGCATGAGGAACGGAGG AAATTCTTTTTACTCTGAGTTGAAGTGGTTGGTGAGCAAAAATAAGGGCCAAAATTTTCCTCAGACGACCAATACCTACCGTAATACTGACTC AGTTGAGCACCTCATCATTTGGGGAATTCATCATCCCAGCAGCACTCAGGAGAAGAATGATCTGTACGGGACCCAATCGTTGAGTATATCCG TGGGATCTAGTACTTATCAGAACAACTTTGTACCTGTAGTGGGTGCCCGCCCACAGGTAAACGGACAGTCGGGGCGTATTGACTTTCATTG GACAATGGTGCAGCCCGGGGATAATATCACTTTTAGCCATAATGGAGGCCTGATCGCACCAAACCGAGTGAGCAAGCTGAAGGGTAGAGG ATTGGGAATCCAGTCTGGAGCCAGCGTTGATAACGACTGTGAGTCTAAATGCTTCTGGAAAGGGGGTAGCATAAATACCAAGCTGCCCTTC CAAAACCTGAGCCCTCGGACCGTGGGCCAGTGCCCAAAGTACGTGAATAAGAAAAGCCTCCTGCTCGCTACAGGCATGAGAAATGTGCCC GAGGTTGCACAGGGCAGAGGGCTCTTCGGCGCTATAGCTGGATTCATTGAAAATGGCTGGGAAGGTATGGTAGATGGTTGGTACGGCTTC AGACACCAGAATGCTCAGGGAACCGGACAGGCTGCTGATTATAAATCTACCCAGGCTGCTATCGATCAGATCACTGGGAAATTGAACCGGC TGATCGAAAAAACAAACACCGAATTCGAGAGTATTGAATCCGAGTTTAGCGAAATTGAACATCAGATCGGCAATGTGATAAACTGGACTAAG GATAGTATAACTGACATCTGGACATACCAAGCTGAGCTCCTGGTGGCCATGGAGAATCAACACACGATCGACATGGCAGACTCTGAAATGTT GAACCTGTACGAGAGGGTGCGGAAGCAGCTCAGACAAAACGCTGAGGAGGACGGCAAAGGTTGCTTCGAAATCTACCATAAGTGCGACGA TAACTGTATCGACAGTATTAGAAACAATACCTACGACCACACGCACTATAGGGAGGACGCTCTGTTGAATACGTTAAATATCAACCCTGTGA AACTATCCAGCGGCTATAAGGACGTGATTTTATGGTTCTCGTTTGGTGCCAGCTGTTTTGTCCTACTGGCAGTGATCATGGGTCTTGTTTTTT TTTGCCTGAAGAACGGCAACATGAGATGCACGATATGTATT
2916	ATGTACAAGATCGTTGTGATTATCGCCCTGCTCGGCGCCGTCAAAGGATTAGACAAGATCTGTCTAGGCCATCACGCCGTTGCGAACGGAA CCATCGTGAAGACCCTGACAAATGAGCAGGAAGAAGTTACCAATGCCACTGAGACCGTAGAAAGCACCGGGATCAACAGACTATGTATGAA AGGCCGCAAGCATAAGGATCTGGGAAACTGCCATCCTATCGGAATGCTCATTGGAACTCCCGCCTGTGACCTGCACCTGACCGCCATGTG GGACACACTGATTGAGCGGGAGAACGCCATCGCTTACTGTTATCCTGGTGCAACTGTGAACGTCGAAGCTTTGCGGCAGAAAATAATGGAA TCGGGGGGGATTAATAAAATAAGTACCGGATTCACATATGGAAGCTCAATTAACTCCGCAGGAACTACAAGAGCATGTATGCGAAACGGCG GCAACTCATTCTACGCTGAACTTAAGTGGCTGGTGTCTAAGTCAAAGGGGCAGAACTTTCCGCAAACCACTAATACATATCGCAATACAGAC ACAGCCGAGCACCTCATCATGTGGGGAATTCACCACCCAAGTAGTACCCAGGAGAAGAACGACCTGTACGGCACTCAGTCGCTTAGCATAA GTGTTGGCTCATCGACTTATCGGAATAACTTCGTCCCAGTGGTTGGTGCTCGTCCCCAAGTGAATGGTCAGTCCGGCAGAATTGATTTTCAT TGGACACTGGTGCAACCCGGTGACAACATCACTTTTAGCCATAATGGTGGTCTAATCGCCCCGTCCCGGGTGTCCAAGTTAATTGGGAGAG GTCTGGGTATCCAGAGTGATGCTCCAATCGATAATAATTGCGAGAGTAAGTGCTTCTGGCGCGGCGGAAGTATCAACACCCGGCTCCCTTT CCAGAATCTATCACCTAGAACTGTTGGGCAGTGCCCTAAATATGTTAATCGCCGCTCACTTATGCTGGCAACCGGGATGCGAAACGTTCCAG AACTGATCCAAGGGCGGGGGCTTTTCGGGGCTATTGCCGGGTTTTTGGAGAACGGCTGGGAAGGGATGGTCGATGGTTGGTATGGCTTCC GCCATCAGAATGCACAGGGTACCGGCCAGGCCGCCGACTACAAGTCAACGCAGGCTGCAATTGACCAAATCACTGGGAAACTGAACCGGC TCGTTGAAAAGACTAACACTGAGTTTGAAAGCATAGAAAGTGAATTCAGTGAAATTGAGCATCAAATCGGCAACGTGATCAATTGGACCAAG GACTCAATCACGGATATCTGGACATACCAGGCCGAGCTGCTCGTTGCGATGGAGAACCAACACACCATCGATATGGCAGACTCTGAGATGC TAAATCTCTACGAAAGAGTGAGAAAGCAGTTGCGACAAAATGCGGAGGAGGATGCTAAGGGTTGCTTCGAAATTTACCATGCCTGCGACGA CAGTTGCATGGAATCCATTAGGAACAATACCTACGATCATTCGCAGTATAGGGAGGAGGCCCTCTTAAACCGCCTGAACATCAACCCAGTAA CCTTGAGCAGCGGGTATAAAGATATAATCCTGTGGTTTAGCTTCGGTGCCTCCTGCTTTGTTCTGCTGGCCGTAGTCATGGGTCTGTTTTTTT TCTGCCTGAAGAATGGTAATATGCGCTGCACCATTTGCATT

2917	ATGTACAAGATCATTGTAATCATCGCCTTGTTGGGAGCCGTCAAAGGGCTCGATAAAATTTGCTTGGGCCATCATGCAGTCGCCAACGGGA CAATAGTCAAGACGCTCACGAATGAACAGGAGGAAGTGACAAACGCCACAGAAACCGTTGAAAGTACAGGGATTAATAGGCTGTGTATGAA AGGGCGCAAACATAAGGATTTGGGCAATTGTCACCCAATAGGGATGCTTATTGGGACACCAGCTTGTGATCTCCATCTCACTGGAACATGG GACACCCTAATCGAGCGCGAAAACGCTATTGCATATTGCTATCCGGGAGCGACGGTTAACGTGGAGGCACTCCGACAGAAAATCATGGAGA GCGGGGGGATCCACAAGATTTCCACAGGCTTTACTTACGGCTCTAGTATAAATTCCGCCGGGACCACAAGGGCCTGTATCAGAAACGGTG GCAACAGTTTTTACGCTGAGTTAAAATGGCTCGTTAGCAAGAGCAAGGGTCAGAACTTTCCGCAGACCACAAACACATACCGGAACACCGAT ACAGCAGAGCATCTAATCATGTGGGGCATACATCATCCTTCCAGTACCCAGGAGAAGAACGATTTATACGGGACCCAATCCCTGAGCATCA GCGTTGGGTCATCTACTTACCGCAATAATTTTGTCCCGGTTGTGGGAGCAAGGCCCCAAGTGAATGGTCAGAGTGGCAGGATTGACTTTCA CTGGACACTGGTGCAACCTGGTGACAACATTACCTTCAGCCATAACGGAGGACTGATCGCGCCTTCACGGGTGAGCAAGCTTATTGGTAGA GGACTCGGGATCCAAAGTGATGCCCCAATCGACAATAACTGTGAGAGCAAATGCTTTTGGCGAGGGGGATCCATCAATACACGCCTCCCCT TTCAAAACCTGAGCCCACGCACTGTGGGGCAATGCCCTAAGTACGTAAACCGCAGGTCTCTAATGCTTGCAACCGGGATGAGAAACGTTCC AGAATTAATTCAGGGACGGGGACTGTTTGGGGCGATCGCCGGCTTTCTGGAAAATGGCTGGGAAGGGATGGTGGATGGCTGGTATGGCTT CCGGCACCAAAATGCCCAGGGCACAGGGCAGGCTGCAGATTACAAATCAACTCAGGCTGCCATCGACCAAATCACAGGCAAGCTGAATAG GCTCGTGGAGAAAACGAACACGGAATTCGAGTCTATTGAATCTGAGTTCTCCGAAATCGAGCACCAGATCGGGAATGTAATTAACTGGACCA AGGACTCAATCACTGACATTTGGACCTATCAGGCAGAATTACTGGTAGCGATGGAGAACCAACACACAATTGACATGGCAGATTCTGAAATG CTCAACCTCTATGAACGAGTGAGGAAACAGCTCCGGCAGAACGCCGAGGAGGATGGAAAGGGATGTTTCGAGATCTATCACGCATGTGAC GATAGCTGTATGGAATCTATACGGAATAACACATACGATCATTCTCAATATCGGGAGGAGGCTCTGCTTAACCGACTCAACATAAACCCAGT CACTCTATCCAGTGGCTACAAGGATATTATCCTGTGGTTTAGCTTTGGCGCCTCGTGCTTCGTGCTTCTCGCTGTTGTCATGGGCCTGTTCT TCTTCTGTCTGAAGAACGGGAATATGCGGTGCACCATTTGCATC
2918	ATGTATAAAATCGTGGTTATTATCGCATTGCTGGGTGCTGTGAAAGGCCTCGATAAAATTTGCCTCGGGCATCACGCCGTGGCCAACGGCA CAATTGTGAAGACACTGACAAACGAACAGGAAGAAGTTACCAACGCTACTGAGACCGTGGAATCCACAGGTATTAATAGGCTTTGTATGAAG GGTCGGAAGCACAAAGACCTGGGGAATTGCCACCCTATCGGAATGCTCATTGGAACTCCGGCCTGTGACCTGCACCTCACAGGTATGTGG GATACTCTGATTGAGCGAGAGAACGCGATTGCCTACTGTTACCCGGGAGCAACCGTGAACGTGGAGGCGCTCCGGCAGAAGATCATGGAG TCTGGCGGCATTAACAAGATCAGTACCGGGTTCACATATGGAAGTTCAATCAATAGTGCCGGCACTACCAGGGCATGTATGCGCAACGGAG GAAATAGCTTCTACGCGGAATTGAAATGGTTAGTCAGCAAGAGCAAGGGACAGAACTTTCCTCAGACTACGAACACTTACCGTAATACTGAT ACTGCCGAACATCTAATTATGTGGGGGATCCATCACCCATCCAGCACCCAGGAAAAGAACGATTTGTACGGCACCCAGTCTTTATCTATTTC CGTGGGGAGTTCGACATATAGAAATAACTTTGTGCCCGTTGTCGGTGCTGGGCCCCAAGTTAACGGCCAATCGGGACGTATCGATTTCCAC TGGACACTGGTTCAGCCAGGGGACAATATCACCTTTTCTCACAACGGAGGTCTTATAGCTCCATCTCGTGTGTCTAAACTCATTGGACGGGG GCTTGGGATCCAAAGCGATGCCCCCATCGACAATAATTGCGAAAGTAAGTGCTTCTGGCGGGGAGGTTCTATTAATACTAGGCTTCCTTTCC AGAACCTCTCACCACGAACAGTCGGCCAATGCCCCAAGTACGTTAATCGACGTTCACTCATGCTGGCCACGGGAATGCGCAACGTGCCCG AGCTTATCCAGGGGCGCGGATTGTTCGGCGCCATCGCTGGCTTCTTGGAGAACGGTTGGGAGGGTATGGTCGATGGGTGGTACGGCTTTC GCCACCAAAACGCTCAAGGGACAGGACAGGCCGCCGATTATAAGAGCACTCAGGCCGCCATAGATCAGATTACGGGCAAGCTAAACAGGC TGGTCGAGAAAACCAATACGGAATTCGAGTCCATTGAATCAGAGTTCTCTGAGATCGAGCACCAGATCGGCAACGTAATCAACTGGACCAAA GATTCTATCACCGATATTTGGACATATCAGGCAGAACTACTGGTCGCCATGGAAAACCAGCATACCATCGATATGGCGGACAGTGAGATGCT TAATCTCTACGAAAGAGTACGAAAGCAGCTGCGACAAAACGCCGAGGAAGACGGTAAAGGCTGCTTCGAAATTTATCACGCGTGCGACGAC TCTTGTATGGAATCTATACCAAACAACACATACGATCACTCCCAATACCGGGAGGAGGCTCTCCTCAATAGACTGAATATCAACCCAGTCAC TTTGAGTAGTGGGTATAAGGACATTATTCTTTGGTTCTCCTTCGGGGCTTCTTGTTTTGTCCTTCTGGCAGTGGTTATGGGGCTCTTCTTTTTT TGCCTCAAGAACGGCAACATGCGCTGTACTATATGTATT
2919	ATGTATAAAGTCGTTGTGATCATCGCTTTGTTGGGAGCTGTCCGGGGTTTGGATAAGATCTGCCTCGGCCATCACGCAGTAGCAAATGGAA CCACTGTGAAGACCCTAACTAACGAACAAGAGGAAGTCACAAATGCCACTGAGACTGTCGAGTCTACTAGCCTGAATAAACTGTGCATGAAG GGGAGGAGGTACAAAGATCTTGGAAATTGCCACCCAATCGGAATGCTGATAGGCACCCCTGTGTGTGATCTCCATCTGACCGGAACATGGG ATACCCTGATTGAGAGAGAGAATGCTACTGCATACTGCTACCCTGGAGTCACTATCAATGAAGAGGCCCTGCGTCAAAAGATCATGGAGTC CGGGGGTATTTCTAAGATGCGCACCGGCTTTACCTACGGACCATCTATCAATTCGGCAGGTACTACCCGGTCATGTATGCGCAATGGAGGC AACTCCTTCTATGCCGAACTGAAGTGGCTGGTCTCGGGCACTAAGGGTCAGAACTTCCCGCAGACTACTAATACATACCGAAATACCGACA CTGCCGAGCACCTGATTATCTGGGGAATTCACCATCCCTCGTCCACTCAAGAGAAGAATGACCTTTATGGCACACAGAGCCTCTCGATCAG CGTGGGCTCCTCGACTTACCAGAATAACTTTGTCCCGGTAATTGGCGCTAGGCCTCAGGTGAACGGCCAGTCCGGCAGAATCGAGTTCCAT TGGACTCTGGTGCGCCCAGGCGACAATATCACATTCAGCCATAATGGGGGTCTGATCGCACCTGACCGAGTTAGTAAACTGATTGGCAAGG GGATCGGGATCCAATCAGGGGCTGTGATCGACAAAGACTGTGAGTCAAAGTGCTTCTGGAGGGGCGGAAGCATAATTACGGAGCTGCCTT TTCAGAACCTGAGCCCAAGAACTGTTGGACAATGCCCTAAGTACGTTAAGAAGCGATCTCTCTTGTTGGCCACTGGAATGAGGAACGTGCC GGAAGTGGTTCAGGGGCGCGGCCTCTTCGGGGCTATAGCCGGATTCATAGAAAACGGCTGGGAGGGGATGGTGGATGGCTGGTACGGCT TCAGACACCAAAATGCGCAGGGCATCGGCCAGGCTGCTGATTACAAGTCTACCCAAACTGCGATTGATCAGATCACCGGAAAACTGAATCG ACTGATCGAAAAAACTAACACTGAGTTCGAAAGCATCGAAAGTGAATTTAGCGAGATTGAACATCAGATCGGCAATGTCATAAATTGGACCA AGGATTCCATTACAGATATTTGGACATACCAGGCCGAGCTCCTCGTGGCAATGGAAAATCAGCATACTATTGATATGGCCGATTCGGAAATG CTGAACCTCTACGAACGTGTCCGGAAACAGCTGCGCCAGAACGCTGAGGAGGACGGAAAGGGATGCTTTGAGATTTACCACACCTGCGAC AATTCTTGTATGGAGAGTATACGCAATAATACATATGACCATTCACAATACCGCGAAGAAGCCCTTCTGAATCGTCTCAATATCAATCCCGTC AAACTGTCCTCAGGGTATAAAGACATTATCCTATGGTTCAGTTTTGGTGCCAGCTGCTTTGTGCTGCTGGCTGTCATCATGGGGCTGGGCTT TTTTTGTCTAAAAAACGGAAACATGCGGTGCACGATCTGCATT

2920	ATGTATAAGATCGTAGTGATCATTGCGCTGCTTGGCGCTGTCAAGGGCCTAGATAAGATATGTCTCGGACACCATGCTGTTGCCAATGGGA CAATTGTGAAGACTCTAACTAACGAACAGGAAGAAGTTACGAATGCTACAGAAACTGTGGAGTCCACTGGAATCAATAGACTCTGTATGAAA GGCCGGAAACACAAGGATCTGGGCAACTGCCACCCCATTGGTATGCTTATAGGAACACCTGCGTGCGATCTGCACCTCACTGGAATGTGG GATACCCTCATCGAGCGGGAGAACGCAATCGCGTACTGTTACCCCGGAGCTACGGTTAACGTCGAAGCTCTTCGGCAGAAGATCATGGAAT CCGGGGGGATCAATAAGATCTCTACTGGATTCACATATGGGTCTTCAATTAACTCCGCTGGGACGACCCGTGCCTGCATGAGGAACGGCGG TAACTCTTTCTATGCTGAGCTGAAGTGGCTGGTCAGTAAAAGTAAAGGACAGAACTTTCCGCAGACCACCAACACGTATCGGAACACTGATA CAGCAGAGCACCTTATCATGTGGGGAATCCACCATCCCTCTTCTACACAAGAAAAGAATGACCTGTACGGCACTCAGAGCCTTTCAATCTCA GTGGGCAGTTCCACCTATCGAAACAACTTCGTGCCCGTGGTTGGAGCACGCCCACAGGTGAACGGCCAATCCGGGCGCATTGATTTTCATT GGACCCTGGTTCAGCCAGGGGATAACATAACTTTCAGTCATAATGGTGGGCTCATCGCACCAAGCCGTGTTTCCAAGTTAATCGGCCGGGG GCTGGGCATCCAGAGCGACGCTCCTATCGACAATAACTGCGAAAGTAAATGCTTCTGGCGCGGAGGTAGCATCAACACCCGACTACCCTTC CAGAATCTTAGCCCACGAACCGTGGGACAATGCCCAAAATACGTCAACCGCAGGTCTCTTATGCTAGCTACCGGTATGCGTAATGTCCCTG AGCTGATTCAGGGCCGCGGGCTTTTTGGGGCCATTGCAGGATTTCTTGAGAATGGATGGGAAGGTATGGTTGACGGATGGTATGGCTTTAG ACACCAGAACGCGCAGGGCACTGGTCAAGCTGCAGATTACAAGTCCACACAGGCTGCTATAGACCAGATAACAGGCAAGCTGAATAGATTA GTGGAGAAAACCAACACAGAATTCGAGAGTATTGAGTCTGAATTTAGCGAGATTGAGCATCAGATTGGGAACGTGATCAATTGGACTAAAGA CAGCATCACGGACATCTGGACATACCAGGCTGAACTTTTGGTGGCCATGGAGAATCAGCACACAATCGATATGGCAGACTCAGAGATGCTC AATTTGTATGAACGGGTGAGAAAGCAATTACGCCAGAACGCGGAAGAGGATGGCAAAGGCTGCTTTGAGATATACCACGCATGTGACGACT CATGCATGGAATCCATCAGGAACAACACCTATGACCATTCCCAGTACCGGGAGGAAGCTCTGTTGAACCGCCTCAACATAAACCCGGTGAC ACTGTCTAGCGGTTATAAAGATATCATCCTTTGGTTCTCATTCGGGGCGTCTTGTTTTGTTCTCCTTGCAGTGGTTATGGGCCTATTCTTCTT CTGCCTGAAGAATGGAAATATGAGATGTACAATCTGTATC
2921	ATGTATAAAGTTGTTGTTATCATAGCGCTGCTAGGCGCTGTCAAGGGCTTAGACAAAATTTGTCTGGGCCATCACGCAGTGGCTAACGGCAC AATAGTTAAGACATTGACCAATGAACAGGAGGAGGTTACAAATGCAACCGAGACCGTGGAATCAACCGGCATTAATAGACTATGTATGAAGG GCCGGAAGCATAAGGATCTGGGGAACTGCCATCCAATCGGTATGCTGATCGGCACGCCAGCTTGCGATTTGCACCTAACGGGAACCTGGG ACACTTTAATCGAGAGGCAGAATGCTATAGCTTATTGCTACCCCGGCGCTACCGTTAATGTTGAGGCACTTCGGCAGAAGATCATGGAAAG CGGGGGGATCGATAAAATCAGCACAGGATTCACCTACGGGAGTTCTATCAACTCTGCCGGGACTACACGTGCATGTATGAGAAACGGCGGT AATTCCTTCTATGCCGAGCTCAAATGGCTTGTTTCCAAGAATAAGGGCCAGAACTTCCCACAGACCACTAATACTTACAGGAATACCGACAC TGCGGAGCACCTCATTATGTGGGGCATACATCACCCTTCTTCGATCCAAGAAAAGAATGACCTCTATGGTACTCAGTCCCTCAGCATATCTG TGGGGAGCAGTACTTACCGGAATAATTTCGTACCCGTTGTGGGAGCACGCCCACAGGTGAACGGCCAGTCTGGTAGAATTGATTTCCATTG GACTTTGGTGCAGCCGGGGGATAACATCACCTTTTCCCATAACGGGGGACTGATCGCTCCAAGCAGAGTTTCCAAACTTATTGGGAGAGGT TTAGGGATCCAGTCAGACGCCCCAATTGATAACAACTGTGAGTCGAAGTGCTTTTGGAGAGGGGGCAGCATCAATACGCGGCTGCCTTTCC AGAACCTGTCACCCAGGACTGTCGGCCAGTGCCCAAAGTACGTGAACAGACGATCCTTGATGCTTGCCACCGGAATGCGTAATGTGCCCG AGCTGATACAGGGCCGTGGACTTTTCGGAGCCATCGCGGGATTTCTGGAGAATGGTTGGGAAGGAATGGTCGACGGCTGGTATGGATTCC GCCATCAAAACGCCCAAGGAACAGGCCAGGCCGCTGACTATAAAAGCACGCAGGCTGCTATTGACCAGATCACAGGTAAGCTGAACCGCC TGGTCGAAAAGACCAACACTGAATTTGAGTCCATCGAGAGCGAGTTTAGCGAAATCGAACACCAAATCGGCAATGTAATCAACTGGACCATG GACAGCATAACCGATATTTGGACTTACCAGGCGGAATTACTGGTCGCCATGGAAAATCAGCACACAATCGATATGGCAGACAGCGAAATGTT GAACCTCTATGAGAGAGTCAGAAAACAATTGCGGCAGAATGCTGAAGAGGATGGCAAAGGCTGCTTCGAAATCTATCACGCATGCGATGAT AGTTGCATGGAATCAATTCGGAATAATACGTATGACCATAGCCACTACAGGGAAGAGGCCCTTCTCAACCGACTTAATATCAATCCTGTCAC ACTTAGTTCTGGATATAAGGACATTATTCTCTGGTTCTCTTTTGGAGCATCATGTTTTGTGCTGCTAGCCGTGGTGATGGGCCTGGTGTTCTT TTGTCTCAAGAACGGTAACATGAGGTGTACTATATGCATC
2922	ATGTATAAAATCGTGGTTATTATCGCCTTGCTAGGGGCCGTCAAAGGCCTGGACAAGATCTGTTTGGGCCACCATGCTGTGGCAAATGGCA CAATTGTTAAGACACTTACCAATGAGCAGGAGGAAGTGACCAATGCTACCGAGACCGTCGAGTCTACCGGCATTAACAGGCTGTGTATGAA GGGGCGGAAACACAAAGACCTGGCAAACTGCCACCCAATCGGAATGCTGATTGGGACTCCTGCTTGCGACTTACACCTGACCGGAATGTG GGATACCCTCATAGAGAGAGAGAACGCGATAGCCTACTGTTATCCTGGGGCGACCGTGAATGTTGAGGCCCTGAGACAGAAGATTATGGAA TCTGGGGGCATCAACAAAATTTCTACCGGATTTACATACGGCAGTTCCATCAACTCAGCCGGCACAACCCGCGCGTGTATGAGAAACGGGG GGAACAGCTTTTACGCCGAGCTGAAGTGGCTGGTTTCCAAAAGCAAAGGGCAGAATTTTCCGCAGACTACGAATACCTACAGAAATACAGAT ACTGCCGAACACCTCATAATGTGGGGCATCCATCATCCGTCCAGTACACAGGAGAAAAATGACTTATATGGTACCCAGTCTTTATCAATTTCT GTAGGCTCCTCAACTTACCGTAACAATTTCGTGCCCGTGGTCGGAGCTCGCCCGCAGGTGAACGGCCAGAGTGGGCGCATTGACTTTCATT GGACCCTCGTACAACCAGGCGACAACATCACTTTTTCTCATAATGGTGGTCTCATCGCCCCAAGCCGAGTCTCTAAGCTGATTGGGCGGGG TCTTGGGATCCAATCAGATGCACCTATCGATAACAACTGCGAAAGTAAATGTTTTTGGCGTGGCGGTTCAATCAACACCCGGCTGCCGTTCC AGAACCTGAGTCCAAGAACAGTTGGTCAGTGCCCCAAGTACGTAAACCGGAGGTCTTTGATGCTGGCCACCGGAATGCGGAACGTGCCAG AACTGATCCAGGGCCGAGGCCTTTTCGGAGCCATTGCTGGCTTTCTCGAGAACGGTTGGGAAGGAATGGTAGATGGTTGGTACGGGTTCA GACACCAGAACGCCCAGGGCACCGGTCAGGCCGCTGATTATAAATCAACCCAGGCCGCTATAGACCAAATCACCGGCAAGCTCAACAGAC TCGTGGAAAAAACGAATACGGAGTTCGAGAGCATCGAGTCAGAGTTTTCGGAAATCGAACACCAAATAGGGAATGTCATTAACTGGACTAAG GACAGCATCACCGACATTTGGACATATCAGGCAGAATTATTAGTGGCGATGGAGAATCAGCATACCATCGATATGGCAGACAGCGAAATGC TCAATCTGTACGAACGAGTACGCAAACAACTTCGCCAAAACGCAGAGGAGGATGGCAAGGGCTGCTTCGAGATATACCATGCATGCGATGA TTCTTGCATGGAGAGTATCCGTAACAATACATACGATCATTCCCAGTACCGAGAAGAGGCCCTGCTGAACCGGCTGAATATCAACCCAGTTA CACTCTCATCTGGCTACAAGGATATTATCCTCTGGTTCAGTTTTGGGGCATCTTGTTTCGTCCTGCTAGCTGTCGTCATGGGCCTGTTCTTTT TCTGTCTCAAAAACGGCAATATGCGGTGTACCATATGCATC

2923	ATGTATAAAATTGTGGTCATTATCGCCCTTCTTGGTGCCGTGAAGGGACTTGACAAAATCTGCCTGGGCCACCACGCCGTCGCAAACGGTA CTATTGTGAAAACTCTTACAAATGAGCAGGAGGAAGTGACAAATGCTACCGAGACCGTTGAAAGCACAGGTATCAACAGACTTTGTATGAAG GGACGCAAACACAAGGATCTCGGTAATTGTCATCCAATCGGTATGCTGATCGGCACTCCAGCATGTGACCTCCACCTCACCGGCATGTGGG ATACTCTAATTGAGAGAGAGAACGCGATCGCTTACTGCTATCCCGGTGCGACTGTCAATGTGGAAGCCCTTCGACAGAAGATCATGGAGTC GGGCGGCATTAACAAAATTTCTACAGGTTTCACCTATGGCAGCAGTATTAACTCTGCCGGAACAACAAGAGCCTGCATGAGGAACGGCGGA AACAGTTTCTACGCTGAACTTAAATGGTTGGTCAGCAAGAGCAAAGGCCAGAATTTCCCGCAGACCACTAACACATATCGCAACACCGACAC CGCTGAGCACCTTATTATGTGGGGAATACACCACCCCTCCAGCACCCAGGAGAAGAACGACTTGTACGGTACTCAGAGCCTCTCTATCTCC GTGGGATCCTCCACATACCGCAATAACTTTGTCCCCGTGGTTGGTGCCCGCCCCCAGGTAAACGGACAGTCAGGCAGGATTGACTTCCACT GGACCCTGGTGCAACCCGGTGACAACATTACATTCAGCCATAACGGCGGCCTAATCGCTCCCTCTCGCGTGAGTAAACTGATTGGACGGG GACTGGGGATTCAATCCGATGCTCCCATCGACAACAATTGCGAATCCAAGTGCTTTTGGCGAGGAGGGTCCATTAACACCAGGCTGCCGTT CCAAAACTTGTCTCCCCGTACGGTGGGGCAGTGTCCCAAATACGTTAACCGCCGGTCCTTAATGCTCGCTACAGGGATGAGAAATGTGCCC GAGCTGATCCAGGGTAGAGGCCTCTTCGGTGCCATTGCTGGATTCCTGGAGAATGGATGGGAGGGTATGGTCGACGGCTGGTACGGTTTT CGGCATCAAAATGCTCAGGGAACAGGACAGGCGGCCGATTATAAGAGCACCCAAGCTGCTATCGATCAGATTACAGGAAAGCTGAATAGGC TGGTCGAGAAGACAAACACGGAGTTCGAAAGTATCGAATCAGAGTTTAGCGAGATTGAACATCAGATTGGCAATGTGATTAATTGGACCAAG GACTCAATTACCGATATCTGGACCTATCAGGCCGAGCTGCTTGTGGCCATGGAGAATCAGCATACAATCGATATGGCCGATTCAGAAATGCT GAACCTGTACGAGCGTGTGCGTAAACAGCTGCGCCAGAATGCCGAGGAGGACGGAAAGGGGTGTTTTGAGATCTACCACGCGTGTGACGA CTCATGTATGGAGAGTATCCGGAACAATACTTACGACCATTCACAGTATCGCGAAGAGGCATTATTAAACCGCCTTAATATCAACCCCGTGA CACTCAGCAGCGGGTATAAGGACATCATTCTGTGGTTCTCATTTGGGGCCTCCTGCTTCGTTCTGCTGGCCGTGGTGATGGGGTTATTCTTT TTTTGTCTAAAGAACGGTAATATGAGGTGCACAATATGTATC
2924	ATGTATAAGGTAGTAGTGATCATCGCCCTGCTGGGGGCCGTGAGAGGCCTTGATAAGATCTGTTTGGGTCACCACGCCGTTGCAAACGGGA CCACTGTTAAGACCCTTACCAATGAGCAGGAGGAGGTGACGAATGCAACCGAGACCGTCGAGTCCACAAGTCTGAACAAGCTGTGTATGAA AGGGAGGAGGTATAAAGATCTGGGGAACTGCCACCCCATCGGAATGTTGATTGGAACCCCAGTCTGCGATCTGCACTTGACCGGCACCTG GGATACACTGATTGAGCGGGAGAACGCTACCGCCTACTGCTATCCCGGGGTCACTATTAATGAGGAAGCCTTGAGACAGAAGATCATGGAG AGTGGAGGGATATCCAAGATGCGGACAGGGTTCACCTACGGTCCCTCCATCAATTCCGCTGGCACCACCCGGTCATGCATGCGCAATGGC GGCAATTCCTTTTACGCCGAGCTGAAATGGCTGGTTAGCGGGACTAAGGGACAGAACTTCCCACAAACCACTAACACCTATAGAAATACTGA TACCGCGGAGCATCTGATCATATGGGGAATTCATCATCCTTCATCGACCCAGGAGAAAAACGACCTGTACGGGACTCAGTCCCTGTCCATAT CCGTGGGAAGTTCCACTTATCAGAACAATTTCGTGCCTGTTATCGGAGCCCGACCTCAGGTTAACGGACAATCGGGACGGATTGAGTTCCA CTGGACTTTAGTGAGACCGGGCGACAACATCACCTTTTCTCATAACGGCGGCCTGATTGCCCCCGATAGAGTCAGCAAATTAATCGGCAAG GGAATCGGCATACAGAGCGGGGCTGTTATTGACAAGGACTGCGAATCTAAATGCTTTTGGCGTGGTGGCTCCATTATCACTGAACTGCCCT TCCAGAATCTCAGCCCTAGAACTGTGGGCCAGTGCCCGAAGTATGTGAAAAAGAGGTCTCTTCTTCTTGCTACAGGTATGAGGAACGTCCC TGAGGTTGTCCAGGGGCGCGGCTTGTTCGGCGCTATCGCCGGGTTCATCGAAAACGGGTGGGAGGGGATGGTGGACGGCTGGTATGGCT TTCGACACCAGAATGCTCAAGGGATCGGGCAGGCAGCCGACTACAAATCCACCCAGACAGCTATAGACCAGATCACGGGGAAACTTAATCG TCTGATCGAAAAGACAAATACGGAGTTTGAGTCTATCGAGTCCGAATTCTCGGAGATTGAGCACCAGATTGGTAATGTCATCAATTGGACTA AGGACAGTATTACTGACATCTGGACTTATCAAGCAGAGCTGCTGGTAGCTATGGAAAACCAACATACAATCGATATGGCCGATAGCGAGATG CTGAACCTTTACGAACGAGTAAGGAAACAGCTGAGGCAGAACGCCGAAGAGGACGGCAAAGGTTGCTTTGAAATTTACCACACCTGTGACA ATTCCTGTATCGAAAGCATACGAAATAATACCTATGATCACTCCCAGTACAGACGAGGAAGCGCTCCTGAACCGACTGAATATCAACCCCGTG AAACTGAGTTCCGGATACAAGGATATCATCCTATGGTTCTCATTCGGCGCCAGCTGTTTCGTTCTGTTAGCGGTTATAATGGGTCTGGGGTT TTTTTGTCTCAAAAATGGCAATATGCGTTGTACCATTTGTATA
2925	ATGTACAAGGTTGTGGTCATCATCGCCCTGCTTGGTGCTGTGAAAGGGCTTGATAAGATATGTTTGGGCCACCATGCCGTGGCTAACGGAA CCATTGTGAAGACCTTGACCAACGAGCAGGAGGAAGTGACTAATGCCACAGAAACTGTAGAGTCTACCGGTATAAACAGGCTGTGCATGAA GGGAACGAAGCACAAAGATCTCGGCAATTGTCATCCCATAGGCATGCTGATCGGTACCCCCGCCTGCGATCTGCACTTGACTGGCACCTG GGACACGCTGATCGAGAGGGAAAACGCTATAGCCTATTGTTACCCTGGGGCTACCGTTAATGTGGAAGCGCTGCGACAAAAGATAATGGAG AGCGGGGGCATTGATAAAATTTCAACCGGATTTACCTACGGCTCTAGTATCAACTCTGCGGGTACTACGCGAGCATGCATGAGAAACGGGG GGAATTCCTTTTATGCAGAGCTTAAGTGGCTGGTTAGCAAGAACAAGGGTCAGAATTTCCCCCAAACTACTAACACTTACAGAAACACAGAT ACCGCCGAACATCTGATCATGTGGGGCATTCATCACCCATCCTCTATTCAAGAGAAGAATGACCTTTATGGCACTCAAAGCCTCTCTATCTC CGTGGGAAGCAGCACTTACCGCAATAATTTCGTGCCCGTCGTTGGGGCACGACCGCAGGTCAATGGCCAGAGTGGCAGGATCGATTTCCA TTGGACCTTAGTCCAGCCGGGCGATAATATTACCTTCAGCCATAACGGAGGACTTATTGCGCCGTCCAGAGTGTCAAAACTGATCGGGAGA GGACTCGGCATCCAGAGCGATGCTCCTATTGATAATAATTGCGAGTCCAAGTGTTTTTGGCGTGGGGGTTCGATCAATACTAGACTACCTTT CCAGAACTTGAGCCCACGCACTGTGGGCCAGTGCCCCAAATATGTAAATAGAAGGTCCCTTATGCTCGCTACAGGTATGAGGAACGTCCCA GAACTCATTCAGGGCCGAGGTTTGTTTGGCGCAATCGCCGGATTTCTGGAGAACGGGTGGGAGGGCATGGTCGACGGCTGGTACGGATTC CGCCACCAGAACGCCCAAGGTACAGGTCAGGCCGCCGACTATAAGAGCACGCAGGCTGCTATAGATCAGATTACAGGAAAGCTAAACCGA CTTGTGGAGAAGACAAATACAGAGTTCGAGTCAATAGAGAGTGAGTTCAGTGAGATAGAGCATCAGATCGGTAATGTCATTAACTGGACAAT GGATAGCATCACCGATATTTGGACTTATCAGGCCGAATTGCTGGTGGCTATGGAAAATCAGCACACCATTGATATGGCAGACTCTGAGATGC TAAATCTTTACGAGCGGGTGCGAAAGCAGTTGCGGCAGAACGCAGAAGAAGACGGAAAAGCGCTGCTTCGAAATATATCACGCTTGCGACGA TTCCTGTATGGAAAGCATCAGAAATAACACATATGACCACAGTCAGTATCGCGAGGAGGCTCTCCTGAACAGACTCAATATCAATCCGGTCA CCCTCTCCTCAGGTTACAAGGACATTATTCTCTGGTTTTCTTTCGGTGCGTCTTGCTTTGTGCTACTCGCTGTGGTGATGGGCTTGGTGTTCT TCTGCCTGAAGAACGGCAATATGCGGTGCACTATCTGTATC

2926	ATGTACAAAATTGTGGTCATCATCGCTTTACTAGGAGCTGTGAAAGGCTTGGATAAAATTTGTCTGGGACATCATGCTGTCGCTAACGGGAC CATAGTGAAAACACTCACGAACGAACAGGAGGAAGTTACCAATGCCACCGAAACGGTCGAGTCAACAGGGATTAACCGGCTGTGTATGAAG GGAAGAAAGCATAAAGATCTGGGGAACTGCCACCCTATAGGAATGTTGATAGGCACGCCTGCTTGTGACCTGCACCTGACTGGGATGTGG GATACCCTGATCGAAAGAGAGAACGCTATAGCCTATTGCTATCCCGGTGCAACAGTGAACGTGGAGGCATTGAGACAGAAAATAATGGAAA GTCGAGCGCATCAATAAGATCTCAACCGGTTTTACCTATGGATCTAGTATAAATAGTGCAGGGACCACACGTGCCTGCATGAGAAATGGGGG CAATTCTTTCTACGCGGAACTGAAGTGGCTGGTGAGCAAATCTAAGGGGCAGAATTTTCCACAGACCACGAATACATATCGTAACACAGACA CCGCGGAGCACTTGATTATGTGGGGTATCCACCACCCCAGCAGCACGCAGGAAAAAAACGATCTCTACGGTACTCAGAGCCTGTCTATCTC AGTAGGCTCATCAACATACCGGAATAACTTTGTTCCAGTTGTGGGGGCCCGGCCTCAGGTGAATGGCCAGTCTGGACGCATTGATTTTCATT GGACGCTGGTACAGCCAGGAGACAATATCACATTTAGTCATAATGGCGGTTTGATCGCCCCGAGTAGAGTGTCTAAGCTGATCGGCCGAGG CCTTGGCATTCAGTCCGACGCCCCCATCGACAATAACTGCGAAAGCAAATGCTTTTGGAGAGGAGGTTCTATTAATACACGCCTGCCATTTC AAAACTTGAGCCCTAGAACTGTGGGTCAGTGTCCCAAGTACGTGAATCGCCGCTCCCTGATGTTAGCCACCGGGATGAGGAATGTGCCTGA GCTGATCCAGGGCAGGGGCCTATTTGGAGCCATCGCTGGTTTCTTGGAGAATGGTTGGGAGGGTATGGTAGATGGATGGTATGGGTTTCG GCATCAGAATGCGCAAGGCACTGGCCAGGCCGCGGATTACAAAAGCACTCAGGCTGCCATTGACCAGATAACCGGAAAACTCAACAGACT GGTTGAGAAGACCAACACTGAATTTGAGAGCATTGAGTCAGAGTTTAGCGAAATAGAACACCAGATCGGGAACGTTATTAACTGGACAAAG GACTCCATTACGGACATATGGACATATCAGGCAGAGCTGCTAGTGGCAATGGAGAACCAACACACTATTGACATGGCCGACTCTGAGATGC TGAACCTCTATGAGCGCGTCAGAAAGCAGCTAAGGCAGAATGCAGAAGAGGATGGTAAAGGGTGTTTTGAAATCTATCATGCATGCGACGA TTCATGCATGGAAAGCATCAGGAACAATACTTATGACCATTCCCAATACCGAGAGGAAGCCCTGCTGAACAGACTGAATATTAACCCAGTGA CCCTATCCAGTGGCTATAAGGATATCATCCTGTCGTTCAGCTTTGGCGCTAGCTGTTTCGTGTTACTCGCGGTCGTCATGGGACTGTTTTTT TTTTGTTTGAAGAACGGGAATATGAGATGCACAATCTGTATT
2927	ATGTATAAAATCGTAGTGATTATTGCGTTGCTTGGCGCTGTGAAAGGGCTAGATAAGATCTGCCTGGGCCACCACGCCGTAGCTAACGGAA CCATCGTCAAAACTTTGACTAACGAGCAGGAAGAGGTCACCAATGCTACTGAGACAGTCGAGTCTACAGGCATAAATAGACTCTGTATGAAG GGTAGAAAGCACAAGGATCTTGGGAATTGCCACCCCATTGGTATGCTGATTGGGACACCCGCCTGCGACCTGCACCTCACAGGGATGTGG GACACCTTAATTGAGCGAGAGAATGCAATCGCCTACTGTTACCCCGGGGCTACTGTTAACGTCCAAGCTCTGCGCCAGAAAATCATGGAGA GCGGGGGGATCAACAAGATCTCCACCGGCTTCACCTATGGCTCATCTATCAATTCTGCAGGCACTACCAGAGCGTGCATGAGAAATGGAGG AAATTCCTTCTATGCGGAATTGAAATGGCTTGTGTCAAAATCTAAAGGGCAAAATTTTCCGCAGACCACCAACACCTACAGGAACACCGATA CTGCAGAGCATCTGATTATGTGGGGGATTCACCACCCTTCATCCACTCAGGAAAAGAACGATCTCTATGGGACACAATCATTGTCCATTAGC GTGGGCTCTTCCACTTACAGAAACAACTTCGTGCCTGTCGTTGGGGCGCGACCGCAAGTTAATGGGCAATCCGGGCGAATTGATTTCCACT GGACGCTGGTTCAACCTGGAGACAATATCACTTTCTCACATAATGGAGGGTTAATTGCTCCCTCCCGCGTCTCCAAACTGATTGGAAGAGGC CTGGGCATTCAGTCCGATGCCCCAATCGACAACAATTGCGAATCCAAGTGTTTCTGGCGCGGGGGCTCAATCAATACCCGCCTGCCCTTCC AAAACCTGAGTCCCCGCACAGTGGGCCAATGCCCTAAATATGTCAACCGTCGATCCCTCATGCTTGCCACAGGCATGAGGAACGTCCCAGA GCTGATCCAGGGCCGCGGACTCTTTGGAGCTATCGCAGGCTTTCTTGAAAACGGGTGGGAGGGCATGGTCGATGGTTGGTATGGATTTCG GCATCAGAATGCACAAGGTACTGGACAGGCCGCTGATTACAAGTCAACCCAGGCGGCCATTGACCAGATAACCGGGAAGCTCAACCGGTT GGTCGAGAAGACTAATACTGAGTTTGAATCCATCGAAAGTGAGTTCTCTGAGATAGAACATCAGATTGGTAATGTCATAAATTGGACTAAAGA TAGCATAACGGACATTTGGACCTATCAGGCAGAATTACTGGTGGCAATGGAAAATCAACACACCATCGACATGGCCGATTCTGAAATGCTCA ACCTATACGAAAGAGTCAGAAAGCAGCTCAGACAGAACGCCGAGGAAGACGGCAAAGGGTGCTTTGAAATTTACCATGCTTGTGACGATTC TTGTATGGAGAGTATACGCAATAACACTTATGACCATTCACAGTACCGAGAGGAGGCACTCCTGAATCGACTGAACATCAACCCCGTCACAC TGTCCTCAGGCTACAAGGATATTATCCTTTGGTTTAGTTTTGGTGCGTCGTGCTTCGTCCTCCTGGCCGTCGTGATGGGGTTATTCTTTTTTT GCTTGAAAAATGGAAATATGCGTTGCACCATCTGTATC
2928	ATGTATAAGATAGTCGTCATAATCGCTCTGCTCGGCGCAGTGAAAGGGCTGGATAAAATTTGTCTGGGTCACCACGCCGTAGCGAATGGCA CCATCGTGAAGACACTGACTAACGAGCAAGAGGAGGTGACTAATGCCACAGAAACCGTCGAGTCTACAGGAATTAACCGCCTCTGCATGAA AGGCACGAAACACAAAGATCTCGGCAACTGTCACCCGATCGGAATGCTCATAGGGACTCCAGCCTGTGATCTCCACCTTACCGGAATGTGG GATACTCTCATAGAGAGGGAGAATGCCATCGCTTATTGTTACCCCGGCGCGACAGTTAACGTGGAAGCTCTCAGACAGAAGATCATGGAAA GTGGGGGCATTAACAAGATATCCACCGGATTCACTTACGGTTCATCCATTAATTCTGCGGGAACCACTCGGGCCTGTATGCGGAACGGTGG CAATTCTTTCTATGCCGAGCTGAAGTGGCTGGTCTCCAAGTCCAAGGGCCAGAATTTCCCACAGACTACCAACACATATAGAAACACTGACA CTGCCGAGCATCTGATCATGTGGGGGATCCATCACCCAAGTTCAACTCAGGAGAAGAATGACCTCTACGGTACCCAGAGCTTATCTATTAGT GTTGGCTCCTCTACTTACCGCAATAATTTTGTGCCAGTAGTCGGTGCTCGCCCTCAGGTGAATGGGCAGTCCGGCAGGATCGACTTCCACT GGACTCTAGTCCAACCCGGGGATAACATCACATTCAGTCATAACGGGGGATTGATCGCTCCAAGTAGAGTTTCAAAACTCATAGGTAGGGG CCTCGGGATTCAGTCGGACGCTCCTATCGACAACAATTGCGAGTCTAAGTGTTTCTGGCGAGGGGGAAGCATAAATACCCGACTCCCCTTC CAGAATCTCAGCCCAAGGACTGTGGGTCAGTGCCCAAAATACGTTAACCGCCGGAGCCTAATGCTGGCGACTGGCATGAGAAACGTCCCC GAACTGATTCAAGGTCGCGGACTCTTCGGGGCAATCGCGGGATTTCTGGAAAACGGCTGGGAGGGAATGGTGGACGGCTGGTACGGCTTC CGACACCAGAACGCTCAAGGGACAGGCCAAGCGGCTGACTACAAGTCCACCCAGGCCGCGATTGACCAAATCACAGGGAAATTGAATAGG CTGGTAGAAAAGACGAATACAGAGTTCGAATCCATAGAGAGTGAATTTTCCGAGATCGAACACCAAATAGGAAATGTCATCAATTGGACGAA GGACAGCATTACGGACATCTGGACATACCAGGCAGAGCTCCTCGTGGCGATGGAAAATCAGCACACAATCGACATGGCTGACTCAGAGAT GCTCAATCTGTACGAGCGGGTCCGCAAACAGTTGAGGCAGAATGCTGAAGAAGACGGTAAAGGGTGTTTTGAAATCTATCATGCCTGCGAC GATTCATGCATGGAATCTATTAGAAACAATACGTACGATCACTCCCAGTATAGGGAGGAAGCTCTGCTGAATAGACTGAACATAAACCCCGT CACCCTGTCTTCCGGTTACAAGGACATCATACTTTGGTTCTCGTTCGGTGCAAGCTGCTTCGTTCTGCTCGCTGTTGTCATGGGCCTGTTCT TCTTCTGTTTGAAAAACGGCAATATGCGATGCACCATTTGTATT

2929	ATGTATAAAATTGTCGTAATCATTGCCCTGCTGGGAGCGGTTAAGGGGCTTGATAAGATATGCTTGGGGCATCACGCCGTAGCTAATGGTAC AATCGTCAAAACACTAACTAATGAACAGGAGGAAGTAACAAACGCAACTGAAACTGTTGAGTCTACGGGAATCAACAGGCTCTGTATGAAAG GCCGAAAGCACAAAGACTTGGGTAATTGTCACCCGATCGGGATGCTCATCGGGACACCTGCATGCGACCTCCACCTCACCGGCATGTGGG ATACACTTATTGAACGGGAGAACGCGATCGCCTACTGCTACCCTGGGGCAACGGTGAATGTAGAAGCTCTCCGGCAGAAGATCATGGAAAG CGGCGGAATTAATAAAATTAGTACCGGATTTACCTACGGGTCATCCATAAACAGCGCCGGACCACACGCGCCTGCATGAGGAACGGCGG CAACAGCTTTTATGCCGAGCTCAAGTGGCTTGTCTCCAAATCCAAAGGACAGAACTTTCCTCAGACAACGAACACATATCGTAATACTGATAC CGCAGAACATCTGATCATGTGGGGAATCCACCACCCCAGTAGCACCCAAGAGAAGAACGATCTATACGGAACTCAGTCTCTTTCCATCAGC GTTGGGTCCTCTACTTACAGGAACAATTTTGTACCGGTCGTGGGAGCCCGCCCCCAGGTGAATGGGCAGTCCGGAAGGATTGATTTTCACT GGACTTTAGTACAGCCTGGAGACAACATAACCTTTTCCCACAACGGAGGTCTCATAGCCCCTTCTCGAGTGAGCAAGTTGATAGGGCGTGG GTTGGGAATCCAGTCTGACGCTCCGATCGACAATAATTGTGAGTCCAAGTGCTTTTGGAGGGGCGGCTCTATCAATACGAGATTGCCTTTCC AGAACCTTAGTCCAAGAACCGTGGGGCAATGTCCAAAGTATGTCAATCGACGCTCACTAATGTTAGCAACCGGCATGAGGAATGTTCCCGA GCTAATCCAAGGCCGGGGACTCTTTGGCGCCATCGCGGGCTTCTTGGAGAACGGTTGGGAAGGAATGGTCGACGGGTGGTATGGCTTCC GGCACCAAAACGCCCAGGGTACAGGCCAGGCAGCCGACTATAAATCCACACAGGCAGCCATCGATCAGATCACCGGCAAGCTCAATAGGC TCGTGGAAAAAACAAACACTGAGTTCGAGTCCATTGAATCCGAGTTCTCGGAGATCGAGCACCAAATCGGGAACGTCATCAATTGGACTAAG GACAGTATTACAGATATCTGGACATATCAGGCCGAACTTTTGGTCGCTATGGAAAATCAACATACGATTGACATGGCAGACTCAGAGATGTT AAACCTGTATGAGAGAGTGCGCAAACAGCTGCGGCAGAATGCCGAAGAAGATGGGAAGGGATGTTTCGAGATTTACCATGCCTGTGACGAT TCCTGCATGGAGTCGATTCGTAATAATACTTATGATCATTCCCAATATAGAGAGGAAGCATTATTAAACCGGCTGAATATTAACCCCGTGACC CTGAGCTCCGGGTACAAGGACATTATTCTGTGGTTCTCATTCGGCGCATCTTGTTTCGTCTTACTCGCAGTGGTCATGGGACTGTTTTTCTTT TGTCTGAAGAATGGGAACATGCGCTGTACCATTTGTATC
2930	ATGTACAAGATTGTCGTCATCATAGCCCTCCTAGGTGCTGTTAAAGGGCTCGATAAGATCTGCCTTGGACATCACGCCGTGGCAAATGGAAC AATAGTAAAGACACTCACAAACGAGCAGGAAGAAGTGACGAATGCCACCGAAACCGTCGAGAGTACCGGCATCAACAGACTGTGCATGAAG GGCAGGAAGCATAAAGACCTGGGCAACTGCCACCCTATCGGCATGCTAATCGGTACACCAGCCTGCGATTTACACCTGACCGGAATGTGG GACACACTGATTGAACGAGAAAATGCCATTGCTTACTGTTATCCTGGCGCTACCGTGAATGTGGAGGCTCTCAGGCAGAAAATTATGGAGA GTGGGGGGATCAACAAGATTAGTACTGGCTTTACCTACGGCTCTAGCATCAACTCTGCTGGCACAACGCGCGCTTGTATGAGAAATGGCGG CAATTCATTCTACGCCGAGCTTAAGTGGCTGGTTAGCAAGTCCAAGGGACAGAACTTTCCTCAAACGACGAATACATACCGCAACACCGATA CTGCTGAGCATCTGATCATGTGGGGCATCCATCATCCCTCCTCAACTCAGGAAAAAAATGACCTCTACGGGACTCAGTCCCTATCAATCTCC GTTGGTAGTAGCACTTATCGGAACAACTTTGTGCCTGTGGTGGGCGCTAGACCTCAGGTCAATGGGCAGTCTGGGCGGATCGACTTTCACT GGACTCTCGTACAACCTGGTGACAACATAACTTTCTCACACAATGGCGGCCTCATCGCCCCAAGCCGAGTCAGTAAATTAATCGGCCGGGG TCTGGGCATCCAGAGCGACGCACCAATTGATAACAATTGTGAATCCAAATGCTTCTGGCGGGGCGGGAGCATTAACACTCGGTTACCCTTT CAGAACTTGAGTCCTCGCACCGTGGGACAATGCCCGAAGTACGTGAACCGCCGCTCACTCATGCTGGCAACCGGCATGCGGAACGTCCCC GAACTCATCCAGGGCAGAGGGCTCTTTGGCGCCATCGCTGGATTTCTGGAGAACGGATGGGAGGGCATGGTCGATGGCTGGTACGGCTTC AGGCATCAGAACGCCCAGGGGACCGGCCAAGCCGCTGATTACAAATCTACTCAGGCTGCCATAGACCAAATCACAGGGAAGCTTAATCGC CTGGTGGAAAAGACCAATACAGAATTTGAAAGCATCGAGTCTGAGTTCAGTGAAATTGAACATCAGATCGGGAACGTGATCAATTGGACCAA AGACAGCATTACCGATATTTGGACCTACCAGGCAGAGTTGCTCGTTGCCATGGAGAACCAGCACACTATTGATATGGCCGACTCCGAGATG CTGAATCTCTATGAGAGGGTTCGCAAACAACTGAGACAAAATGCGGAAGAAGATGGAAAGGGCTGTTTTGAGATCTATCACGCCTGCGATG ATAGTTGCATGGAAAGTATCAGAAATAACACATACGATCACTCTCAGTACCGGGAAGAGGCCCTCCTGAACAGGCTTAATATTAACCCCGTG ACGCTTTCATCTGGCTATAAGGACATCATTTTGTGGTTCTCATTTGGCGCCAGCTGCTTTGTGCTGCTGGCTGTAGTTATGGGCTTATTCTTC TTCTGCCTGAAGAACGGAAACATGAGATGTACCATTTGCATT
2909	ATGTATAAAATTGTGGTCATAATTGCCCTGCTTGGGGCAGTGAAGGGGTTAGACAAAATTTGCTTGGGGCACCACGCAGTGGCCAACGGGA CAATTGTCAAAACACTGACAAATGAACAGGAGGAAGTGACAAACGCCACAGAAACCGTTGAATCAACCGGAATCAATAGACTTTGTATGAAG GGCAGAAAGCACAAAGATTTGGGCAATTGCCATCCAATTGGAATGTTGATCGGCACCCCTGCCTGTGACCTGCATCTGACTGGGATGTGGG ACACCCTGATCGAACGGGAGAACGCAATTGCGTACTGTTATCCCGGCGCTACAGTTAATGTGGAGGCCCTCCGACAGAAGATCATGGAGA GTGGCGGCATTAACAAGATAAGCACCGGCTTCACCTATGGCTCCAGCATAAACTCAGCTGGCACTACTCGGGCATGCATGAGGAACGGCG GGAACTCCTTCTACGCTGAACTCAAGTGGCTGGTGTCGAAATCTAAGGGACAGAACTTCCCGCAAACCACCAACACTTACCGGAATACGGA CACAGCAGAGCATCTGATCATGTGGGGCATACACCATCCATCCAGCACGCAGGAAAAAAACGACCTGTATGGAACACAGAGCTTGTCTATC TCGGTGGGGAGCAGCACCTACAGGAACAATTTCGTCCCTGTTGTTGGAGCCGGGCCCCAAGTTAATGGCCAGAGTGGCAGAATCGACTTT CATTGGACACTGGTGCAGCCTGGAGATAATATTACTTTTAGCCACAACGGTGGACTCATTGCTCCCAGTAGGGTCTCGAAGTTGATCGGTC GTGGCCTGGGTATTCAGAGCGACGCTCCCATTGACAATAATTGCGAGAGCAAATGTTTCTGGCGGGGAGGTAGTATTAATACTAGGCTCCC TTTCCAGAATCTGTCACCGCGCACCGTTGGCCAGTGTCCGAAATATGTCAACCGGCGATCCCTAATGCTAGCCACTGGCATGCGGAACGTC CCTGAACTCATTCAAGGCAGGGGACTGTTCGGCGCTATCGCTGGGTTTCTGGAAAATGGATGGGAGGGCATGGTTGATGGCTGGTATGGC TTCAGGCACCAGAACGCTCAGGGCACCGGGCAAGCCGCCGATTATAAATCCACTCAGGCGGCCATCGATCAAATAACTGGCAAACTAAATA GGCTGGTCGAAAAGACAAACACTGAGTTTGAGAGCATTGAGAGCGAATTTTCTGAAATTGAGCACCAGATCGGTAACGTTATTAACTGGACC AAGGATTCAATTACAGATATTTGGACCTACCAGGCAGAGCTTCTGGTCGCCATGGAAAACCAGCACACAATTGATATGGCCGACAGCGAAAT GTTGAACCTTTACGAGAGAGTGCGGAAGCAGTTGAGACAGAACGCTGAGGAGGATGGGAAAGGGTGCTTCGAGATTTACCATGCCTGCGA TGATTCTTGTATGGAGAGCATCAGGAACAATACCTATGACCACAGCCAGTATCGCGAAGAGGCCCTGCTGAACAGACTGAACATCAACCCC GTGACACTCTCTTCTGGCTATAAGGATATAATCCTGTGGTTTAGCTTCGGAGCCAGCTGCTTTGTCCTTCTAGCAGTTGTCATGGGGCTCTT CTTTTTCTGCCTGAAGAATGGAAACATGAGATGTACAATTTGCATT

2910	ATGTACAAAGTGGTGGTCATCATTGCCTTGTTGGGTGCTGTCCGGGGGGTTAGACAAGATCTGTCTGGGGCATCACGCCGTAGCCAATGGAA CTACTGTCAAGACCTTAACCAACGAGCAAGAAGAGGTCACAAATGCGACGGAGACCGTGGAAAGCACATCGCTTAATAAGCTGTGCATGAA GGGGCGTCGCTACAAAGATCTGGGAAATTGCCATCCAATCGGCATGCTGATTGGAACTCCCGTCTGTGATCTACACCTGACTGGAACTTGG GATACTTTGATTGAAAGAGAGAACGCTACGGCTTACTGTTATCCAGGGGTAACCATCAATGAGGAGGCCCTGCGTCAGAAGATTATGGAGA GTCGGGGAATCAGCAAAATGCGGACAGGATTCACCTATGGGCCGTCCATTAATTCTGCCGGTACCACGAGAAGCTGCATGCGGAACGGTG GAAACTCGTTTTACGCAGAACTGAAATGGTTGGTCTCTGGCACGAAAGGACAGAACTTCCCACAGACTACAAACACCTATCGAAATACCGAT ACCGCCGAGCACCTGATCATTTGGGGGATCCACCACCCTAGCAGCACCCAGGAAAAAAATGACTTATATGGCACTCAATCCCTGAGCATCT CTGTGGGCTCCTCTACTTATCAAAACAATTTTGTGCCTGTGATTGGAGCCAGGCCCCAGGTGAATGGACAATCCGGACGCATTGAGTTCCA CTGGACACTTGTCAGACCCGGGGATAATATCACCTTTTCTCACAACGGCGGTCTTATAGCGCCAGACCGTGTTAGCAAGCTGATTGGCAAG GGGATCGGCATCCAATCAGGTGCTGTGATCGACAAGGATTGTGAGTCTAAGTGCTTTTGGAGGGGTGGGTCCATCATTACCGAGCTGCCTT TTCAGAACCTAAGCCCACGCACCGTCGGACAGTGCCCCAAATATGTGAAAAAAAGATCTTTACTGTTGGCCACCGGAATGAGAAATGTGCC AGAGGTGGTCCAAGGTCGGGGCTTGTTCGGGGCGATTGCAGGCTTTATAGAGAACGGCTGGGAGGGCATGGTGGACGGTTGGTACGGTT TCCGGCACCAGAACGCACAGGGAATTGGGCAAGCCGCCGACTACAAGTCAACCCAAACAGCCATAGACCAGATTACGGGAAAACTGAACA GATTAATCGAGAAAACTAATACAGAATTTGAGAGCATCGAATCTGAGTTTTCCGAAATCGAGCACCAGATAGGGAATGTCATCAATTGGACTA AGGACTCCATTACAGACATCTGGACTTACCAGGCCGAGTTGCTGGTCGCCATGGAGAATCAGCATACAATTGATATGGCTGACAGCGAAAT GCTCAACCTGTACGAGCGCGTAAGGAAGCAACTGCGCCAGAATGCTGAGGAAGACGGGAAGGGTTGCTTTGAGATTTACCACACCTGTGA CAACTCATGTATGGAAAGCATCCGGAACAACACTTACGATCACTCTCAATATCGCGAGGAGGCCCTTCTGAATCGGCTGAATATAAACCCAG TTAAATTGTCTTCAGGCTATAAAGACATCATCCTGTGGTTCTCGTTCGGGGCTAGCTGTTTCGTGCTGCTGGCCGTGATAATGGGCCTTGGA TTCTTTTGTCTCAAGAACGGCAATATGAGGTGCACCATATGTATC
2911	ATGTACAAGGTCGTTGTTATCATCGCCCTGTTGGGCGCGGTGAGGGGACTCGATAAGATTTGTCTGGGCCACCACGCCGTCGCCAACGGC ACGACTGTGAAGACATTGACCAACGAGCAGGAGGAGGTTACTAATGCCACCGAAACTGTGGAGAGTACCAGTCTGAACAAGTTGTGCATGA AGGGCCGGCGGTATAAGGATCTGGGAAATTGCCACCCCATCGGAATGCTCATTGGTACCCCAGTGTGTGATCTTCACCTCACAGGAACCTG GGATACCCTCATCGAGCGTGAGAACGCAACCGCGTATTGCTATCCCGGCGTGACCATAAATGAGGAAGCCCTGAGACAGAAGATCATGGA GAGCGGCGGCATAAGTAAGATGAGGACCGGCTTCACTTATGGCCCCTCAATTAACAGTGCGGGAACAACCCGGTCCTGCATGAGAAATGG CGGCAATAGTTTCTATGCTGAACTGAAGTGGCTTGTGAGCGGAACAAAGGGACAGAATTTTCCTCAGACAACGAACACGTATCGCAACACT GATACCGCTGAACACCTGATCATCTGGGGCATCCATCATCCCAGTTCTACTCAGGAGAAAAATGACTTGTATGGTACCCAATCTTTGTCTATA TCCGTCGGGAGTTCCACATACCAGAACAACTTCGTCCCTGTGATTGGGGCGCGACCTCAAGTGAACGGCCAGTCGGGCCGCATAGAGTTT CACTGGACCTTGGTCCGACCAGGCGATAACATTACATTTTCGCACAATGGCGGCTTGATCGCCCCCGATCGCGTGAGCAAACTTATCGGGA AGGGGATCGGAATCCAGAGCGGCGCAGTGATAGACAAGGACTGCGAAAGCAAATGCTTCTGGCGCGGTGGATCTATCATCACTGAATTGC CTTTCCAAAACCTGTCCCCCCGAACAGTGGGTCAGTGCCCGAAGTATGTGAAGAAGCGCAGTTTGCTATTAGCGACTGGCATGCGGAACGT GCCTGAGGTGGTTCAGGGGCGCGGTCTCTTTGGCGCCATCGCAGGATTTATTGAGAATGGCTGGGAGGGCATGGTCGACGGCTGGTACG GATTTAGACATCAGAATGCTCAGGGAATTGGGCAGGCCGCCGATTACAAGAGTACCCAGACAGCAATTGACCAGATCACAGGCAAACTTAA CAGGCTTATTGAGAAAACCAACACCGAGTTCGAGTCCATCGAGTCAGAGTTTAGCGAGATCGAACATCAAATCGGCAATGTGATTAATTGGA CAAAAGATTCTATTACCGATATCTGGACCTACCAGGCAGAGCTGCTGGTGGCTATGGAAAACCAGCACACCATAGACATGGCAGACAGTGA GATGCTGAACCTATATGAGCGTGTGAGGAAACAATTGAGACAAAATGCCGAGGAAGACGGGAAGGGATGCTTCGAAATATACCACACATGT GATAACTCCTGTATGGAATCAATTCGGAACAATACTTACGACCACAGCCAGTATCGCGAGGAAGCCTTACTTAATCGTTTAAATATTAATCCC GTTAAGCTGAGCTCCGGTTATAAAGACATCATTCTGTGGTTCTCCTTTGGAGCCAGTTGCTTCGTTTTGCTCGCCGTTATCATGGGCCTGGG GTTTTTCTGTCTCAAGAATGGAAATATGCGATGCACCATTTGCATC
2912	ATGTATAAGATTGTCGTGATTATCGCTCTCCTTGGTGCCGTTAAGGGCCTGGACAAGATCTGTCTTGGACACCACGCTGTGGCCAATGGCA CTATTGTCAAAACCCTCACTAATGAGCAGGAGGAGGTCACCAACGCAACGGAGACAGTAGAAAGCACCGGGATCAACCGTCTGTGCATGAA GGGGCGGAAACATAAAGACCTCGGAAACTGCCATCCGATTGGAATGCTGATCGGGACCCCAGCTTGCGACCTCCATCTGACAGGGATGTG GGACACGCTTATCGAACGCGAGAATGCCATTGCCTATTGTTATCCTGGGGCTACTGTGAACGTAGAAGCCCTGCGTCAGAAGATTATGGAG TCCGGCGGGATCAATAAAATCAGCACAGGGTTCACCTATGGCTCTTCAATCAACAGTGCCGGAACAACCCGCGCCTGCATGCGGAACGGC GGGAATTCCTTCTACGCCGAATTAAAGTGGTTGGTTTCTAAAAGTAAAGGGCAAAACTTCCCTCAGACCACGAACACCTATCGTAACACAGA CACCGCCGAGCATCTCATCATGTGGGGGATCCACCATCCCAGCAGTACACAGGAAAAGAATGATCTGTACGGCACACAGTCGCTGTCGATC TCAGTTGGGAGTAGCACTTACCGCAACAACTTTGTGCCCGTCGTGGGGGCGAGGCCGCAGGTAAATGGGCAATCTGGACGTATAGACTTC CACTGGACTCTAGTGCAACCCGGCGACAACATCACATTTTCTCATAACGGGGGCCTTATAGCCCCGTCTCGAGTGTCCAAACTGATTGGCC GCGGCCTCGGAATCCAGTCCGATGCCCCCATCGACAATAATTGCGAGTCTAAATGTTTTTGGCGAGGGGGGAGCATTAACACACGGCTCCC ATTTCAAAATTTGAGTCCGCGCACCGTCGGTCAGTGTCCAAAATATGTGAACAGACGCAGCTTGATGCTCGCAACGGGCATGAGAAACGTT CCTGAATTGATCCAGGGTCGGGGACTGTTTGGGGCCATCGCAGGCTTTTTGGAGAACGGTTGGGAAGGAATGGTGGACGGATGGTACGGC TTTAGACACCAGAACGCACAGGGAACCGGTCAGGCAGCTGACTACAAGTCAACTCAGGCCGCCATTGACCAGATCACAGGCAAGCTGAAT CGACTCGTCGAGAAGACCAACACCGAATTTGAGTCTATCGAATCAGAGTTTTCGGAGATCGAGCACCAGATCGGCAATGTGATCAATTGGA CAAAGGATTCTATCACCGATATCTGGACGTATCAGGCGGAATTGCTGGTAGCCATGGAGAATCAACACACAATAGATATGGCCGATTCTGAG ATGCTTAACCTGTACGAACGGGTCCGGAAGCAACTCCGCCAGAATGCAGAGGAGGATGGGAAGGGCTGCTTCGAGATTTACCACGCTTGT GATGACTCCTGCATGGAGTCAATTCGAAACAACACTTACGATCACTCACAGTACCGCGAGGAAGCACTTCTTAATAGGCTGAACATCAATCC TGTTACTCTCTCCAGCGGATATAAAGATATCATTCTGTGGTTCAGCTTCGGTGCCAGCTGCTTTGTGCTACTGGCAGTGGTCATGGGGCTCT TCTTCTTCTGCCTTAAAAATGGGAATATGCGTTGCACCATTTGCATT

2913	ATGTATAAAGTGGTAGTGATCATCGCTCTGCTGGGTGCCGTGCGAGGCCTTGATAAGATCTGTCTTGGCCACCATGCAGTGGCCAACGGGA CCATAGTTAAGACATTAACCAACGAACAGGAGGAGGTTACCAACGCTACCGAAACTGTCGAGAGCAAATCTCTGGGTAAGCTGTGTATGAA GGGCCGTAGCTACAATGATCTTGGTAACTGTCATCCTATCGGCATACTAATTGGGACCCCAGCCTGTGACTTGCACCTTACTGGTACTTGGG ACACCTTGATAGAACGTGAAAATGCCGTGGCTTACTGTTACCCGGGCGCCACTGTGAACGAGGAGGCCCTGCGGCAGAAAATTATGGAGT CACGTGGCATCTCTAAGATTTCAACAGGCTTCACCTACGGGACAAGTATTAATTCAGCTGGGACTACCAAAGCCTGTATGCCAAATGGGGG TAACTCTTTTTATGCCGAGCTTAAGTGGCTGGTTTCTAAAAATAAAGGACAGAATTTTCCACAAACTACAAACACTTATCGGAACACGGACAC CGCCGAGCACCTGATTATTTGGGGAATCCACCATCCATCTAGCACTCAGGAAAAAAATGATCTGTATGGTACTCAGAGCCTGAGCATTAGCG TAGGCAGCTCTACATATCAGAATAACTTTGTGCCTGTTGTAGGGGCTAGGCCTCAAGTGAACGGGCAGAGCGGTAGGATTGATTTCCACTG GACCCTGTTGCAGCCCGGAGACAATATAACCTTCTCGCATAACGGAGGGCTGATAGCCCCGTCCAGAGTGAGTAAGCTGATCGGTAGAGG ACTGGGAATTCAGAGCGAAGCCCCAATTGATAACGGCTGCGAGAGTAAGTGCTTCTGGAAGGGCGGTAGCATTAACACGAAGTTGCCATTT CAGAATCTCAGTCCACGTACGGTGGGCCAGTGTCCCAAATACGTCAACAAACGCTCTCTGATGCTTGCTACTGGGATGAGGAATGTCCCCG AGATAATGCATGGACGCGGGTTGTTTGGCGCCATCGCCGGGTTTATTGAGAATGGCTGGGAGGGCATGGTAGATGGATGGTACGGTTTTC GGCATCAGAATGCCCAGGGGACCGGACAGGCTGCGGATTACAAGTCTACACAGGCAGCCATCGATCAGATCACAGGCAAGTTGAACCGAC TGATCGAAAAGACAAACACAGAGTTCGAATCAATTGAGAGCGAGTTTTCCGAGATAGAGCATCAGATTGGCAACATTATTAATTGGACTAAA GATAGCATTACCGATATATGGACCTATCAGGCAGAACTGCTCGTCGCAATGGAGAATCAGCACACTATCGATATGGCCGATTCAGAGATGCT AAACCTGTACGAACGCGTTCGCAAGCAACTCAGACAGAATGCTGAGGAGGATGGCAAGGGTTGTTTCGAGATCTACCATGCGTGCGACGAT AGTTGTATGGAGAGCATTCGGAATAATACCTACGATCACTCCCAGTACAGGGAAGAGGCTTTGTTAAATAGACTGAATATAAACCCTGTCAA GCTATCATCAGGTTACAAAGACATCATTCTTTGGTTCTCTTTTGGCGCTTCCTGTCTGATCCTTCTAGCCGTGGTGATGGGTCTCGTTTTCTT TTGCCTCAAAAATGGAAACATGCGTTGTACGATATGTATA
2914	ATGTATAAGATAGTTGTGATAATTGCGCTGTTGGGGGCCGTGAAGGGACTCGACAAAATCTGCCTCGGCCACCATGCAGTCGCCAACGGTA CTATTGTGAAAACCCTAACCAACGAACAGGAGGAGGTTACTAATGCCACCGAGACTGTCGAATCAACCGGCATCAACCGCCTCTGTATGAA AGGCCGCAAACACAAGGATCTGGGAAACTGCCACCCCATCGGCATGTTAATAGGCACCCCCGCATGCGATCTGCACCTGACAGGAATGTG GGACACCTTGATCGAGCGTGAAAACGCGATCCCTTATTGTTACCCTGGCGCGACTGTGAATGTCGAGGCTCTGAGACAGAAGATTATGGAA AGCGGTGGCATTAATAAGATTAGTACTGGATTTACTTACGGTTCCAGCATAAATTCTGCAGGGACCACTCGCGCGTGCATGCGCAATGGTG GCAATAGTTTCTACGCTGAACTTAAGTGGCTTGTATCTAAGTCGAAGGGACAAAATTTTCCACAGACGACCAACACATATAGGAACACAGATA CAGCAGAGCACCTTATCATGTGGGGCATTCATCATCCGAGCAGTACTCAGGAGAAGAACGACCTGTACGGGACTCAGTCTTTGAGCATTAG CGTGGGAAGCAGCACTTACCGGAACAATTTTGTTCCTGTAGTGGGCGCTCGTCCTCAGGTCAATGGGCAGTCAGGACGGATAGACTTTCAC TGGACACTCGTGCAGCCCGGCGATAACATTACTTTTTCACATAATGGGGGTCTGATAGCACCAAGTCGCGTCTCTAAACTGATTGGTAGGG GCCTGGGAATTCAGTCCGACGCTCCCATCGACAATAATTGCGAATCCAAATGCTTTTGGCGGGGAGGGTCTATTAACACCAGGTTGCCATTT CAGAATCTGTCCCCGCGGACAGTTGGCCAGTGCCCCAAGTATGTCAACCGCAGGTCCCTGATGCTCGCTACAGGAATGCGCAATGTGCCA GAGTTGATCCAGGGACGGGGCTTGTTCGGTGCAATTGCAGGCTTTCTGGAAAACGGCTGGGAGGGGATGGTCGATGGGTGGTACGGCTTT CGCCACCAGAACGCCCAGGGAACAGGCCAGGCCGCCGATTACAAATCTACCCAGGCTGCCATCGATCAAATTACAGGGAAATTAAACCGA CTAGTCGAGAAGACCAACACTGAATTTGAATCCATTGAATCTGAGTTTTCTGAGATAGAGCACCAGATTGGAAATGTTATTAATTGGACAAAG GATAGCATCACAGACATATGGACATACCAAGCCGAATTGCTCGTGGCGATGGAAAACCAGCACACAATTGATATGGCCGACAGCGAAATGC TAAACCTTTACGAGAGAGTCCGAAAGCAGCTAAGGCAGAATGCCGAGGAAGACGGTAAAGGCTGTTTCGAAATCTACCATGCATGCGACGA TTCATGCATGGAAAGTATACGCAACAATACCTATGATCATTCCCAGTACCGCGAAGAGGCCTTGCTGAACCGACTCAACATCAACCCTGTAA CCCTGAGTTCTGGGTATAAGGACATCATCCTGTGGTTTAGTTTTGGAGCATCGTGCTTTGTGCTGCTCGCTGTGGTGATGGGATTGTTCTTC TTTTGCTTAAAGAACGGCAACATGAGGTGCACCATCTGTATA
2915	ATGTATAAGATTGTCGTCATCATCGCCCTACTGGGCGCGGTGAAAGGGCTAGATAAGATATGTCTCGGGCACCATGCAGTGGCTAACGGAA CAATTGTTAAAACCTTAACGAATGAACAGGAGAAGGTTACCAACGCTACTGAAACAGTTGAGTCAACTGGCCTCAACAGACTGTGCATGAAG GGTCGCAAACACAAAGATTTAGGCAACTGCCACCCCATTGGTATGTTAATCGGAACCCCAGCCTGTGATCTCCATCTGACCGGCACATGGG ATACGATAATTGAGAGGGAGAATGCTATTGCCTACTGTTACCCTGGCGCCACCGTCAATGAAGAGGCCCTCAGACAGAAGATCATGGAGTC CGGGGGTATCGACAAAATTTCCACCGGTTTCACCTACGGTAGCTCCATCAATAGTGCCGGCACTACACGCGCATGCATGCGCAATGGGGG CAACTCATTCTACGCGGAGCTAAAGTGGCTCGTTAGTAAGAGCAAAGGACAGAACTTCCCACAGACTACTAACACTTATAGGAACACTGATA CGGCAGAACATTTAATTATGTGGGGCATACACCATCCTAGCTCAACGCAGGAGAAGAATGACCTATACGGTACCCAATCCCTCTCCATCAGT GTGGGGAGCTCTACTTATAGAAACAACTTCGTGCCCGTCGTAGGCGCTCGACCTCAGGTCAATGGACAGTCTGGCAGAATCGACTTCCACT GGACTCTGGTACAACCTGGCGATAATATTACCTTTAGCCACAATGGCGGCTTAATAGCACCATCTCGTGTGAGCAAGTTGATTGGGCGAGG ATTGGGCATTCAGTCGGACGCGCCCATTGATAACAACTGTGAATCCAAATGCTTCTGGCGTGGAGGGTCAATCAATACCCGCCTCCCCTTC CAGAACCTGTCCCCGAGGACTGTGGGCCAATGTCCCAAATATGTGAACAAGAGGTCTCTGATGCTGGCAACCGGGATGAGAAATGTGCCC GAACTGATCCAGGGACGCGGGCTATTCGGGGCTATAGCAGGTTTTCTCGAAAATGGATGGGAGGGCATGGTGGACGGGTGGTACGGCTTT CGCCACCAGAATGCACAGGGAACCGGCCAGGCCGCAGATTACAAATCCACCCAGGCGGCCATCGATCAGATAACCGGCAAATTGAACAGG CTGGTTGAAAAAACAAATACAGAATTCGAATCCATTGAGTCTGAGTTTAGCGAGATAGAGCACCAGATAGGCAATGTCATAAACTGGACCAA AGATTCCATTACAGATATCTGGACATATCAGGCAGAGCTGCTCGTGGCAATGGAGAATCAGCATACTATTGATATGGCAGACAGCGAGATGC TCAATCTGTATGAACGGGTAAGGAAACAGCTGCGTCAGAATGCTGAGGAGGACGGTAAGGGGTGTTTTGAAATATACCACGCTTGCGATGA CAGTTGCATGGAATCTATTCGCAACAATACGTACGACCACTCTCAGTACCGCGAGGAGGCCCTTCTTAACAGGCTGAACATTAATCCCGTCA CACTAAGCTCCGGTTATAAGGACATCATTCTTTGGTTTTCATTTGGCGCTAGCTGCTTCGTTCTGCTTGCCGTGGTTATGGGCCTGGTCTTTT TCTGCCTTAAGAACGGAAATATGAGGTGTACAATCTGCATT

2916	ATGTACAAGATAGTGGTCATTATTGCTCTGTTGGGAGCTGTGAAAGGGCTTGATAAGATTTGCCTGGGCCACCACGCGGTGGCGAATGGGA CGATCGTGAAAACTCTCACAAACGAGCAGGAAGAAGTGACTAACGCTACAGAGACCGTGGAAAGCACGGGGATCAATAGGCTGTGCATGA AGGGCCGGAAGCATAAGGATCTAGGCAACTGCCACCCAATTGGCATGCTCATCGGGACGCCCGCATGCGATCTGCACCTGACAGGGATGT GGGACACCCTTATCGAGCGTGAGAACGCCATCGCCTATTGCTACCCCGGGGCAACAGTGAATGTGGAAGCACTGAGGCAGAAGATCATGG AGAGCGGGGGAATCAACAAGATATCCACAGGCTTCACATATGGATCATCAATCAACAGCGCCGGCACCACCCGGGCATGTATGCGCAACG GGGGCAATAGTTTCTACGCTGAGCTCAAATGGCTTGTTAGTAAGTCTAAGGGGCAGAACTTTCCACAGACTACGAATACGTACCGCAATACG GATACTGCAGAACACTTAATAATGTGGGGAATCCACCACCCAAGTTCCACCCAGGAGAAAAACGATTTGTACGGAACCCAATCCCTGTCAAT ATCCGTGGGAAGCTCTACCTATAGGAACAATTTCGTCCCGGTCGTGGGAGCTCGCCACAGGTGAACGGACAGAGTGGAAGGATCGACTT CCACTGGACGCTCGTGCAACCAGGTGATAATATCACGTTCAGTCACAACGGCGGACTGATTGCTCCAAGCCGCGTCAGCAAGCTGATAGG CCGGGGGCTGGGGATCCAGTCAGACGCCCCTATTGACAATAATTGTGAGTCCAAATGTTTCTGGAGAGGTGGCTCTATCAATACCCGCCTT CCCTTTCAGAACTTGTCCCCACGGACAGTGGGACAGTGCCCCAAGTATGTCAATCGCAGGTCACTGATGCTGGCAACCGGGATGAGGAAC GTTCCCGAACTCATTCAAGGCCGGGGTCTATTTGGCGCAATCGCCGGCTTCCTGGAAAATGGCTGGGAAGGAATGGTGGACGGGTGGTAT GGTTTCCGGCACCAGAATGCTCAGGGCACAGGCCAGGCTGCTGATTATAAGTCTACTCAGGCAGCCATTGACCAGATCACCGGTAAGCTCA ATAGATTGGTAGAAAAAACAAACACGGAGTTCGAGAGCATCGAGTCTGAATTTTCCGAGATCGAACATCAGATCGGGAACGTCATCAATTGG ACCAAGGACTCTATTACGGACATTTGGACTTACCAAGCTGAGCTCCTGGTGGCAATGGAGAACCAGCATACCATTGATATGGCCGACTCTG AGATGCTGAACTTATACGAACGGGTGCGAAAGCAACTGAGGCAGAACGCCGAGGAGGATGGTAAGGGCTGCTTTGAGATATATCATGCCT GTGATGATTCTTGCATGGAGAGCATCCGTAATAATACTTACGACCATAGCCAGTATCGAGAGGAAGCGCTACTCAATCGATTAAATATCAAC CCTGTGACCCTATCTTCGGGTTACAAGGACATAATTCTGTGGTTCTCATTCGGCGCATCTTGTTTCGTATTGCTCGCAGTGGTGATGGGCCT TTTTTTCTTCTGTCTGAAAAATGGTAACATGCGCTGTACTATATGCATT
2917	ATGTACAAAATAGTGGTGATTATCGCCCTCCTGGGTGCTGTCAAGGGCCTGGATAAGATCTGTCTTGGCCATCACGCTGTCGCAAATGGCA CTATAGTCAAAACTCTGACAAACGAGCAGGAGGAGGTGACCAACGCAACCGAGACGGTGGAATCTACTGGAATTAACAGGCTGTGTATGAA GGGTAGGAAGCATAAGGATCTAGGTAATTGCCATCCCATCGGTATGCTGATCGGAACTCCCGCGTGCGACTTACATCTCACTGGAATGTGG GACACGTTGATCGAGCGGGACAATGCTATTGCGTACTGCTACCCCGGCGCTACGGTGAACGTCGAGGCTTTGCGTCAGAAGATCATGGAG AGCGGGGGTATCAACAAAATAAGTACCGGATTCACATACGGAAGCAGTATCAATTCCGCCGGAACAACCCGCGCATGCATGCGGAACGGT GGTAATTCCTTCTATGCTGAACTCAAGTGGCTCGTGTCAAAGAGCAAGGGACAAAACTTTCCCCAAACGACCAATACCTATAGAAATACAGA CACTGCTGAGCATCTGATCATGTGGGGCATTCACCATCCATCATCTACTCAGGAAAAAAACGACCTGTACGGTACCCAGAGCCTGTCTATTA GCGTCGGATCATCGACCTACAGGAATAACTTCGTGCCAGTCGTTGGGGCCCGCCCTCAGGTGAACGGCCAGAGCGGACGGATTGATTTTC ACTGGACCCTGGTCCAGCCAGGTGATAACATCACCTTCTCACATAACGGAGGCCTGATCGCCCCCAGCCGAGTTAGTAAACTAATCGGAAG GGGACTTGGTATCCAGAGTGACGCACCCATCGACAACAACTGTGAGTCTAAGTGTTTTTGGCGAGGCGGGAGTATCAATACTAGACTGCCG TTTCAGAATTTGAGTCCCAGAACAGTGGGCCAGTGCCCTAAGTACGTGAACCGGAGGTCTCTGATGCTCGCAACCGGGATGCGGAATGTTC CCGAGCTGATCCAGGGACGCGGTCTCTTCGGTGCCATCGCTGGGTTTCTCGAAAACGGCTGGGAGGGAATGGTGGATGGTTGGTACGGCT TCAGACATCAAAACGCTCAAGGAACTGGCCAGGCCGCTGACTATAAGTCAACTCAGGCTGCAATCGATCAGATCACCGGGAAACTGAATCG CCTCGTGGAGAAGACAAACACCGAGTTCGAGTCAATCGAGTCCGAGTTCTCCGAGATCGAACACCAAATAGGCAACGTTATAAACTGGACG AAGGACTCCATTACGGACATCTGGACATACCAGGCCGAGCTCCTCGTTGCTATGGAGAACCAACACACAATCGATATGGCTGATTCCGAGA TGCTCAATCTCTACGAGAGGGTGAGGAAACAGTTGAGGCAGAATGCAGAAGAGGATGGAAAGGGATGCTTTGAAATCTACCATGCCTGCGA CGACTCATGCATGGAGTCCATTAGAAATAACACTTATGACCATTCGCAATACACAGAAGAAGCTCTTCTCAACAGGCTCAACATCAACCCGG TCACCCTTAGCTCCGGCTACAAGGACATTATCCTCTGGTTCTCTTTCGGGGCATCATGCTTTGTTCTCCTGGCAGTGGTGATGGGTCTCTTC TTCTTCTGTCTCAAAAACGGCAACATGAGATGTACGATTTGTATC
2918	ATGTATAAGATCGTGGTGATCATAGCGCTCCTGGGCGCCGTGAAAGGACTGGATAAAATTTGTCTGGGGCATCATGCAGTCGCCAACGGAA CCATCGTAAAGACTTTGACTAATGAGCAGGAAAAGGTTACCAATGCAACCGAAACCGTGGAGAGCACCGGTCTGAACCGCCTCTGCATGAA GGGCCGTAAACACAAAGATCTGGGCAATTGTCATCCCATCGGAATGCTAATCGGCACTCCTGCCTGTGACCTACACTTGACCGGGACTTGG GATACCATTATAGAGAGAGAGAATGCTATTGCCTACTGTTACCCCGGGGCAACCGTGAACGAAGAAGCTTTGAGACAGAAAATTATGGAGA GTGGCGGAATAGATAAGATTAGCACAGGGTTCACCTACGGAAGCTCTATTAATAGTGCGGGCACCACAAGGGCATGCATGCGGAACGGCG GTAACTCTTTCTATGCTGAGTTGAAGTGGCTCGTGAGCAAGTCTAAAGGGCAGAACTTTCCACAAACCACCAACACTTACAGAAACACGGAT ACTGCCGAGCACCTCATCATGTGGGGCATCCACCATCCTTCATCAACACAAGAGAAGAACGACTTATATGGTACTCAGAGTCTGTCGATTAG CGTCGGCTCTTCAACGTACAGGAACAATTTCGTTCCTGTAGTCGGGGCCCGGCCACAAGTAAATGGACAGAGCGGGAGGATTGATTTCCAT TGGACGCTGGTCCAGCCCGGAGATAACATAACTTTCTCTCATAACGGAGGACTCATAGCTCCATCAAGGGTGAGCAAACTGATTGGCAGGG GATTGGGAATCCAGTCTGACGCTCCCATCGACAACAATTGTGAGAGTAAGTGCTTTTGGCGGGGCGGTTCCATCAACACACGACTGCCTTT CCAGAATCTGAGTCCCAGAACTGTGGGGCAGTGCCCAAAGTATGTGAATAAAAGATCTTTGATGTTAGCTACTGGAATGAGAAATGTGCCTG AGCTAATACAGGGTAGAGGTCTGTTCGGAGCAATAGCAGGGTTCCTGGAGAATGGGTGGGAAGGCATGGTCGACGGATGGTACGGTTTCC GGCATCAGAACGCACAAGGTACAGGGCAGGCGGCTGATTACAAATCTACACAAGCCGCTATTGACCAAATCACCGGCAAATTGAACCGGCT CGTGGAGAAGACGAATACGGAGTTTGAATCTATTGAGAGTGAGTTCAGTGAGATTGAACATCAGATTGGCAACGTTATAAATTGGACCAAGG ACAGCATAACTGATATATGGACCTATCAAGCCGAGCTTCTCGTGGCCATGGAAAATCAGCATACTATCGACATGGCTGACTCTGAGATGCTC AACTTATACCAACGTGTAAGGAAGCAGCTGCGGCAAAACGCCCAGGAAGACGGTAAGGGCTGCTTCGAGATTTATCACGCGTGCGATGAC TCTTGTATGGAGTCCATCAGAAACAACACCTACGATCACTCACAGTATAGAGAAGAGGCTTTGTTGAATCGGCTCAACATCAATCCAGTTAC CCTTTCAAGTGGTTACAAGGACATCATCCTATGGTTTAGCTTCGGTGCCTCTTGTTTCGTGTTGCTGGCGGTCGTTATGGGGTTGGTCTTTTT TTGCTTGAAAAATGGTAATATGCGGTGTACCATTTGCATC

2919	ATGTACAAGATTGTGGTGATCATCGCGCTGCTTGGTGCTGTGAAGGGACTCGACAAAATATGCCTGGGGCATCATGCTGTCGCCAATGGGA CAATTGTGAAGACATTGACGAACGAGCAGGAGGAAGTCACTAACGCAACAGAGACAGTTGAGAGCACCGGGATTAACCGACTGTGTATGAA GGGTCGGAAGCACAAGGACTTGGGTAACTGTCACCCGATCGGCATGTTAATCGGAACCCCTGCCTGTGACCTGCATCTGACCGGCATGTG GGACACACTGATAGAAAGGGAAAACGCCATTGCATACTGCTACCCCGGGGCTACGGTAAATGTGGAGGCTCTCCGGCAGAAGATAATGGA GTCAGGCGGCATCAACAACATCTCTACCGGGTTCACGTACGGCTCTTCTATCAATTCAGCTGGAACTACCCGCGCATGCATGAGAAACGGT GGGAACTCCTTTTACGCCGAACTGAAATGGCTAGTGTCCAAATCTAAGGGCCAGAATTTCCCACAGACCACTAATACTTACAGAAACACGGA TACAGCCGAACACTTGATAATGTGGGGGATTCATCACCCATCATCTACACAGGAAAAAAATGATTTGTACGGTACCCAGAGCCTTAGCATTT CTGTGGGCAGCTCTACCTACAGAAATAACTTCGTGCCAGTAGTAGGTGCTCGACCGCAGGTTAATGGCCAAAGCGGTAGAATCGATTTTCA CTGGACATTGGTGCAGCCGGGCGACAACATAACCTTTTCCCATAATGGGGGACTTATTGCACCTTCCAGGGTGTCCAAACTCATAGGCCGC GGCCTCGGCATTCAAAGCGATGCTCCCATTGACAATAATTGTGAATCAAAGTGCTTCTGGCGTGGAGGAAGTATAAACACACGGCTGCCCT TCCAGAACTTAAGCCCAAGAACCGTGGGGCAGTGCCCTAAGTACGTTAATAGGCGGTCGCTTATGCTTGCAACAGGTATGAGGAACGTCCC GGAGCTTATCCAAGGCAGGGGACTTTTCGGAGCTATCGCCGGTTTTTTAGAGAACGGATGGGAAGGAATGGTGGACGGTTGGTATGGATT CCGGCACCAAAATGCTCAGGGCACCGGGCAGGCGGCTGACTACAAATCTACACAGGCGGCTATCGATCAGATTACGGGAAAGCTGAATAG GTTAGTGGAAAAAACAAACACAGAGTTTGAGAGCATCGAAAGCGAGTTTTCTGAGATAGAGCATCAGATCGGTAATGTAATCAACTGGACCA AAGACAGCATCACAGACATATGGACGTACCAGGCTGAGCTTCTAGTCGCCATGGAAAACCAGCACACTATTGATATGGCAGATTCTGAGAT GCTTAATCTGTATGAACGAGTGCGAAAACAGCTGCGCCAGAACGCCGAAGAGGATGGGAAGGGCTGTTTCGAAATCTACCACGCCTGCGA CGACTCCTGCATGGAAAGCATTAGAAATAATACCTACGATCATTCGCAATATCGGGAAGAGGCCCTCCTGAATCGCCTAAATATCAATCCTG TTACACTAAGCAGCGGCTACAAAGACATTATTTTATGGTTCTCCTTCGGCGCCTCCTGCTTCGTACTATTAGCCGTAGTTATCGGTCTGTTCT TCTTTTGCCTTAAGAACGGTAACATGAGATGCACAATCTGTATA
2920	ATGTACAAAATTGTTGTGATCATTGCCCTCCTGGGGGCAGTGAAGGGACTGGACAAAATCTGCCTGGGTCACCACGCCGTGGCGAACGGC ACCATTGTGAAGACCCTGACCAATGAACAGGAAGAGGTGACCAATGCTACCGAGACTGTAGAGAGTACAGGCATCAATAGGTTATGCATGA AAGGCCGCAAACACAAGGACTTGGGAAACTGCCACCCTATAGGCATGTTGATCGGGACGCCAGCTTGCGACTTGCACCTTACAGGAATGTG GGATACCTTGATAGAAAGGGAAAACGCTATCGCTTACTGCTACCCGGGCGCTACCGTCAATGTGGAAGCACTGCGACAGAAAATTATGGAG AGTGGTGGCATAAATAAAATATCAACTGGGTTCACATACGGAAGCTCAATCAATAGCGCGGGCACAACACGAGCCTGCATGCGTAACGGAG GAAACAGCTTTTATGCAGAGCTCAAATGGTTAGTTTCAAAATCTAAGGGGCAGAACTTTCCTCAAACCACTAACACGTATCGGAATACCGACA CAGCAGAGCACTTAATAATGTGGGGCATACATCATCCCTCAAGCACACAGGAAAAGAACGACCTGTACGGAACGCAGTCCTTATCAATCTCT GTGGGGTCCAGCACATACCGAAACAACTTTGTACCAGTCGTGGGTGCTCGCCCACAAGTGAACGGCCAGTCTGGGCGCATCGACTTTCAC TGGACTTTGGTTCAGCCTGGTGATAATATCACCTTTAGCCATAATGGCGGACTCATAGCGCCAAGCAGGGTGTCCAAACTTATCGGACGTG GGTTGGGTATTCAGAGCGACGCACCAATCGACAATAACTGCGAGTCAAAGTGTTTTTGGAGAGGCGGGTCAATCAACACCCGCCTGCCCTT CCAGAACCTGTCTCCCCGTACGGTGGGGCAGTGTCCAAAGTACGTGAATAGGCGGAGCCTGATGCTGGCCACAGGGATGCGCAACGTGC CTGAGCTCATTCAAGGCCGGGGCTTGTTTGGAGCCATCGCGGGTTTTCTCGAAAACGGGTGGGAAGGGATGGTCGATGGCTGGTACGGCT TTCGGCACCAGAATGCTCAGGGAACGGGGCAGGCTGCCGACTACAAGTCAACGCAGGCGGCCATCGATCAAATCACCGGCAAGCTCAACA GGCTCGTGGAAAAAACCAATACGGAGTTCGAAAGTATCGAAAGCGAGTTCAGCGAAATTGAGCACCAGATCGGCAATGTAATCAATTGGAC TAAGGATAGCATCACCGATATCTGGACATACCAAGCTGAGCTGTTAGTCGCCATGGAGAATCAGCATACAATAGACATGGCTGATAGTGAAA TGTTGAATCTGTACGAGAGGGTTAGGAAACAGCTGCGCCAGAATGCTGAGGAAGACGGAAAGGGCTGCTTTGAAATCTATCACGCTTGCGA CGATTCCTGTATGGAGTCCATACGAAATAATACCTACGACCATTCCCAGTACCGCGAAGACGCTCTCCTGAATCGGCTGAACATCAACCCTG TGACACTTAGCTCGGGCTATAAGGACATAATTTTGTGGTTCTCTTTTGGGGCATCATGTTTTGTCTTGCTGGCCGTCGTTATGGGGCTCTTCT TCTTCTGTCTAAAAAATGGAAACATGCGGTGCACCATTTGTATT
2921	ATGTATAAAATCGTGGTGATCATTGCTCTGTTAGGCGCCGTGAAGGGCCTGGACAAGATTTGCCTAGGACATCATGCCGTCGCTAACGGCA CTATCGTGAAGACTCTGACCAATGAGAAAGAGGAGGTGACTAACGCAACAGAAACAGTTGAGTCAACTGGCCTGAACCGGCTCTGCATGAA AGGGACAAAGCATAAAGACCTAGGCAATTGCCATCCTATTGGTATGCTCATTGGCTCTCCCGCCTGTGACCTGCATCTTACTGCAACGTGG GACACCCTGATTGAAAGAGAAAACGCCATCGCGTATTGCTACCCGGGGGCCACCGTGAACGGAGAAGCGTTACGCCAAAAGATCATGGAG TCAGGCGGCATCGATAAGATCAGCACTGGCTTTACCTACGAAAGTAGCATCAACAGCGCTGGAACCACCCGCGCCTGTATGAGAAACGGG GGTAATTCCTTCTATGCGGAGCTGAAATGGCTTGTGTCTAAATCCAAAGGCCAGAACTTTCCACAGACAACGAACACTTATCGGAATACTGA CACAGCAGAGCATCTTATCATGTGGGGGATCCACCATCCTAGTTCCACGCAGGAGAAGAATGACCTGTATGGGACACAAAGCCTGTCTATT AGCGTGGGATCATCAACTTACAGGAACAATTTTGTCCCTGTGGTGGGCGCCAGGCCCCAGGTAAACGGACAGAGTGGACGCATCGACTTT CATTGGACTCTCGTGCAGCCTGGGGATAATATTACCTTCAGCCATAATGGTGGCCTGATAGCACCCTCAAGAGTGTCAAAACTCATCGGTCG CGGATTGGGGATTCAATCTGACGCCCCTATAGATAACAATTGCGAGAGCAAGTGCTTTTGGAGAGGAGGCAGCATCAACACCCGCCTTCCC TTTCAGAATCTGAGTCCCCGGACGGTTGGGCAGTGTCCTAAATATGTCAATAAGCGGTCACTCATGTTGGCAACTGGCATGCGCAACGTAC CTGAGCTCATGCAGGGAAGGGGCCTCTTTGGAGCTATTGCAGGGTTCCTGGAGAACGGCTGGGAGGGGATGGTAGACGGATGGTACGGA TTCCGGCATCAAAATGCGCAAGGCACTGGTCAAGCTGCTGATTATAAGTCCACCCAGGCCGCTATTGATCAGATTACCGGAAAGCTGAACC GGCTGGTGGAAAAGACCAATACAGAATTTGAGAGCATTGAATCGGAGTTTAGCGAAATCGAACATCAAATTGGGAACGTCATAAACTGGACA AAAGACAGCATCACCGATATCTGGACATACCAGGCGGAGCTGCTAGTGGCGATGGAGAATCAGCACACGATTGATATGGCCGACTCTGAGA TGCTGAATCTGTATGAGCGGGTTAGAAAGCAGCTGAGGCAAAACGCAGAAGAAGATGGTAAAGGATGCTTTGAGATCTATCATGCCTGCGA TGACTCATGTATGGAATCTATCCGTAACAATACATACGATCACAGTCAGTATCGGGAGGAAGCTCTGCTGAATAGGCTAAACATCAATCCCG TGACCCTCTCGAGTGGCTATAAGGATATCATCTTGTGGTTCTCTTTCGGCGCCAGCTGTTTCGTGCTACTGGCCGTCGTGATGGGGCTGGT ATTCTTCTGCCTGAAGAACGGTAATATGCGCTGCACCATATGTATT

2922	ATGTACAAGATCGTGGTCATCATTGCACTGCTTGGCGCTGTGAAAGGATTAGACAAGATTTGCCTCGGCCACCACGCCGTCGCCAATGGGA CCATCGTTAAGACATTAACAAATGAACAAGAAGAGGTAACCAACGCCACCGAGACCGTGGAGTCGACAGGCATCAACCGGTTATGCATGAA AGGGCGGAAACACAAAGATCTGGGGAACTGCCATCCAATCGGAATGCTCATTGGCACCCCAGCGTGCGATCTCCATCTCACTGGCATGTG GGATACATTAATTGAGAGGGAGAATGCAATCGCATATTGCTACCCTGGCGCCACTGTCAACGTCGAGGCCCTGCGCCAGAAAATAATGGAA TCAGGTGGCATAAACAAGATTTCCACCGGATTTACTTATGGCTCTTCCATCAACAGCGCTGGCACCACACGGCGCTTGTATGCGAAACGGGG GCAACTCATTTTATGCAGAACTGAAGTGGCTTGTCTCAAAAAGTAAGGGTCAGAACTTTCCACAGACAACAAATACATACAGGAATACCGAC ACTGCAGAGCATCTGATCATGTGGGGCATCCACCATCCTAGTTCGACACAGGAGAAGAACGACCTGTATGGCACCCAATCACTGAGTATCT CTGTGGGGAGTAGTACTTATCGGAACAATTTTGTGCCAGTCGTGGGCGCAAGACCACAAGTCAATGGCCAGAGTGGCCGTATCGACTTCCA TTGGACCCTGGTACAGCCTGGTGATAACATCACCTTTAGCCACAACGGAGGACTGATTGCACCGTCAAGAGTTTCAAAACTGATCGGCCGA GGCCTGGGAATCCAGAGCGACGCTCCCATCGATAATAATTGTGAAAGTAAGTGCTTCTGGCGGGGTGGATCCATCAACACCCGTCTACCCT TTCAGAATCTGTCACCCCGGACCGTCGGACAGTGCCCAAAATACGTGAACAGGCGCAGTTTGATGTTAGCTACAGGGATGAGAAACGTGCC AGAACTGATTCAGGGACGCGGACTGTTCGGCGCCATCGCCGGCTTCCTCGAAAACGGTTGGGAGGGTATGGTGGACGGTTGGTATGGCTT CAGGCATCAGAACGCCCAGGGGACAGGCCAGGCCGCTGACTATAAGAGCACTCAGGCGGCTATCGACCAGATCACTGGGAAGCTCAATC GTCTCGTCGAGAAAACGAATACCGAGTTTGAGAGCATCGAAAGTGAATTCTCTGAGATAGAGCACCAGATCGGTAATGTGATCAATTGGACA AAGGACTCGATTACTGACATCTGGACGTACCAAGCCGAGCTGCTCGTCGCCATGGAAAATCAGCATACCATTGATATGGCTGACTCAGAGA TGCTTAACTTGTACGAACGGGTCCGGAAACAACTGAGACAGAATGCCGAAGAGGACGGGAAAGGCTGTTTTGAAATCTATCACGCCTGTGA TGATTCTTGTATGGAAAGCATTAGAAACAATACATACGACCATTCACAGTACAGGGAGGAGGCTTTACTCAATCGCCTGAACATCAATCCCG TAACATTGAGCTCTGGATATAAAGACATTATCCTCTGGTTCTCCTTTGGTGCCTCTTGTTTTGTACTCCTGGCAGTGGTTATGGGTCTCTTCTT CTTTTGTCTAAAGAACGGTAATATGAGGTGTACCATTTGCATA
2923	ATGTACAAGATCGTGGTGATTATTGCCCTGTTGGGGGCCGTGAAGGGCTTGGACAAGATATGCCTTGGTCACCATGCTGTGGCCAACGGCA CTATCGTCAAAACACTGACAAACGAGCAGGAAGAGGTGACCAACGCCACGGAGACTGTAGAAAGCACCGGGATTAACCGACTATGTATGAA AGGCCGCAAACACAAGGACCTCGGGAACTGCCATCCCATCGGCATGTTAATAGGCACCCCCGCTTGCGATCTGCACCTGACAGGGATGTG GGATACCCTAATTGAGCGCGAGAACGCAATTGCCTATTGTTACCCAGGAGCCACAGTAAATGTAGAAGCTTTAAGGCAGAAGATTATGGAGA GTGGAGGGATTAACAAGATCAGCACGGGCTTCACATACGGTTCCTCCATCAATTCTGCTGGAACCACCAGGGCATGCATGCGTAATGGGGG AAACTCATTCTACGCAGAGCTAAAGTGGCTGGTTTCCAAGTCTAAGGGACAGAACTTCCCGCAGACAACGAATACTTATCGGAACACCGACA CTGCCGAACATTTAATCATGTGGGGGATCCACCACCCCAGTTCTACCCAGGAGAAAAATGATCTCTACGGTACTCAGTCCCTGTCCATCTCC GTTGGTTCCTCCACCTATCGGAATAATTTTGTCCCTGTTGTGGGAGCACGGCCACAGGTGAATGGCCAGTCCGGGAGGATAGACTTTCACT GGACCCTTGTTCAGCCCGGGGACAATATCACTTTCTCACATAATGGGGGCCTGATTGCTCCTTCCAGAGTGAGTAAATTGATAGGCCGGGG GCTGGGCATCCAATCAGATGCACCCATTGATAACAACTGCGAGTCCAAATGCTTCTGGCGCGGAGGCAGCATCAATACCCGACTGCCCTTC CAGAACCTGTCTCCTAGGACAGTTGGCCAGTGTCCCAAATATGTAAACCGCAGATCACTGATGCTGGCTACCGGTATGAGAAACGTCCCAG AGCTAATTCAAGGCCGAGGACTTTTTGGCGCTATTGCAGGATTCCTTGAGAACGGGTGGGAGGGCATGGTCGACGGCTGGTACGGGTTCA GACACCAAAACGCTCAGGGAACAGGCCAAGCCGCGGACTATAAATCAACACAAGCGGCCATTGATCAGATAACCGGCAAGCTGAATAGGC TGGTGGAAAAGACTAACACTGAGTTTGAAAGCATCGAGTCGGAATTCTCTGAGATTGAACACCAGATTGGAAATGTAATCAACTGGACAAAG GATTCCATCACCGACATCTGGACTTATCAAGCAGAACTGTTAGTGGCCATGGAAAATCAGCATACCATCGACATGGCCGACAGCGAGATGC TGAATCTGTACGAACGGGTTCGTAAACAGCTGAGGCAGAACGCAGAGGAGGACGGTAAGGGATGCTTTGAAATTTATCACGCTTGTGACGA TAGTTGTATGGAGAGCATTAGGAACAACACTTACCACCACTCTCAGTATAGGGAAGAAGCACTCCTTAACAGGCTGAACATCAACCCTGTGA CGCTCTCTTCCGGGTATAAAGACATCATCCTGTGGTTTTCATTTGGCGCTTCATGTTTTGTACTGCTAGCAGTGGTGATGGGTCTCTTCTTTT TTTGCCTCAAAAACGGGAACATGAGGTGCACTATTTGCATC
2924	ATGTACAAGATCGTAGTGATAATCGCTCTACTAGGAGCTGTCAAAGGTCTCGATAAGATATGTCTGGGGCACCATGCGGTAGCCAATGGAA CGATCGTGAAAACACTTACTAACGAGCAGGAGGAGGTCACCAACGCCACAGAAACGGTGGAATCTACGGGGATCAACCGCTTATGCATGAA GGGGAGGAAGCATAAGGATTTGGGCAATTGCCACCCGATTGGAATGCTGATTGGGACTCCTGCTTGCGATCTGCACTTAACCGGCATGTGG GATACGCTAATAGAGCGCGAAAACGCCATCGCCTACTGTTATCCCGGCGCGACAGTAAACGTAGAGGCACTAAGACAGAAGATTATGGAGT CCGGGGGAATAAACAAGATAAGCACGGGCTTTACCTATGGATCATCCATCAATTCTGCTGGAACAACACGCGCCTGCATGCGGAACGGAGG CAATTCTTTCTATGCAGAGCTGAAATGGCTCGTGTCCAAAAGTAAAGGTCAGAACTTCCCGCAGACTACAAACACTTATAGGAACACTGACA CAGCTGAGCATCTAATCATGTGGGGTATTCACCACCCAAGTTCAACCCAAGAAAAAAATGATCTCTACGGCACGCAGTCACTTTCCATCTCT GTGGGCTCTAGCACCTACAGAAACAACTTTGTCCCAGTGGTGGGAGCTCGGCCTCAAGTGAACGGACAGTCCGGTAGGATCGATTTTCATT GGACCCTCGTACAGCCCGGGGATAACATCACTTTTAGTCACAATGGTGGCCTGATTGCCCCCTCCAGGGTGTCTAAGTTGATTGGGCGCGG CCTAGGTATACAGAGTGACGCCCCCATAGATAATAATTGCGAGAGCAAATGTTTCTGGAGGGGCGGAAGTATCAATACTCGGCTGCCTTTTC AGAACCTTAGCCCACGTACTGTGGGCCAGTGTCCCAAGTACGTTAATCGAAGAAGCCTGATGCTTGCAACCGGAATGAGGAACGTGCCCGA ACTGATCCAAGGTCGTGGCTTATTCGGAGCCATCGCCGGATTCCTGGAGAACGGATGGGAAGGGATGGTGGATGGGTGGTACGGGTTCCG ACACCAGAACGCCCAGGGAACAGGTCAAGCCGCCGATTACAAATCAACACAAGCGGCCATCGATCAGATTACTGGAAAACTGAACAGGTTG GTGGAGAAGACTAACACAGAGTTCGAGTCAATCGAAAGCGAATTTAGCGAGATTGAGCACCAAATCGGGAACGTGATCAACTGGACAAAGG ATTCCATTACCGACATATGGACCTACCAGGCTGAGCTGTTGGTGGCAATGGAAAATCAGCATACTATCGATATGGCCGATTCTGAGATGCTG AATCTGTACGAGCGGGTCAGAAAGCAGCTGACACAAAATGCTGAGGAAGACGGTAAGGGTTGTTTCGAGATCTACCATGCCTGCGACGACT CTTGCATGGAGAGCATCAGGAACAACACATACGATCACTCCCAGTATAGAGAAGAAGCTCTGTTGAATAGGCTTAACATCAACCCAGTGACG CTTAGTAGCGGTTACAAAGATATCATTCTGTGGTTTAGTTTTGGGGCCTCATGTTTCGTGCTTTTGGCAGTAGTTATGGGGCTGTTTTTCTTC TGTCTCAAAAATGGTAATATGCGATGCACGATTTGTATT

2925	ATGTACAAAATCGTGGTCATTATCGCCCTGCTGGGCGCGGTGAAAGGCTTGGATAAAATCTGCCTCGGGCATCACGCCGTGGTGAATGGAA CGATCGTGAAGACTCTGACCAACGAACAGGAAGAGGTTACCAACGCAACGGAAACAGTCGAATCAACCGGACTCAATCGTTTATGTATGAA GGGCAGAAATCACAAGGACCTTGGAAACTGCCACCCAATAGGGATGCTTATAGGAACCCCAGCTTGCGACCTGCACCTAACCGGCACCTG GGACACTCTTATTGAGAGAGAGAACGCAATTGCTTACTGTTACCCCGGCGCAACAGTGAACGAAGAGGCCCTCAGGCAGAAGATTATGGAG TCTGGAGGGATAAACAAGATATCGACCGGTTTCACCTATGGATCCTCTATCAATTCTGCAGGGACAACCCGCGCCTGTATGAGAAATGGCG GAAACTCCTTTTATGCCGAACTGAAGTGGCTTGTTTCCAAAAGCAAAGGTCAGAACTTTCCTCAAACAACTAATACGTACAGGAACACCGATA CCGCCGAGCACCTGATCATGTGGGGAATCCATCACCCTAGCAGTACCCAAGAAAAAAACGACCTATATGGTACCCAGTCCCTCAGCATCTC AGTGGGCTCATCAACCTATCAAAACAACTTCGTGCCTGTTGTCGGCGCACGACCCCAGGTGAACGGTCAAAGCGGAAGGATCGATTTCCAT TGGACCCTGGTTCAGCCCGGCGACAATATCACATTTAGTCACAATGGCGGTCTGATTGCACCTTCCCGCGTGAGTAAACTGATCGGTCGGG GGCTTGGAATTCAAAGCGACGCCCCCATCGACAACAATTGTGAGAGCAAGTGTTTTTGGAGAGGCGGGTCAATAAACACCAGGCTGCCCTT TCAGAACCTAAGTCCCAGGACTGTAGGCCAGTGTCCCAAATACGTCAATAAGCGTTCACTAATGTTGGCCACGGGAATGAGAAACGTGCCC GAGCTGATGCAAGGCAGAGGCCTGTTCGGGGCCATCGCGGGGTTCATCGAGAACGGTTGGGAAGGCATGGTTGATGGTTGGTATGGATTT AGGCACCAGAACGCTCAGGGCACAGGACAGGCCGCTGACTATAAGAGTACTCAAGCCGCAATTGACCAGATCACCGGGAAGTTGAACCGA CTCATCGAGAAGACGAACACAGAATTCGAGAGCATAGAGTCAGAGTTCTCGGAGATCGAACACCAAATTGGCAACGTTATCAATTGGACTAA GGACTCGATCACTGACATCTGGACATACCAGGCTGAGCTGTTGGTGGCCATGGAAAACCAGCACACCATAGACATGGCCGACAGCGAGAT GCTGAATCTCTACGAAAGAGTTCGCAAACAGCTCAGGCAGAATGCCGAAGAAGACGGTAAAGGATGCTTTGAAATCTATCACGCTTGTGAC GATTCCTGCATGGAGAGCATCAGAAACAATACCTACGATCACAGCCAGTACCGGGAAGAAGCACTCCTGAATAGGCTGAACATAAATCCAG TAACTCTTTCGTCCGGATACAAAGACATAATACTCTGGTTTTCGTTTGCTGCATCCTGCTTCGTGCTGCTCGCAGTGGTGATGGGCTTGTC TTTTTCTGTTTGAAGAACGGAAATATGCGGTGCACCATATGCATC
2926	ATGTACAAGATCGTTGTGATAATTGCCTTGCTGGGCGCAGTCAAAGGACTCGATAAGATCTGTCTCGGCCATCACGCCGTTGCCAATGGAA CCATTGTGAAGACTCTCACTAACGAGAAAGAGGAGGTTACAAACGCTACCGAAACTGTGGAGAGTACTGGTCTGAACCGACTGTGTATGAA GGGCAGAAAGCATAAGGATCTGGGTAATTGTCACCCCATCGGAATGTTAATCGGTAGCCCAGCCTGTGATTTGCACCTTACAGGGACCTGG GATACACTCATAGAGAGGGAAAACGCCATCGCATATTGTTATCCCGGGGCCACCGTGAACGGAGAAGCTTTACGCCAGAAGATCATGGAGT CCGGGGGGATCGATAAGATCAGCACCGGATTCACCTACGAGAGTAGTATAAACTCAGCGGGCACCACCAGAGCGTGCATGCGGAATGGCG GCAATAGCTTTTATGCTGAGCTTAAGTGGTTGGTGTCCAAATCTAAGGGACAGAATTTCCCCCAGACCACCAATACGTACCGCAATACAGAC ACAGCCGAGCATTTAATCATGTGGGGAATCCACCACCCATCGTCTACCCAGGAAAAGAACGACTTGTATGGGACTCAGAGTCTTTCCATCTC CGTCGGATCCTCAACCTACCGGAACAATTTCGTGCCCGTGGTAGGGGCGCGTCCCCAAGTTAACGGCCAATCTGGCAGGATAGATTTCCAT TGGACTCTGGTGCAGCCTGGCGACAACATTACCTTCTCGCATAACGGGGGACTGATCGCCCCCTCTCGCGTTTCTAAGCTGATCGGCCGC GGTCTGGGCATCCAGTCTGACGCTCCCATTGACAATAACTGCGAATCAAAGTGCTTTTGGAGAGGAGGGTCTATTAATACTAGGTTGCCCTT TCAGAATCTGTCCCCCCGAACCGTCGGGCAGTGTCCGAAATATGTCAACAAACGCTCATTAATGCTTGCTACTGGCATGAGGAATGTCCCT GAGTTGATGCAGGGCAGGGGGCTGTTCGGCGCCATAGCAGGGTTTCTCGAGAACGGGTGGGAGGGCATGGTAGATGGGTGGTACGGCTT TAGGCATCAGAATGCTCAAGGTACCGGCCAAGCCGCCGACTACAAATCCACACAGGCCGCAATTGACCAGATCACAGGTAAATTGAATCGG CTTGTGGAGAAGACCAATACTGAATTCGAAAGTATTGAATCCGAATTTTCTGAGATAGAGCACCAAATCGGTAACGTGATCAACTGGACTAA AGATTCTATCACCGACATTTGGACATATCAGGCGGAATTGCTGGTGGCTATGGAGAACCAGCACACGATTGATATGGCTGACAGCGAGATG CTGAACCTTTACGAGCGAGTGCGTAAGCAACTGAGGCAGAACGCAGAGGAGGATGGTAAGGGGTGCTTCGAGATATATCACGCATGTGAT GATTCATGCATCGAGTCCATTCGCAACAATACATACGATCACAGCCAGTATCGGGAGGAAGCTCTGCTAAACCGATTGAACATTAATCCGGT AACTTTGTCCTCCGGTTACAAGGACATTATCCTCTGGTTCTCGTTCGGGGCAAGTTGCTTCGTTCTTTTGGCAGTCGTTATGGGTCTGGTTTT TTTTTGCCTGAAGAACGGCAATATGCGCTGCACAATCTGCATC
2927	ATGTATAAGATTATAGTAATCATAGCTCTTCTAGGCGCTGTCAAAGGACTAGACAAGATCTGTTTAGGCCATCATGCTGTTGCAAATGGAACA ATCGTGAAAACATTAACAAACGAACAGGAAGAGGTAACAAATGCTACTGAGACAGTCGAAAGCACTGGCATCAACAGGCTGTGCATGAAGG GGCGTAAACACAAGGACCTTGGCAACTGCCATCCCATCGGAATGCTCATCGGCACCCCCGCATGCGATCTGCACCTAACCGGAACCTGGG ACACTCTCATCGAGCGCGAAAACGCAATCGCCTACTGCTACCCCGGCGCTACGGTGAACGTCGAGGCGTTGCGACAGAAGATTATGGAAT CAGGTGGAATCGATAAGATCTCCACTGGATTTACATACGGATCGTCTATCAACAGCGCCGGTACAACTCGCGCCTGTATGCGTAACGGCGG AAACTCATTTTACGCCGAATTAAAGTGGCTGGTGAGCAAAAGTAAAGGGCAGAACTTTCCGCAAACAACAAATACCTATAGAAATACAGACA CTGCCGAGCACCTCATTATGTGGGGTATACACCACCCTAGCAGCACGCAGGAGAAAAACGACCTCTACGGAACCCAGTCACTGAGCATCTC AGTTGGTTCCTCCACATACCGGAACAATTTCGTGCCCGTCGTGGGCGCACGCCCTCAGGTCAATGGACAGTCTGGTCGTATAGACTTTCAT TGGACGCTTGTGCAGCCAGGCGATAACATCACATTTAGCCATAACGGGGGGTTGATTGCTCCCTCACGGGTGAGCAAGCTAATAGGTAGG GGCCTCGGTATCCAGTCCGATGCACCTATTGATAACAACTGCGAGTCTAAATGTTTTTGGCGAGGTGGGTCCATCAATACGCGGCTGCCTTT TCAGAATCTTTCTCCTCGAACTGTCGGGCAATGCCCAAAATACGTCAACCGTCGTAGTCTTATGTTGGCTACAGGGATGCGGAACGTACCG GAACTTATTCAGGGCAGAGGCCTGTTTGGTGCGATAGCCGGCTTCCTGGAAAACGGATGGGAGGGAATGGTCGATGGATGGTACGGTTTT CGACATCAAAATGCCCAGGGAACCGGGCAGGCAGCGGACTATAAGTCTACCCAGGCAGCCATAGACCAGATTACAGGAAAGCTCAATAGA CTCGTGGAGAAAACTAACACAGAGTTTGAATCAATCGAGAGCGAGTTCAGCGAGATTGAACATCAAATCGGGAACGTGATCAACTGGACCA AAGATTCTATCACAGACATTTGGACATACCAAGCTGAACTCCTAGTTGCCATGGAAAATCAACACACTATCGATATGGCCGATTCAGAAATGT TGAATCTGTATGAGCGCGTGAGAAAACAACTCCGCCAAAACGCAGAAGAGGATGCCAAGGGCTGTTTTGAGATCTACCACGCCTGTGATGA CTCTTGTATGGAGTCCATTCGGAATAATACCTACGATCATAGCCAGTATCGGGAAGAGGCTCTGCTTAACCGATTGAACATTAATCCCGTGA CCTTGTCAAGCGGTTACAAAGACATAATTCTATGGTTCTCATTTGGCGCCTCCTGCTTTGTGCTGCTCGCTGTTGTCATGGGCCTCTTTTTTT TTTGTCTGAAAAATGGAAATATGCGATGTACCATCTGCATT

2928	ATGTATAAAATTGTCGTGATCATAGCTCTCCTTGGGGCTGTGAAAGGCCTGGATAAGATCTGTCTGGGACACCACGCGGTGGCAAATGGCA CTATCGTCAAAACCCTGACAAACGAGCAGGAGGAGGTTACTAACGCCACGGAAACGGTTGAGTCCACCGGGATTAACAGACTTTGCATGAA GGGCAGGAAACACAAGGATCTCGGTAACTGTCACCCCATTGGCATGCTGATTGGGACTCCCGCATGTGACCTTCACTTAACCGGAATGTGG GACACTCTGATCGAGCGTGAAAACGCCATTGCCTACTGTTATCCCGGGGCAACAGTGAACGTCGAGGCGTTGAGGCAAAAGATCATGGAAA GCGGCGGCATCAACAAAATCAGCACTGGGTTCACATACGGATCTAGCATTAACAGCGCCGGCACTACACGTGCATGTATCAGGAATGGCG GCAATTCTTTTTACGCCGAACTGAAGTGGCTTGTTTCCAAATCCAAGGGTCAGAACTTTCCACAAACCACGAATACCTATCGGAATACTGACA CTGCAGAGCATCTAATTATGTGGGGCATCCACCACCCCAGTTCAACTCAGGAGAAGAATGACTTGTACGGGACCCAGAGCCTGAGTATTTC TGTTGGGAGTTCAACCTACAGAAATAACTTCGTCCCTGTCGTCGGGGCTAGACCTCAAGTTAATGGTCAGTCCGGCCGTATCGACTTTCACT GGACTCTGGTACAGCCGGGGGATAACATTACATTCTCACACAATGGGGGTCTCATCGCACCCAGCCGGGTGTCCAAACTCATTGGCCGAG GATTGGGGATCCAATCCGATGCCCCCATAGACAACAATTGCGAGTCTAAATGCTTTTGGAGGGGCGGCTCTATAAACACACGCCTGCCATT CCAGAACCTCAGTCCCCGGACCGTCGGGCAGTGCCCAAAATATGTGAATCGAAGGAGTCTTATGCTGGCTACTGGCATGCGTAACGTGCCT GAGCTGATACAGGGAAGGGGCCTTTTCGGGGCTATCGCCGGATTTTTAGAGAATGGATGGGAGGGGATGGTTGACGGGTGGTATGGATTT CGGCACCAGAATGCACAGGGTACAGGACAAGCCGCCGATTACAAGTCTACCCAAGCTGCAATAGATCAGATCACCGGCAAACTCAACCGG CTTGTTGAAAAGACCAACACAGAATTCGAATCAATCGAGAGCGAGTTCTCTGAAATAGAGCATCAGATTGGCAACGTGATCAACTGGACAAA AGACTCTATTACAGACATCTGGACTTACCAGGCTGAGTTGCTCGTGGCCATGGAGAACCAACATACAATCGATATGGCAGACTCAGAGATG CTTAACCTGTATGAACGTGTGAGGAAGCAGCTCCGCCAGAATGCTGAAGAGGACGGAAAGGGGTGTTTTGAGATATACCATGCCTGCGACG ATAGCTGCATGGAGTCCATCAGGAACAATACGTACGATCACTCCCAATACAGAGAGGAGGCCTTGCTGAATCGGCTTAATATCAACCCCGT CACGCTGAGTTCCGGATACAAGGATATTATACTATGGTTCTCATTTGGCGCTTCATGCTTCGTGCTCCTCGCCGTGGTGATGGGCCTGTTTT TCTTTTGCCTGAAGAACGGTAACATGAGGTGTACAATCTGCATT
2929	ATGTACAAAATCGTGGTCATTATTGCCCTGCTGGGCGCCGTAAAAGGCCTCGATAAGATTTGTCTGGGACATCACGCGGTGGCTAACGGAA CAATTGTTAAAACCCTGACTAACGAGCAGGAAGAAGTCACAAACGCTACGGAGACCGTGGAGAGTACCGGAATCAATCGCCTGTGCATGAA GGGCCGTAAGCACAAGGACCTGGGTAACTGCCATCCAATCGGGATGCTTATCGGCACGCCTGCTTGCGATTTGCATCTGACTGGGATGTG GGACACTCTGATCGAGCGTGAGAACGCTATAGCTTACTGCTACCCCGGAGCCACCGTCAATGTTGAAGCACTGCGTCAGAAGATCATGGAA AGCGGTGGCATCAACAAGATCTCTACCGGATTCACTTATGGCTCTTCCATAAACTCAGCAGGCACCACACGGGCATGCATGCGGAATGGCG GCAACTCTTTCTACGCCGAACTAAAATGGCTGGTGAGTAAGAGCAAGGGGCAAAACTTTCCACAGACCACTAATACATACAGGAATACTGAC ACCGCAGAGCACCTCATCATGTGGGGAATCCACCATCCTTCAAGTACCCAAGAGAAAAACGATCTCTACGGGACACAGTCACTTAGTATATC CGTTGGCTCAAGCACTTACAGAAATAATTTTGTGCCCGTGGTCGGCGCCCGCCCCCAGGTTAATGGACAGAGCGGACGGATTGACTTTCAT TGGACCCTGGTCCAGCCCGGAGACAACATAACTTTTAGCCACAATGGCGGGTTGATCGCACCCAGCAGAGTTTCCAAGCTGATAGGAAGAG GGCTGGGTATCCAATCCGATGCCCCCATCGATAACAATTGTGAGTCAAAATGCTTCTGGCGCGGAGGGTCAATCAACACACGCCTTCCCTT TCAAAACCTCTCACCTCGGACCGTGGGTCAGTGTCCGAAATATGTTAATCGACGGAGCCTGATGCTGGCCACCGGGATGCGCAACGTGCC CGAGTTGATCCAGGGTAGGGGCTTATTTGGTGCAATCGCCGGGTTCCTGGAGAACGGATGGGAAGGCATGGTAGACGGCTGGTATGGATT CCGCCATCAGAACGCCCAGGGGACAGGCCAGGCAGCCGATTATAAAAGCACCCAGGCTGCAATCGACCAGATCACGGGAAAGCTGAACC GGCTGGTGGAGAAGACTAATACGGAATTCGAATCTATTGAGTCTGAATTCAGTGAGATTGAGCATCAGATCGGCAACGTGATTAATTGGACT AAGGATAGCATCACCGACATTTGGACCTATCAGGCCGAGCTGCTGGTGGCTATGGAAAATCAGCATACAATCGACATGGCTGACTCTGAGA TGCTGAACCTCTACGAGAGAGTCAGAAAACAGTTACGTCAGAACGCCGAGGAGGATGGCAAGGGGTGCTTTGAGATCTATCACGCGTGCG ACCATAGTTGCATGGAGTCAATTAGGAACAATACCTATGACCATAGTCAATACCCCGACGAGGCCCTGTTGAACCGCCTGAACATCAACCC TGTGACACTGTCCTCGGGGTATAAGGACATCATCCTTTGGTTCTCATTTGGAGCTAGCTGCTTCGTGTTATTGGCCGTTGTGATGGGCCTTT TCTTCTTCTGTCTCAAAAATGGAAACATGCGCTGTACCATTTGTATT
2930	ATGTACAAAATCGTGCTCGTTCTGGCTCTTCTTGGGGCTGTGCACGGGCTGGATAAAATTTGCCTGGGCCACCATGCCGTCCCAAATGGCA CTATTGTTAAGACTTTAACCAACGAGAAAGAGGAAGTGACTAATGCTACCGAGACAGTGGAAAGCAAAAGCCTGGACAAGCTTTGTATGAAA AATAGGAATTACAAGGATCTGGGCAATTGTCATCCAATTGGCATGCTCGTCGGCACTCCCGCCTGCGATCTGCACCTTACAGGCACGTGGG ATACCCTCATCGAGCGCGATAACAGCATTGCATATTGTTACCCAGGGGCCACGGTTAGTGAGGAAGCATTGAGACAGAAGATCATGGAGAG TGGTGGCATCGATAAGATTTCAACAGGGTTCACCTATGGATCTTCTATCAATAGTGCAGGTACCACTAAGGCTTGCATGCGCAACGGCGGC AACTCATTTTACTCTGAGTTGAAGTGGCTCGTCAGTAAAAATAAGGGCCAGAACTTTCCGCAGACAACAAATACTTACCGAAACACGGATAG TGTGGAACACCTCATAATTTGGGGAATCCACCACCCCTCCAGCACACAGGAGAAGAACGACTTGTATGGCACTCAAAGCCTATCAATTAGC GTGGGATCATCTACTTACCAAAACAACTTTGTCCCAGTCGTGGGTGCCAGGCCACAAGTGAATGGCCAGAGTGGTCGGATTGATTTCCACT GGACGATGGTCCAACCCGGCGATAACATTACATTTTCTCATAATGGAGGCCTCATTGCTCCTAACAGGGTGTCTAAGCTGAAGGGCAGAGG ACTCGGAATCCAATCCGGCGCCTCAGTGGATAATGATTGTGAGAGCAAGTGTTTTTGGAAGGGGGGTTCTATCAATACTAAGCTCCCCTTCC AGAACCTGAGCCCACGCACCGTAGGACAATGCCCCAAATACGTGAACAAGAAGTCCCTGCTCCTCGCAACAGGCATGCGGAACGTTCCAG AAGTGGCACAGGGCAGGGGGTTATTTGGGGCTATCGCTGGTTTTATTGAGAACGGATGGGAGGGGATGGTGGATGGCTGGTACGGTTTCC GCCATCAGAACGCCCAGGGGACTGGCCAAGCGGCAGACTACAAGTCCACACAGGCAGCCATAGATCAGATAACTGGAAAGCTTAATAGGC TCATTGAGAAGACAAATACCGAGTTTGAAAGCATCGAATCCGAATTTTCCGAGATTGAGCATCAAATTGGCAACGTAATCAACTGGACAAAG GACAGCATTACTGACATCTGGACGTACCAGGCCGAACTGCTGGTGGCGATGGAAAACCAGCACACTATCGATATGGCAGATAGTGAAATGC TGAACTTATATGAGCGCGTGCGAAAGCAACTGAGGCAAAACGCTGAAGAGGACGGCAAAGCGTGTTTCGAAATTTACCATAAGTGCGATGA TAATTGTATGGAGAGTATCCGGAATAACACGTATGATCATACCCAGTATCGCGAAGAAGCTCTCCTTAACAGGCTTAATATTAATCCAGTCAA GCTAAGTTCCGGATATAAGGACGTCATTCTCTGGTTTAGTTTTGGCGCTTCCTGCTTTGTATTGCTGGCTGTGATCATGGGCCTGGTCTTCTT CTGTCTCAAGAACGGGAACATGCGATGCACCATATGCATC

2931	ATGTATAAAATTGTGGTCATCATCGCTTTATTGGGAGCAGTTAAGGGGCTCGATAAGATCTGTCTGGGCCACCACGCCGTTGCAAACGGGA CCATTGTGAAGACTCTCACCAATGAGCAGGAGAAAGTTACTAATGCAACAGAGACCGTCGAATCTACAGGCCTGAATCGGCTTTGTATGAAG GGACGCAAGCATAAGGACCTGGGCAATTGCCACCCTATAGGCATGCTAATTGGGACGCCGGCCTGCGACTTGCACCTCACAGGTACGTGG GACACTATCATCGAAAGGGAGAACGCTATCGCCTACTGTTACCCAGGCGCTACCGTGAACGAGGAAGCGCTGAGACAGAAAATCATGGAAT CCGGTGGAATCGATAAGATATCTACTGGCTTTACTTACGGCTCATCCATTAATTCCGCTGGCACCACTCGCGCCTGTATGCGCAACGGCGG CAATTCCTTCTACGCTGAACTTAAATGGCTCGTGTCCAAGAGCAAAGGCCAGAACTTTCCACAGACCACCAATACTTATAGAAATACAGACA CGGCCGAGCACCTGATCATGTGGGGGATCCATCACCCTAGTAGTACACAGGAGAAAAATGACTTATACGGCACCCAATCCCTTAGTATATCT GTTGGGTCAAGCACGTACAGAAACAATTTCGTGCCTGTAGTGGGCGCCAGGCCACAGGTCAACGGGCAGTCAGGCAGGATTGATTTCCATT GGACACTGGTGCAACCTGGTGATAATATCACTTTCTCCCACAACGGCGGTTTAATTGCTCCCAGCAGAGTAAGTAAGCTCATCGGCCGAGG ACTCGGCATTCAGAGCGACGCACCCATAGATAATAACTGCGAATCTAAATGTTTTTGGAGGGGGGGCTCTATCAATACACGCCTCCCTTTTC AGAACCTGTCTCCTAGGACTGTAGGGCAGTGCCCCAAGTACGTAAACAAAAGATCGCTGATGTTAGCCACCGGGATGCGGAACGTCCCTGA GCTCATTCAGGGAAGGGGCCTGTTCGGAGCCATTGCAGGGTTCTTGGAGAATGGATGGGAAGGAATGGTTGACGGGTGGTACGGGTTCC GACATCAGAATGCCCAAGGCACCGGACAGGCCGCTGACTACAAAAGCACCCAGGCAGCCATAGACCAGATTACCGGCAAACTAAATCGTTT AGTGGAGAAGACCAATACCGAATTTGAATCCATAGAGAGCGAATTCAGCGAGATAGAGCATCAAATCGGCAATGTGATTAACTGGACAAAAG ATTCCATAACCGACATTTGGACTTATCAGGCCGAGTTGCTAGTCGCAATGGAGAATCAGCATACAATCGACATGGCTGACTCTGAGATGTTG AATCTGTACGAGAGGGTTCGGAAGCAGCTGCGGCAGAACGCCGAGGAAGACGGGAAAGGGTGTTTTGAGATCTACCATGCGTGTGATGAT AGCTGTATGGAAAGCATTCGCAATAACACCTATGACCACTCACAGTATCGGGAAGAGGCGTTGTTAAACAGGCTGAACATCAACCCAGTAAC ACTGTCTAGCGGTTATAAGGACATAATATTGTGGTTCTCCTTTGGTGCAAGCTGCTTTGTCTTGCTGGCGGTCGTGATGCGCCTGGTGTTCT TTTGTCTGAAAAACGGTAACATGAGGTGTACTATTTGTATC
2932	ATGTATAAGATCGTTCTCGTGTTGGCTCTACTGGGTGCCGTCCATGGTTTGGACAAGATCTGCCTGGGCCACCATGCAGTGCCAAACGGTA CCATAGTGAAAACCTTGACTAATGAAAAAGAAGAGGTGACTAATGCCACCGAGACCGTGGAGTCGAAATCCCTCGATAAACTGTGTATGAAG AACAGGAACTACAAGGACCTGGGCAATTGCCATCCAATTGGCATGGTGGTGGGGACACCTGCCTGCGACTTACACCTCACCGGGACATGG GACACTTTGATCGAGAGGGATAATTCAATTGCGTACTGCTATCCCGGGGCCACTGTGAGTGAGGAGGCGCTGAGGCAAAAGATCATGGAGT CTGGGGGCATAGACAAGATTAGCACCGGCTTCACATATGGAAGCTCCATCAACTCAGCTGGCACGACGAAAGCTTGCATGCGGAACGGTG GGAACTCCTTTTATTCTGAGCTGAAGTGGTTAGTTAGCAAGAACAAAGGCCAGAACTTCCCCCAGACTACAAACACATATAGAAACACAGAC TCAGTGGAGCATTTGATCATTTGGGGGATTCACCATCCATCGTCCACACAGGAAAAGAACGACCTGTATGGCACCCAATCGCTGAGCATCT CAGTCGGGTCCAGTACTTATCAGAATAACTTCGTGCCCGTGGTGGGCGCTCGCCCACAGGTAAACGGGCAAAGCGGGAGAATCGACTTCC ATTGGACTATGGTGCAGCCTGGGGATAATATTACCTTCTCCCACAATGGAGGTCTTATAGCCCCCAATCGAGTTAGCAAACTGAAAGGCCGA GGACTGGGTATCCAGAGCGGCGCAAGCGTGGATAATGATTGTGAGTCCAAATGTTTTTGGAAGGGAGGATCAATTAATACCAAATTGCCCT TCCAGAACTTGAGCCCTCGCACTGTTGGCCAGTGTCCAAAGTACGTTAACAAGAAATCATTACTGCTAGCCACCGGCATGCGTAACGTGCC TGAGGTCGCCCAGGGGCGCGGTCTCTTCGGAGCAATAGCAGGATTCATAGAGAATGGTTGGGAGGGAATGGTGGACGGCTGGTATGGATT CCGCCATCAGAATGCTCAAGGAACCGGCCAGGCTGCGGACTATAAAAGCACGCAAGCTGCGATAGACCAAATTACGGGAAAGCTGAACCG CCTGATTGAAAAGACCAATACTGAGTTTGAATCTATTGAGTCTGAGTTCTCCGAAATCGAACATCAGATTGGGAATGTTATCAACTGGACTAA AGACAGTATAACTGACATTTGGACGTACCAGGCGGAGCTTCTAGTGGCCATGGAAAATCAGCACACCATAGATATGGCAGATTCTGAAATGC TGAACCTCTACGAACGAGTGCGAAAGCAATTGCGTCAGAACGCCGAGGAGGACGGCAAGGGATGTTTCGAGATTTACCACAAATGTGACGA CAACTGCATGGAAAGCATTAGGAATAACACTTACGACCATACACACTACAGGGAAGAGCCCCTGCTCAACCGCCTCAATATTAACCCGGTG AAACTATCTTCTGGGTACAAGGATGTTATTCTGTGGTTTAGCTTCGGCGCCTCCTGTTTTGTACTGCTTGCCGTGATTATGGGTTTGGTTTTC TTCTGCCTCAAGAATGGCAATATGAGATGTACAATTTGCATA
2933	ATGTACAAGATCGTGGTCATCATTGCGCTCCTGGGTGCCGTGAAAGGGCTTGACAAGATCTGCCTGGGTCACCATGCTGTCGCCAATGGGA CAATAGTGAAGACTCTTACCAACGAGAAGGAAGAAGTCACCAACGCAACAGAGACTGTCGAGAGTACCGGCTTAAACAGACTTTGTATGAAA GGGCGCAAGCCACAAGGACCTGGGGAACTGTCACCCAATTGGCATGCTGATAGGGTCTCCAGCCTGTGATCTGCACCTAACTGGCACATGG GACACCCTGATTGAGAGAGAAAACGCCATTGCATACTGTTATCCAGGTGCCACCGTTAATGGGGAGGCATTGAGGCAGAAAATCATGGAGT CCGGAGGAATAGACAAGATATCAACAGGGTTCACATATGAGAGCTCTATCAACTCGGCTGGGACAACCAGGGCATGTATGCGCAACGGAG GAAATTCGTTCTATGCCGAGCTGAAATGGCTTGTTAGCAAGTCTAAGGGCCAGAATTTTCCTCAGACTACCAATACATACCGCAACACCGAC ACTGCCGAGCATCTGATTATGTGGGGGATTCACCACCCCTCTAGTACTCAGGAGAAGAACGACTTGTACGGGACACAAAGCCTGTCAATTA GTGTCGGGAGCTCTACATACCGGAATAATTTCGTTCCTGTTGTCGGGGCGAGGCCACAGGTGAACGGCCAGAGTGGGAGGATCGACTTCC ACTGGACTCTCGTCCAACCAGGGGACAACATCACTTTCAGCCACAACGGAGGCCTTATTGCTCCTTCGAGAGTCAGTAAGCTGATTGGCAG GGGTTTGGGCATTCAAAGTGATGCCCCAATTGATAATAACTGCGAAAGCAAGTGCTTCTGGCGAGGGGGCTCTATTAATACTCGATTGCCTT TTCAGAACCTGAGCCCGAGAACAGTTGGGCAATGTCCCAAGTACGTTAATAAGAGGTCTCTGATGCTGGCGACAGGGATGAGAAATGTCCC TGAGCTGATGCAAGGGCGGGGACTGTTTGGCGCTATAGCGGGCTTCCTCGAAAACGGATGGGAGGGAATGGTTGATGGATGGTATGGGTT CCGACACCAGAATGCACAGGGCACCGGGCAAGCTGCTGATTACAAGTCTACTCAAGCCGCCATCGACCAGATCACCGGCAAACTTAACCG CCTGGTCGAGAAAACAAATACAGAATTTGAATCTATTGAATCCGAGTTCTCTGAGATAGAACACCAAATCGGCAACGTGATTAATTGGACCAA GGATAGTATCACAGATATCTGGACCTATCAAGCGGAGCTGCTGGTGGCCATGGAGAACCAGCACACAATCGATATGGCTGACTCAGAGATG CTTAATTTGTATGAGCGGGTTCGCAAGCAGCTCCGACAAAATGCTGAGGAGGATGGAAAGGGATGTTTTGAGATCTATCATGCCTGCGACG ACTCATGCATGGAAAGCATTCGCAATAATACCTATGATCATTCACAGTACAGAGAAGAGGCTCTCCTGAACAGGCTGAACATAAACCCCGTT ACACTTTCTAGCGGGTACAAAGACATTATCTTATGGTTCAGTTTCGGAGCATCTTGCTTCGTACTACTGGCCGTTGTCATGGGCCTGGTGTT CTTCTGTCTAAAGAATGGGAACATGCGCTGCACCATTTGCATT

2934	ATGTACAAGATCGTCGTGATTATCGCCTTATTGGGAGCAGTCAAGGGGCTCGATAAGATCTGTTTAGGGCATCATGCGGTTGCAAACGGCA CTATCGTTAAGACCTTGACAAATGAACAGGAGGAAGTGACGAATGCCACTGAAACCGTTGAGTCCACTGGGATTAATCGGCTGTGCATGAA AGGACGAAAGCATAAGGACCTTGGAAACTGCCACCCCATTGGCATGCTGATTGGGACCCCTGCTTGCGACCTGCATCTGACTGGAACCTG GGATACCCTTATTGAGCGAGAAAATGCCATCGCCTATTGCTATCCAGGCGCCACAGCAAACGTCGAGGCCCTCAGACAGAAAATCATGGAG TCCGGTGGAATTGACAAAATCTCTACCGGCTTTACCTACGGGTCTAGCATCAATTCCGCAGGGACTACGAGGGCATGTATGAGAAATGGCG GCAACAGCTTCTATGCCGAGCTTAAATGGCTTGTTTCAAAATCCAAGGGACAGAACTTCCCTCAGACTACAAACACTTATCGGAATACAGATA CAGCAGAACACTTGATTATGTGGGGCATTCACCACCCAAGCAGTATCCAGGAGAAAAATGACCTCTACGGCACGCAGTCACTGAGTATATC CGTGGGCTCTTCAACCTACAGGAACAATTTTGTCCCCGTCGTCGGTGCGAGGCCTCAAGTGAACGGGCAATCGGGAAGGATCGACTTCCAT TGGACACTGGTGCAACCTGGAGATAATATCACGTTTAGCCACAACGGAGGCTTGATTGCTCCATCCAGAGTGAGCAAACTGATCGGCAGGG GCCTAGGGATCCAATCAGACGCACCAATCGATAACAACTGTGAGTCTAAGTGTTTCTGGAGAGGTGGCAGCATTAACACCCGGTTGCCCTT CCAGAATTTGAGCCCCCGGACTGTCGGTCAGTGCCCAAAGTACGTGAATCGGCGCTCTCTTATGCTGGCCACCGGAATGAGGAACGTCCC CGAGCTCATACAGGGCCGGGGTTTATTCGGGGCAATTGCTGGGTTCCTGGAGAACGGCTGGGAAGGCATGGTGGATGGCTGGTACGGATT TAGGCACCAGAACGCACAAGGCACTGGCCAAGCCGCGGACTACAAATCAACTCAAGCTGCCATCGATCAGATAACCGGGAAGTTGAACAG ACTGGTAGAAAAAACCAATACCGAGTTCGAGTCAATTGAGTCCGAGTTTAGTGAGATAGAACACCAGATCGGTAACGTGATCAACTGGACAA TGGATAGTATTACTGACATTTGGACCTATCAAGCAGAACTGCTGGTGGCGATGGAGAACCAGCACACTATCGACATGGCAGACAGCGAAAT GCTTAACCTGTACGAACGGGTGCGGAAACAGCTAAGACAAAACGCCGAAGAGGACGGCAAAGGTTGCTTCGAAATCTACCACGCCTGTGA CGACTCTTGTATGGAAAGCATAAGGAACAATACCTACGATCACTCCCAGTACCGAGAGGAGGCGTTACTCAATAGACTGAACATCAACCCTG TGACCCTGAGCTCTGGATATAAAGACATCATTCTCTGGTTTTCTTTTGGGGCCAGCTGCTTCGTGCTTCTCGCAGTCGTTATGGGACTCGTG TTCTTCTGCTTGAAGAACGGTAATATGAGATGCACAATCTGCATA
2935	ATGTATAAGATTGTCGTGATTATTGCCCTGTTAGGGGCCGTGAAAGGACTAGATAAAATTTGCCTGGGCCACCATGCTGTCGCCAATGGCAC GATCGTTAAAACTTTAACAAACGAGCAGGAGGAAGTAACGAACGCAACAGAGACAGTTGAGAGTACAGGAATCAATAGACTGTGTATGAAG GGGAGGAAACACAAGGATTTAGGAAATTGCCACCCTATCGGAATGCTCATCGGAACTCCAGCATGTGATCTGCATCTGACAGGCATGTGGG ACACACTGATCGAGCGGGAAAATGCGATTGCCTATTGCTATCCCGGGGCTACAGTGAATGTCGAGGCTCTGCGGCAGAAGATCATGGAATC CGGCGGCATCAATAAAATTTCTACTGGGTTCACTTACGGGAGTTCTATTAATAGTGCCGGCACGACCAGAGCCTGCATGCGGAACGGCGGC AACTCGTTCTATGCAGAGCTGAAATGGCTGGTTTCAAAATCTAAAGGACAGAACTTTCCACAAACAACAAATACTTATAGGAACACCGACACC GCCGAGCATCTGATTATGTGGGGAATCCATCACCCAAGTAGCACACAGGAAAAGAACGACCTTTACGGCACTCAGTCACTGAGCATAAGCG TTGGGTCCTCTACTTACCGCAATAATTTCGTGCCAGTCGTCGGCGCTAGACCACAAGTCAATGGGCAATCAGGGAGAATCGATTTTCACTGG ACCCTTGTGCAGCCTGGTGATAATATCACTTTCAGCCACAACGGGGGCCTGATTGCTCCAAGTAGAGTCTCCAAGCTTATTGGACGGGGAC TCGGGATTCAGTCTGATGCTCCCATAGATAATAATTGTGAGTCGAAGTGCTTTTGGCGAGGAGGGTCCATCAACACGCGGTTGCCCTTCCA GAACCTCTCACCTCGGACCGTGGGGCAGTGCCCCAAATACGTGAACAGGAGGAGTCTGATGCTGGCCACCGGGATGCGCAACGTGCCCG AATTGATCCAGGGGAGAGGCCTTTTCGGCGCTATAGCGGGGTTTTTAGAAACGGGTGGGAGGGCATGGTGGACGGCTGGTACGGATTTC GCCATCAGAATGCCCAGGGTACTGGTCAAGCCGCCGATTACAAATCTACCCAGGCTGCCATCGACCAGATTACAGGAAAGTTGAACCGACT GGTGGAAAAGACCAACACTGAGTTTGAGAGTATTGAAAGTGAATTCTCTGAGATCGAACACCAGATTGGGAACGTTATCAACTGGACGAAAG ATTCCATCACAGACATCTGGACCTACCAAGCCGAACTCCTTGTGGCCATGGAGAATCAACACACGATCGACATGGCTGACTCAGAAATGCT GAATTTGTACGAAAGAGTGAGAAAACAGCTGAGGCAAAATGCCGAGGAGGATGGCAAAGGCTGTTTCGAGATTTATCACGCCTGCGATGAT TCCTGTATGGAATCAATCAGGAATAATACCTACGACCACTCTCAGTACCGCGAGGAAGCCCTATTGAACCGACTTAACATAAACCCTGTCAC TTTGTCCAGTGGTTACAAGGATATCATTCTTTGGTTTTCGTTCGGAGCTTCCTGCTTCGTACTCCTCGCGGTGGTAATGGGCTTATTCTTTTT TTGCCTAAAGAACGGAAATATGAGGTGTACGATCTGCATC
2936	ATGTACAAAATTGTGGTGATTATTGCTCTGCTCGGAGCCGTCAAGGGCCTGGATAAGATCTGTCTGGGCCACCACGCCGTGGCCAACGGAA CAATTGTGAAGACACTCACAAACGAGCAGGAAGAAGTGACTAATGCCACCGAGACCGTTGAAAGTACCGGGATAAACCGCCTTTGCATGAA GGGTCCGAAACATAAACACCTGGGGAACTGCCACCCTATAGGTATGCTTATTGGAACACCCGCTTGTGATCTACACCTTACTGGGATGTGG GACACCCTTATAGAGCGCGAGAACGCCATCGCATACTGTTATCCTGGTGCCACGGTTAACGTGGAGGCTCTTAGGCAGAAGATCATGGAGT CAGGCGGTATTAATAAGATTTCCACAGGCTTTACTTACGGGAGCTCAATAAACAGCGCCGGGACCACGAGGGCCTGCATGCGGAACGGCG GGAACTCCTTTTACGCGGAATTGAAATGGCTGGTGTCAAAGTCCAAAGGACAAAACTTCCCTCAAACCACTAATACCTACAGGAATACCGAC ACTGCAGAACATCTTATCATGTGGGGCATACACCACCCGTCGTCTACACAAGAGAAGAACGATCTTTATGGGACCCAGTCTCTGTCCATCAG CGTGGGGTCATCCACTTACCGAAATAATTTTGTGCCTGTGGTTGGGGCTAGGCCACAGGTGAATGGGCAGTCTGGCCGTATTGATTTCCAC TGGACACTTGTGCAGCCAGGGGATAATATAACATTCAGCCACAATGGGGGCCTAATCGCTCCATCCCGGGTGTCTAAGTTGATCGGTCGCG GACTTGGAATCCAATCTGACGCTCCAATCGACAACAATTGTGAGTCTAAATGCTTTTGGCGCGGCGGCTCTATCAACACTAGACTGCCATTC CAAAATCTGTCTCCTCGCACCGTTGGGCAGTGTCCTAAGTATGTAAACAGACGGTCCCTGATGCTTGCGACAGGAATGCGGAATGTACCTG AACTCATCCAGGGAAGAGGGCTGTTCGGAGCTATTGCCGGCTTTCTGGAGAATGGTTGGGAGGGAATGGTAGACGGCTGGTACGGCTTTC GGCATCAAAATGCGCAGGGAACAGGCCAGGCAGCAGACTACAAATCAACCCAGGCTGCCATTGACCAGATTACAGGGAAGCTCAATAGAC TTGTGGAGAAAACAAACACCGAGTTCGAATCCATCGAGTCCGAGTTCAGTGAGATCGAGCACCAGATTGGGAACGTGATAAACTGGACGAA AGATTCTATCACCGATATTTGGACATACCAGGCTGAGCTCCTGGTCGCCATGGAGAATCAACATACAATTGATATGGCGGACAGCGAGATG CTGAACCTATATGAGCGCGTACGAAAGCAATTGAGACAGAACGCCGAGGAGGACGGTAACGGGTGCTTTGAAATATATCATGCTTGTGACG ACTCATGCATGGAATCAATTAGAAATAATACATATGACCACTCTCAATATAGAGAGGAGGCCCTCTTGAATCGCTTGAACATCAATCCGGTGA CTCTCTCTAGTGGGTACAAAGACATTATACTGTGGTTCAGTTTCGGGGCCAGCTGCTTCGTGCTGCTGGCAGTAGTTATGGGGCTCTTCTTT TTTTGCCTGAAAAATGGCAATATGCGATGTACCATTTGTATC

2937	ATGTATAAAATTGTCGTGATTATTGCCTTGCTAGGCGCCGTCAAGGGCCTGGACAAAATCTGCCTGGGACACCACGCTGTGGCGAACGGCA CTATCGTTAAGACTCTGACCAATGAGCAGGAAAAGGTCACAAACGCTACAGAAACGGTCGAATCTACAGGCCTGAACAGGCTGTGTATGAA AGGTAGAAAGCACAAAGACCTGGGAAATTGCCACCCAATAGGGATGTTAATCGGCACCCCAGCATGCGATTTGCACCTGACAGGTACATGG GACACCATCATCGAAAGGGAGAATGCCATTGCTTACTGTTATCCCGGAGCCACAGTGAATGAGGAGGCCCTCAGACAAAAAATAATGGAAT CCGGCGGAATCGACAAGATAAGTACAGGATTTACCTATGGCAGCTCTATTAATTCCGCCGGCACTACAAGAGCTTGTATGAGGAACGGTGG AAATTCGTTTTACGCAGAGCTTAAATGGCTAGTGAGCAAGTCCAAAGGACAGAACTTTCCTCAGACAACAAATACTTATCGGAACACCGATA CGGCTGAACACTTAATAATGTGGGGGATCCACCATCCGTCGTCAACGCAGGAGAAAAATGATCTGTACGGTACGCAATCACTGTCCATCTC CGTAGGCAGCTCCACCTACAGAAATAATTTTGTTCCAGTTGTCGGTGCTCGGCCACAAGTGAATGGCCAGAGCGGCCGAATCGACTTTCAC TGGACCCTGGTGCAGCCAGGGGATAACATCACCTTCAGCCACAACGGTGGCCTTATCGCCCCCTCTAGGGTGAGTAAGCTGATCGGGCGG GGCCTCGGAATCCAATCTGACGCACCCATTGATAACAACTGCGAGAGCAAGTGTTTCTGGCGGGGTGGTTCTATCAATACCCGACTCCCGT TTCAGAATCTATCTCCTAGGACCGTCGGCCAGTGCCCAAAATATGTGAACAAGCGCAGCCTTATGCTAGCAACTGGAATGAGGAATGTTCCC GAGCTGATTCAGGGCCGAGGTCTCTTCGGAGCTATTGCCGGGTTTCTCGAGAACGGATGGGAAGGCATGGTGGACGGGTGGTACGGTTTT CGGCACCAGAACGCCCAGGGAACCGGACAAGCAGCCGATTACAAGAGCACCCAGGCTGCTATCGATCAGATTACCGGCAAGCTCAATCGC TTGGTGGAGAAAACCAATACTGAATTTGAGTCCATTGAATCAGAGTTCAGCGAGATCGAGCATCAGATCGGGAACGTCATTAATTGGACAAA GGATAGCATCACCGATATCTGGACATACCAGGCTGAGCTTCTGGTTGCAATGGAAAATCAGCATACTATCGACATGGCTGATAGCGAAATGT TGAATTTGTATGAAAGGGTGCGTAAGCAGCTGCGGCAGAACGCCGAGGAGGACGGCAAGGGATGCTTCGAGATCTACCACGCATGTGACG ACTCATGCATGGAGAGCATTCGGAACAACACTTATGATCACTCTCAATATCGTGAAGAGGCACTCCTAAACAGACTAAATATAAACCCGGTC ACCCTCAGCAGCGGCTATAAGGACATCATTCTTTGGTTCTCTTTTGGTGCAAGCTGTTTCGTCCTCCTCGCCGTGGTTATGGGCCTGGTGTT TTTTTGCCTGAAGAATGGCAATATGCGGTGCACCATTTGCATC
2938	ATGTATAAGATTGTGGTGATTATTGCGTTGCTGGGGGCTGTGAAAGGCCTGCATAAGATCTGTCTTGGGCATCATGCCGTAGCTAACGGCA CAATTGTGAAAACTCTGACAAATGAGCAGGAGGAGGTGACAAACGCAACCGAGACCGTTGAGAGCACAGGGATTAATCGCCTGTGCATGAA GGGGCGCAAACACAAGGACCTGGGGAACTGTCACCCTATCGGAATGCTGATCGGGACCCCAGCCTGTGATTTACACCTAACAGGAATGTG GGACACACTCATTGAAAGAGAGAACGCGATAGCATACTGCTACCCCGGAGCTACTGTGAATGTGGAAGCACTGCGCCAGAAGATTATGGAA TCCGGAGGCATAAATAAGATTTCCACTGGGTTCACCTATGGCTCCTCTATCAACAGTGCAGGAACGACGAGAGCCTGCATGCGCAATGGGG GAAATAGTTTCTACGCCGAGCTGAAGTGGCTGGTGTCTAAGTCAAAGGGTCAAAACTTTCCCCAAACTACCAATACGTACCGAAATACTGAC ACGGCAGAACACCTAATAATGTGGGGGATTCACCATCCCAGCAGCACCCAGGAGAAGAATGATTTGTATGGGACTCAGTCTCTGTCGATTT CCGTGGGCAGCAGCACGTACCGAAATAATTTCGTACCCGTGGTTGGAGCTCGACCTCAGGTGAACGGCCAGTCCGGCAGAATTGATTTTCA CTGGACACTAGTTCAGCCAGGCGATAATATAACATTTTCGCACAACGGAGGTCTGATCGCACCTTCACGCGTGTCCAAGCTTATCGGTCGG GGCCTGGGGATCCAGAGTGATGCTCCCATCGATAACAATTGCGAGTCTAAATGTTTTTGGCGAGGTGGATCCATCAATACCCGTCTGCCCT TCCAGAACCTGTCTCCACGCACAGTCGGTCAGTGCCCCAAGTATGTGAATAGAAGGTCCCTGATGCTCGCCACAGGAATGAGAAACGTTCC GGAACTAATTCAGGGCCGTGGACTATTCGGGGCCATAGCAGGGTTCCTGGAGAATGGCTGGGAGGGTATGGTTGACGGTTGGTACGGCTT TCGGCATCAGAACGCCCAAGGGACCGGGCAGGCCGCCGACTACAAAAGTACCCAGGCTGCAATCGACCAGATCACGGGGAAACTGAACC GACTCGTGGAGAAGACTAACACTGAGTTTGAGTCCATCGAGTCTGAGTTTAGCGAAATCGAGCATCAGATTGGGAACGTTATTAACTGGACA AAGGACAGCATCACCGATATATGGACCTACCAGGCAGAACTATTAGTGGCAATGGAGAACCAGCACACAATTGACATGGCGGACAGCGAGA TGCTGAATCTTTATGAGCGTGTCCGGAAACAGCTGAGACAGAACGCCGAAGAAGATGGCAAAGGCTGCTTTGAGATATACCATGCGTGCGA TGATTCATGCATGGAGTCTATTCGCAATAATACATACGATCACAGTCAGTACAGAGGAAGCTCTACTGAATCGGCTCAACATCAATCCAG TCACCCTGTCTTCCGGCTACAAGGACATCATTCTGTGGTTCAGCTTTGGAGCCTCCTGTTTTGTACTTCTAGCCGTTGTGATGGGCTTGTTTT TTTTCTGTTTGAAGAACGGGAATATGAGATGTACCATCTGCATC
2939	ATGTACAAGATCGTGGTAATTATCGCTCTGCTAGGAGCCGTGAAGGGGCTAGACAAGATATGTCTCGGGCACCACGCCGTGGTTAACGGAA CCATTGTGAAGACATTGACTAACGAACAGGAGGAGGTGACCAATGCGACGGAAACAGTTGAGTCAACAGGATTGAATAGGCTGTGTATGAA GGGGCGTAACCACAAAGACCTGGGCAATTGCCATCCAATCGGAATGCTTATTGGAACTCCCGCTTGCGACTTACACCTGACCGGCACATGG GATACCCTGATAGAAAGAGAAAACGCTATTGCGTATTGCTACCCAGGAGCTACCGTAAACGAAGAGGCCCTTCGGCAGAAAATAATGGAGT CCGGGGGAATTAACAAAATATCTACTGGCTTTACTTATGGATCGTCCATCAACAGCGCTGGAACTACACGCGCTTGCATGCGTAACGGAGG TAATTCATTTTATGCCGAACTGAAGTGGCTGGTGTCAAAGTCAAAGGGACAGAACTTCCCCCAAACTACAAATACATACAGGAATACCGACA CTGCTGAACACCTGATTATGTGGGGTATTCACCACCCCTCAAGCACCCAGGAAAAGAATGACCTGTACGGAACTCAGAGTTTGTCTATCAGC GTGGGAAGTTCAACCTACCAAAATAACTTCGTTCCGGTGGTTGGCGCCAGGCCCCAGGTGAACGGACAGTCGGGGCGCATCGACTTTCAT TGGACTTTAGTCCAGCCTGGCGACAATATCACATTCTCCCACAACGGAGGGCTTATCGCCCCGAGTCGCGTGAGCAAGCTCATTGGTCGGG GGTTGGGAATTCAGAGCGATGCCCCTATTGACAACAACTGCGAAAGCAAGTGCTTCTGGAGAGGCGGTTCTATTAACACAAGACTCCCATT CCAAAACCTGAGCCCACGGACAGTCGGGCAGTGCCCCAAGTACGTAAATAAAAGGTCACTCATGTTGGCTACCGGGATGCGGAATGTCCC AGAACTGATGCAGGGAAGGGGGCTGTTCGGCGCAATCGCCGGCTTTATTGAGAATGGCTGGGAGGGAATGGTTGACGGATGGTATGGTTT CAGGCACCAGAATGCCCAGGGTACCGGTCAGGCAGCCGATTATAAATCCACCCAGGCAGCGATTGACCAGATTACTGGAAAGTTGAACCG GCTTATTGAAAAAACCAACACTGAGTTCGAGAGCATCGAGAGTGAGTTCAGCGAGATCGAACATCAGATCGGTAACGTGATAAACTGGACC AAGGACTCCATCACTGATATTTGGACCTACCAGGCTGAGTTATTGGTTGCCATGGAGAACCAGCACACCATCGACATGGCCGATTCTGAAAT GCTGAATCTCTATGAGAGAGTTCGGAAACAGCTGAGACAAAATGCAGAGGAGGATGGGAAGGGCTGTTTTGAGATTTACCACGCCTGTGAC GACAGTTGTATGGAGTCTATTAGAAATAATACCTACGATCATTCTCAATATCGAGAAGAGGCCTTGTTAAATAGGCTGAATATTAACCCAGTC ACCCTAAGCAGCGGGTACAAAGATATCATACTGTGGTTCTCGTTTGGAGCCTCTTGCTTTGTGCTATTGGCAGTCGTGATGGGTTTGGTGTT CTTCTGCCTCAAAAACGGGAATATGAGATGTACGATTTGCATC

2940	ATGTATAAGATTGTCCTTGTGCTCGCCCTGCTCGGAGCAGTGCACGGCCTCGATAAAATCTGTCTGGGGCACCACGCCGTGCCTAACGGCA CCATCGTCAAGACTCTCACAAACGAGAAGGAGGAAGTGACAAACGCTACTGAAACAGTAGAAAGTAAGAGCCTGGACAAACTTTGCATGAA AAACAGGAATTATAAAGACTTGGGTAATTGTCACCCGATTGGCATGGTCGTCGGAACGCCTGCTTGTGACCTCCATCTGACGGGAACCTGG GATACCTTGATCGAGCGGGACAACTCAATTGCATACTGCTATCCAGGCGCCACCGTTAGTGAGGAGGCTTTGAGACAGAAGATCATGGAGT CTCGAGGCATTGACAAAATTTCCACCGGCTTTACATATGGTAGCTCAATCAACTCGCCTGGGACTACAAAAGCTTGTATGAGGAACGGTGG AAATAGCTTTTACTCGGAACTAAAATGGTTGGTGTCTAAAAACAAGGGTCAGAACTTTCCTCAGACGACCAACACTTATCGAAATACTGACTC CGTGGAGCACCTGATTATTTGGGGGATCCATCATCCCAGCAGCACACAAGAGAAGAATGACCTGTATGGGACGCAGTCTCTGTCCATTAGC GTCGGCAGTTCAACTTACCAGAATAATTTCGTGCCTGTGGTGGGAGCGCGCCCCCAAGTCAATGGGCAGAGCGGAAGGATTGATTTCCACT GGACTATGGTCCAGCCCGGCGATAACATTACTTTCAGCCACAACGGAGGGCTCATTGCACCAAACCGTGTCTCAAAACTGAAGGGCAGGG GGCTTGGTATTCAGAGCGGAGCTAGTGTGGACAATGACTGCGAATCTAAGTGTTTTTGGAAAGGAGGATCAATTAACACTAAGCTTCCGTTT CAAAACCTGAGTCCACGCACAGTCGGACAGTGCCCTAAATATGTGAACAAAAAGTCGCTTCTGCTAGCTACCGGGATGAGGAACGTGCCAG AGGTCGCCCAGGGCAGAGGTTTGTTCGGCGCGATCGCCGGTTTCATCGAAAACGGATGGGAGGGAATGGTGGATGGCTGGTACGGATTTC GTCATCAAAACGCACAGGGAACGGGTCAGGCCGCAGATTATAAATCTACTCAGGCCGCTATAGATCAGATCACAGGCAAGCTGAATAGGTT AATAGAGAAGACGAATACCGAGTTTGAGAGCATCGAAAGCGAGTTCAGTGAAATAGAGCATCAGATTGGCAATGTGATCAATTGGACAAAAG ATTCGATCACTGATATTTGGACATATCAGGCAGAACTTCTTGTGGCAATGGAGAACCAGCATACGATTGATATGGCCGATTCGGAAATGCTG AATCTTTACGAGAGAGTCAGAAAACAGCTGCGCCAAAATGCTGAGGAAGACGGGAAGGGGTGCTTTGAAATATACCATAAATGTGACGACA ACTGCATGGAGTCGATCCGCAACAACACATATGACCATACCCAGTACCGTGAGGAAGCTCTGTTAAATAGACTTAACATCAATCCAGTTAAA CTGTCCTCTGGGTATAAAGATGTCATTTTGTGGTTCTCTTTCGGCGCTTCATGCTTTGTCCTCCTCGCCGTCATCATGGGACTGGTGTTTTTT TGCCTCAAAAACGGGAATATGCGATGTACGATATGCATC
2941	ATGTATAAAATAGTAGTGATCATCGCTCTGTTAGGAGCGGTGAAAGGGCTCGACAAGATTTGTCTGGGCCACCACGCCGTTGCTAACGGCA CCATCGTCAAGACCCTCACTAATGAACAGGAGGAGGTCACTAACGCCACTGAGACAGTTGAATCGACCGGGATTAACCGGTTGTGTATGAA AGGCCGGAAGCATAAAGATCTCGGGAATTGCCACCCAATAGGGATGCTTATCGGCACACCAGCATGCGACCTCCATCTCACGGGCATGTG GGACACCCTTATCGAAAGAGAAAATGCCATTGCCTATTGCTATCCCGGCGCCAACCGTGAATGTCGAGGCCCTTCGACAAAAAATCATGGAA AGCGGCGGCATTAACAAAATCAGCACGGGATTCACCTACGGGTCTAGTATCAATTCAGCAGGCACGACCAGAGCTTGTATGCGTAACGGGG GTAATTCTTTCTACGCAGAGCTGAAATGGCTCGTGTCTAAGAGCAAAGGTCAGAATTTTCCCCAGACTACTAATACATATAGGAATACCGACA CTGCGGAGCATCTTATAATGTGGGGCATCCACCACCCATCCTCCACACAGGAAAAGAACGACCTGTATGGTACGCAGTCACTGAGTATAAG CGTTGGGTCCAGCACTTATCGAAATAACTTTGTGCCCGTAGTGGGAGCCCGCCCACAGGTTAACGGCCAGAGTGGCAGAATCGACTTCCAT TGGACCCTGGTGCAGCCCGGCGACAACATCACTTTCTCACATAACGGAGGGTTAATAGCTCCATCCAGGGTATCCAAACTCATTGGTAGGG GGCTGGGCATCCAATCCGACGCCCCCATTGACAATAATTGTGAGAGCAAATGTTTCTGGAGGGGTGGAAGCATCAATACCAGACTCCCATT CCAGAATTTATCTCCCCGCACAGTAGGGCAGTGCCCCAAATATGTTAACAGAAGATCCCTGATGCTCGCGACAGGAATGCGGAATGTGCCC GAGCTGATCCAGGGCCGCGGGTTGTTTGGCGCCATAGCTGGTTTTCTCGAGAATGGTTGGGAAGGGATGGTTGACGGCTGGTATGGGTTT CGCCATCAGAACGCTCAGGGAACAGGACAAGCTGCCGACTACAAGAGCACTCAAGCAGCAATCGATCAAATCACGGGGAAGCTCAACAGG CTTGTTGAAAAGACTAATACTGAGTTCGAAAGCATTGAGTCCGAGTTTTCCGAGATTGAGCATCAGATTGGGAACGTCATCAACTGGACCAA AGACAGTATCACGGACATATGGACTTATCAAGCCGAACTCCTGGTGGCCATGGAAAACCAGCACACAATCGATATGGCTGATTCTGAGATG CTTAACCTGTACGAGCGCGTGCGGAAACAACTGCGGCAAAACGCAGAGGAAGACGGAAAAGGATGTTTTGAAATATACCACGCATGTGATG ATAGCTGCATGCAGAGCATTCGTAACAACACTTATGATCATTCTCAGTACCGCGACGAAGCTCTCCTAAACAGACTCAACATTAATCCAGTG ACCCTGAGCTCTGGATATAAAGACATTATTCTGTGGTTTAGTTTCGGCGCCAGCTGCTTTGTGCTGCTGGCGGTGGTTATGGGACTGTTCTT CTTCTGTTTGAAAAACGGAAATATGAGATGTACTATTTGCATC
2942	ATGTATAAGATAGTAGTAATTATTGCTCTCTTGGGGGCGGTGAAAGGGCTGGATAAAATCTGCCTGGGGCATCACGCCGTGGCCAATGGCA CCATTGTAAAAACGCTTACAAATGAGCAGGAAGAAGTGACCAACGCCACAGAAACTGTGGAGAGCACTGGCATCAACCGGCTCTGTATGAA GGGTACAAAGCACAAGGATCTGGGGAATTGCCACCCCATCGGCATGTTAATAGGCACTCCGGCTTGCGATTTGCACCTTACCGGAATGTGG GACACACTAATCGAGAGGGAAAATGCCATTGCGTACTGCTACCCAGGTGCCACCGTGAATGTCGAGGCACTGAGGCAGAAAATCATGGAAT CAGGAGGAATAAACAAGATCTCTACCGGCTTTACGTATGGCTCATCCATCAACTCTGCAGGTACCACTCGAGCTTGCATGCGGAACGGCGG CAATTCCTTCTACGCAGAACTAAAGTGGCTTGTGTCCAAAAGTAAGGGTCAGAATTTCCCCCAAACTACCAATACGTACAGGAACACTGACA CGGCAGAACACCTGATCATGTGGGGCATCCATCATCCCTCCTCTACCCAGGAAAAAAATGACCTGTACGGGACACAGAGCCTGTCCATATC TGTAGGCTCAAGTACATATCGTAACAACTTCGTGCCCGTTGTGGGCGCACGTCCACAGGTTAACGGCCAAAGCGGACGCATTGATTTCCAT TGGACCCTAGTCCAGCCCGGAGACAATATAACCTTCTCGCATAATGGAGGACTGATCGCTCCCTCTCGCGTGTCTAAGCTGATCGGTCGGG GATTGGGGATTCAGAGCGACGCTCCTATTGATAACAACTGCGAATCCAAGTGCTTTTGGAGAGGGGGCAGCATCAATACTAGATTGCCTTTT CAGAACCTTTCTCCCCGCACAGTGGGGCAATGCCCTAAATACGTCAATAGAAGGTCTCTTATGCTTGCCACCGGAATGCGTAATGTACCAGA ACTCATACAAGGTAGAGGGTTATTCGGCGCCATCGCTGGCTTTCTGGAAAATGGCTGGGAGGGGATGGTTGACGGCTGGTATGGTTTCAGA CACCAAAATGCTCAGGGTACAGGTCAGGCCGCCGATTACAAGTCCACACAGGCAGCTATTGATCAGATCACAGGCAAGTTGAACCGCCTCG TGGAAAAGACTAACACAGAGTTTGAGTCTATTGAATCCGAATTCTCTGAGATTGAACACCAAATTGGAAATGTTATCAACTGGACAAAGGACT CTATTACCGATATTTGGACCTATCAAGCCGAGTTGTTGGTCGCCATGGAGAATCAACATACTATTGACATGGCGGACTCTGAAATGCTAAAT CTCTACGAGAGAGTAAGAAAGCAGCTTCGACAGAACGCAGAGGAAGACGGAAAGGGATGTTTTGAGATCTATCATGCTTGTGACGATTCGT GCATGGAATCCATTCGGAATAATACATATGACCATTCTCAGTATAGAGAAGAAGCCCTGCTTAACCGGCTAAACATCAATCCCGTGACGCTA TCATCCGGGTATAAAGATATCATCCTGTGGTTCTCATTCGGGGCTTCGTGTTTTGTGCTCCTGGCGGTCGTGATGGGCCTGTTCTTTTTCTG TCTTAAAAACGGGAACATGCGATGCACCATATGCATC

2943	ATGTACAAGGTCGTAGTTATTATCGCGCTGCTAGGTGCAGTTCGGGGGCTCGATAAGATTTGCCTGGGGCACCATGCGGTCGCCAACGGA ACTATCGTTAAGACATTAACCAACGAACAGGAGGAGGTGACAAATGCAACCGAGACCGTGGAATCTAAATCCCTGGGTAAGCTGTGCATGA AAGGGAGGAGTTATAACGACCTTGGAAACTGTCACCCAATCGGCATTCTTATCGGGACCCCCGCTTGTGATCTCCATTTAACCGGCACCTG GGATACACTCATTGAGCGCGAAAATGCTGTTGCCTACTGTTACCCAGGAGCTACAGTAAACGAGGAAGCTCTTAGACAGAAGATAATGGAG TCCGGAGGGATTTCCAAGATCAGCACTGGCTTCACCTATGGCACATCAATCAACAGCGCAGGGACGACAAAGGCCTGTATGAGGAACGGC GGCAACTCATTCTACGCCGAGCTGAAATGGCTGGTGTCAAAGAACAAAGGCCAGAACTTCCCACAAACCACGAACACCTACAGGAACACCG ATACCGCAGAGCATCTTATAATATGGGGAATCCACCACCCCTCTTCCACCCAGGAGAAAAATGATCTTTATGGCACTCAATCTTTATCTATAT CAGTAGGGTCTTCAACTTACCAGAACAATTTCGTGCCTGTGGTGGGAGCGCGGCCGCAAGTGAACGGTCAGAGCGGTCGTATAGACTTTCA CTGGACCCTGCTGCAGCCAGGTGACAACATCACCTTCTCTCACAATGGTGGGCTAATTGCTCCCTCCAGAGTCAGCAAGCTTATAGGGAGA GGCCTCGGGATACAGTCAGAAGCCCCTATCGACAATGGGTGCGAGTCGAAGTGTTTCTGGAAGGGTGGATCTATTAACACAAAACTTCCTT TTCAGAACTTATCTCCGAGGACTGTGGGCCAATGCCCTAAATACGTTAACAAGCGCAGTTTAATGCTTGCTACGGGTATGAGGAATGTCCCA GAGATCATGCACGGCCGCGGGCTCTTCGGGGCCATTGCCGGATTTATTGAGAACGGGTGGGAGGGCATGGTAGATGGATGGTACGGGTT CCGCCACCAAAACGCACAGGGAACGGGACAGGCAGCTGACTACAAATCAACACAAGCGGCTATCGACCAGATAACAGGCAAGCTGAATAG ACTGATCGAGAAAACTAATACAGAGTTCGAGAGCATAGAGAGCGAATTTTCCGAGATCGAGCATCAGATCGGTAACATTATAAATTGGACTA AGGATAGCATTACTGATATTTGGACCTATCAGGCTGAGCTGCTGGTTGCCATGGAAAACCAACATACAATCGATATGGCCGATTCAGAAATG CTGAATCTATATGAGAGAGTGAGGAAGCAGCTGAGACAGAACGCTGAGGAGGACGGTAAAGGCTGTTTCGAAATTTACCATGCATGCGACG ACTCTTGTATGGAATCTATCCGCAACAACACATACGACCACTCCCAGTACAGGGAGGAAGCCCTGCTGAATCGACTGAACATTAATCCAGTC AAGCTTTCCTCTGGTTATAAGGACATCATCTTATGGTTCAGCTTCGGGGCTTCGTGTCTCATTCTCTTGGCTGTGGTAATGGGCTTAGTTTTC TTTTGTCTTAAGAACGGCAACATGAGGTGCACAATTTGCATC
2944	ATGTATAAAATCGTGGTTATTATTGCACTGCTAGGGGCCGTCAAGGGCTTAGATAAGATTTGCCTGGGCCACCACGCGGTGGCCAATGGGA CCATTGTGAAAACTCTGACCAACGAGCAGGAAGAAGTGACTAACGCCACCGAGACTGTGGAATCAACAGGAATCAATCGTCTGTGCATGAA AGGCAGAAAGCATAAGGACCTTGGAAACTGTCATCCTATTGGGATGCTGATCGGCACCCCAGCTTGTGATTTGCACCTCACCGGTATGTGG GATACTCTTATCGAAAGGGAAAACCCTATCGCTTATTGTTACCCAGGAGCGACTGTCAATGTCGAGGCCCTACGCCAGAAAATAATGGAAAG TGGAGGGATTAATAAGATAAGCACTGGTTTCACGTATGGCAGTTCGATCAATAGTGCCGGAACAACTCGCGCTTGCATGCGTAATGGTGGA AACTCATTTTACGCAGAGCTCAAATGGCTAGTTTCGAAGTCCAAAGGTCAGAACTTTCCACAGACGACCAACACCTATCGCAATACCGACAC TGCAGAGCACCTGATCATGTGGGGAATTCATCACCCGTCAAGCACTCAAGAGAAGAACGATCTTTATGGAACCCAGAGTCTCAGTATTTCCG TTGGCTCGAGTACGTACCGCAATAACTTCGTTCCTGTCGTGGGTGCCAGGCCTCAGGTTAACGGGCAGTCCGGGCGCATCGATTTTCACTG GACTCTCGTTCAGCCCGGGGATAATATCACATTCAGTCACAACGGTGGGCTGATCGCGCCGTCCAGAGTGTCAAAACTGATCGGCAGGGG CCTGGGCATTCAGAGCGACGCACCAATTGATAACAATTGTGAGAGCAAATGTTTCTGGCGTGGAGGCAGCATCAATACCCGACTCCCGTTT CAGAATCTGAGCCCCAGAACTGTCGGACAATGTCCTAAATATGTGAATCGGCGCTCCCTGATGCTCGCAACTGGCATGCGGAACGTGCCCG AGTTGATTCAAGGACGCGGGCTGTTCGGTGCCATCGCTGGGTTCCTGGAAAACGGCTGGGAAGGCATGGTAGATGGATGGTACGGGTTCA GGCACCAGAACGCCCAAGGGACAGGACAGGCCGCCGACTACAAGAGTACTCAAGCTGCTATTGACCAAATTACCGGAAAACTGAATAGGC TGGTGGAAAAGACCAATACAGAGTTCGAGTCTATCGAGAGCGAGTTTAGCGAGATCGAGCATCAGATTGGTAATGTGATTAACTGGACGAA AGATAGTATCACCGACATCTGGACATATCAAGCAGAACTTCTTGTGGCTATGGAGAATCAACATACTATTGACATGGCTGATAGCGAGATGT TGAATCTTTACGAGAGAGTGCGCAAACAGCTGCGGCAGAATGCTGAGGAAGATGGCAAAGGTTGTTTCGAGATCTACCATGCCTGCGACGA CAGTTGTATGGAGAGCATTAGGAATAACACTTACCACCACTCACAATATCGAGAACAGCGCTCTACTCAACAGGCTAAACATCAATCCAGTCA CCCTCTCCTCCGGTTATAAGGATATCATTCTCTGGTTCTCCTTCGGAGCTTCCTGTTTTGTCCTTCTGGCAGTGGTGATGGGATTATTTTTTTT CTGCCTTAAGAATGGGAATATGAGATGTACAATCTGTATT
2945	ATGTACAAGATCGTAGTGATTATCGCATTACTTGGCGCCGTGAAGGGACTCGATAAGATTTGCTTGGGCCATCATGCTGTGGCCAATGGCA CTATAGTGAAGACACTCACCAATGAACAGGAAGAAGTGACCAACGCCACCGAGACCGTGGAGTCTACAGGCATCAACAGGCTCTGTATGAA GGGCCGTAAGCACAAGGATCTTGGCAACTGCCATCCTATCGGAATGCTCATTGGAACACCAGCTTGTGATCTCCACTTAACAGGCATGTGG GACACACTGATTGAGCGGGAGAATGCGATTGCCTACTGTTATCCCGGGGCCACCGTGAACGTTGAGGCTCTCAGACAGAAAATCATGGAGA GCGGCGGGATAAACAAGATAAGCACAGGCTTCACTTACGGTTCTAGCATTAATTCGGCAGGAACCACACGGGCCTGCATGCGCAACGGCG GCAATTCATTTTACGCCGAACTTAAGTGGCTGGTGTCCAAGAGCAAGGGCCAGAACTTCCCCCAGACCACTAATACTTACCGCAACACCGA CACAGCAGAGCACCTTATCATGTGGGGTATACACCATCCTAGCAGTACCCAAGAAAAAAACGATCTGTACGGAACACAAAGCCTTTCAATTT CTGTAGGAAGTTCCACATATAGGAACAACTTTGTCCCGGTGGTGGGCGCTGGCCCTCAGGTTAATGGCCAGTCCGGGAGGATCGATTTCCA TTGGACACTCGTGCAGCCTGGGGACAACATTACTTTCTCTCACAACGGGGGTTTAATCGCCCCCTCAAGGGTCTCTAAGCTCATCGGGCGC GGTCTGGGAATCCAATCCGACGCTCCTATTGACAACAACTGCGAGTCAAAGTGCTTCTGGAGAGGTGGATCCATCAACACGCGCCTCCCAT TTCAGAACTTGTCCCCGCGTACAGTGGGCCAGTGTCCCAAGTATGTGAATCGACGGAGCTTGATGCTGGCCACTGGGATGAGGAACGTGC CTGAACTTATCCAGGGAAGGGGCCTCTTCGGAGCCATCGCTGGCTTCCTGGAAAACGGCTGGGAGGGGATGGTGGACGGGTGGTACGGC TTTCGACATCAGAATGCCCAGGGGACTGGACAGGCAGCAGATTATAAGAGTACCCAAGCCGCCATTGACCAGATCACAGGGAAGCTAAACA GGTTGGTCGAAAAGACTAACACAGAATTCGAATCCATCGAGAGCGAGTTTTCCGAGATTGAGCACCAGATCGGTAACGTGATTAACTGGAC GAAAGACTCAATAACCGATATCTGGACATATCAGGCGGAGTTGCTTGTGGCGATGGAGAACCAGCACACTATTGATATGGCAGACTCTGAG ATGCTCAACCTGTATGAGCGGGTTCGGAACCAGCTGCGACAAAATGCTGAGGAAGACGGAAAGGGGTGTTTCGAAATCTATCACGCCTGTG ATGACTCTTGCATGGAGTCGATTAGGAACAATACCTATGATCACTCTCAATATAGAGAGGAGGCTTTACTAAATCGGCTCAACATAAATCCAG TGACCCTGTCCAGCGGATATAAAGACATTATTCTATGGTTCTCTTTCGGAGCGAGCTGCTTTGTGCTTCTCGCTGTGGTGATGGGTCTCTTC TTTTTTTGCCTGAAGAATGGTAACATGCGCTGCACTATTTGTATT

2946	ATGTACAAGGTGGTAGTGATTATTGCCCTGCTGGGAGCAGTTAGAGGCTTAGATAAAATTTGCCTCGGGCATCATGCTGTTGCTAATGGTAC AATTGTCAAGACCTTAACAAATGAGCAGGAGGAGGTAACCAATGCGACCGAGACGGTGGAGTCTAAGTCTTTGGGGAAACTGTGTATGAAG GGAAGATCATATAACGACCTAGGCAATTGCCACCCCATCGGGATTCTGATTGGGACCCCCGCCTGTGATCTGCACCTGACAGGCACATGGG ACACCCTGATCGAACGGGAAAACGCGGTGGCCTATTGCTACCCTGGGGCGACAGTCAACGAGGAAGCTTTGAGGCAGAAAATCATGGAGA GCGGGGGCATCTCTAAAATCTCCACCGGGTTCACGTATGGTACCTCCATTAATTCAGCCGGAACCACCAAGGCATGTATGCGTAATGGAGG CAACTCCTTCTACGCCGAGCTCAAGTGGCTTGTTTCCAAAAATAAGGGGCAGAATTTCCCTCAGACAACGAACACATACCGGAATACTGACA CTGCAGAGCACCTAATCATCTGGGGAATTCACCACCCATCCAGTACGCAGGAAAAAAACGACCTCTACGGGACTCAGTCTCTGTCAATTTCT GTGGGCTCCTCAACATACCAAAATAACTTCGTTCCAGTGGTGGGCGCCCGGCCCCAGGTCAATGGACAGAGCGGAAGGATAGACTTCCAC TGGACACTCCTTCAGCCGGGGGACAATATCACGTTCTCTCACAACGGGGGACTGATAGCTCCATCTAGGGTGAGCAAACTGATTGGACGG GGATTGGGCATACAATCAGAAGCGCCCATAGACAATGGATGCGAAAGCAAATGCTTTTGGAAGGGAGGTTCAATAAACACAAAACTCCCATT CCAGAATCTGTCTCCTAGAACAGTGGGGCAGTGCCCCAAATATGTTAATAAGAGATCTCTCATGCTTGCTACTGGCATGCGGAACGTCCCC GAAATTATGCATGGGAGGGGCCTCTTCGGGGCGATTGCCGGCTTCATTGAAAATGGCTGGGAAGGAATGGTCGATGGATGGTATGGCTTT CGACATCAGAACGCACAGGGCACTGGGCAGGCAGCAGACTACAAAAGTACTCAGGCCGCAATAGACCAGATAACCGGAAAGCTCAATCGC CTCATCGAAAAGACAAACACTGAATTCGAAAGTATTGAATCCGAATTCTCTGAAATAGAGCACCAGATTGGTAATATCATCAACTGGACCAAG GATAGTATTACTGATATCTGGACATACCAAGCCGAACTGCTGGTAGCAATGGAGAATCAGCACACGATTGATATGGCAGATTCTGAAATGCT GAACCTGTATGAGCGAGTGCGCAAGCAGCTGCGCCAAAACGCTGAAGAGGATGGGAAAGGCTGCTTCGAAATCTACCACGCTTGCGACGA TTCCTGCATGGAGAGTATCCGGAATAACACCTACGATCATAGCCAGTACAGGGAGGAAGCCCTTCTGAACAGACTGAACATTAACCCAGTTA AGCTTAGTAGTGGATACAAAGACATAATCCTCTGGTTCTCCTTTGGCGCTTCATGCTTAATTTTACTTGCCGTCGTAATGGGACTGGTGTTTT TCTGCCTTAAGAACGGCAATATGAGGTGCACGATTTGCATC
2947	ATGTACAAGATCGTGGTTATTATTGCCTTGTTGGGCGCCGTGAAGGGTCTCGATAAGATTTGTCTGGGCCATCATGCTGTAGCCAATGGCAC CATTGTGAAGACGCTCACTAACGAGCAGGAAGAGGTTACGAACGCAACCGAGACAGTGGAGAGCACTGGGATAAACAGACTGTGCATGAA AGGACGTAAACACAAGGACCTTGGTAATTGCCACCCTATCGGGATGCTGATTGGGACACCCGCATGCGACTTGCATTTGACCGGAATGTGG GATACATTGATTGAGCGCGAGAACGCAATTGCCTACTGTTACCCAGGAGCCACCGTCAATGTTGAGGCCCTGCGGCAGAAGATCATGGAGA GCGGCGGCATTAACAAGATATCTACAGGCTTTACATATGGATCATCCATAAATTCAGCAGGAACTACGCGCGCATGCATGCGCAACGGCGG CAACAGCTTTTACGCTGAGCTTAAGTGGCTCGTCAGTAAGTCAAAGGGCCAGAATTTCCCCCAAACCACCAATACATACAGAAACACCGACA CGGCAGAGCACCTGATCATGTGGGGAATCCATCACCCATCTTCCACGCAAGAGAAGAATGACCTATACGGGACTCAGAGCCTCTCCATATC CGTCGGGTCCTCCACCTACAGGAACAACTTTGTACCTGTTGTAGGGGCTCGCCCCCAGGTGAATGGCCAGAGCGGACGGATCGACTTTCA TTGGACACTTGTTCAGCCCGGCGACAACATTACTTTCTCCCACAACGGAGGTCTGATCGCCCCCAGCAGGGTAAGTAAACTGATTGGCCGC GGCTTAGGAATACAGAGCGACGCCCCAATCGATAACAATTGTGAATCTAAGTGCTTTTGGAGGGGGGCAGTATAAACACCCGGCTACCCT TTCAGAATCTGAGTCCACGCACCGTGGGGCAGTGTCCAAAGTATGTGAACCGGAGAAGTTTAATGCTGGCAACCGGCATGAGAAATGTCCC TGAGCTAATTCAGGGTCGCGGGCTTTTCGGCGCAATCGCAGGCTTTCTCGAAAACGGTTGGGAGGGCATGGTGGACGGGTGGTACGGGTT CCGGCACCAAAACGCCCAGGGAACTGGGCAGGCCGCCGATTACAAAAGCACTCAGGCAGCTATAGACCAGATCACAGGCAAACTGAATCG GCTGGTGGAAAAGACAAATACAGAGTTCGAGAGCATTGAGAGCGAGTTTAGCGAGATCGAACATCAGATCGGGAACGTTATTAACTGGACC AAGGATAGCATCACTGATATTTGGACCTACCAGGCTGAACTGTTGGTCGCCATGGAGAACCAGCACACAATCGACATGGCAGACTCGGAGA TGCTTAACCTCTATGAGCGGGTGAGAAAGCAGCTCCGACAGAACGCCGAAGAGGACGGGAAGGGGTGCTTCGAAATTTATCATGCATGTG ACCATAGTTGTATGGAGTCAATCCCCAATAATACTTATGATCATAGCCAGTATCGTGAGGAGGCCCTGCTCAATCGTCTGAATATAAATCCC GTGACCCTGAGTTCGGGCTACAAAGACATAATACTCTGGTTTTCGTTTGGCGCGAGTTGCTTCGTGCTGTTAGCTGTTGTAATGGGATTGTT CTTTTTCTGCCTAAAAAACGGCAACATGAGGTGTACTATTTGCATC
2948	ATGTACAAGATTGTCCTGGTGCTTGCTCTGCTGGGAGCCGTGCACGGCCTCGATAAAATCTGCTTAGGCCACCACGCTGTCCCTAATGGCA CCATTGTTAAGACCTTGACAAACGAAAAGGAAGAGGTGACCAACGCCACTGAGACTGTGGAATCCAAGTCACTCGACAAATTATGTATGAAG AACAGGAACTATAAAGATCTTGGTAACTGCCATCCAATTGGCATGGTGGTAGGAACTCCTGCTTGCGATCTACACCTGACGGGCACATGGG ACACCCTGATTGAGAGGGACAACTCTATTGCATACTGCTATCCAGGTGCGACCGTTTCCGAAGAAGCACTCCGGCAGAAGATTATGGAGTC CGGCGGCATCGATAAAATCTCCACCGGCTTCACATACGGAAGTAGCATCAACTCTGCTGGGACGACCAAAGCGTGTATGAGGAACGGGGG GAATAGTTTCTACTCAGAACTCAAGTGGCTGGTCAGTAAGAACAAGGGGCAGAACTTCCCTCAAACGACCAACACATACCGAAATACAGATT CTGTGGAGCACTTAATCATCTGGGGGATTCATCATCCATCCAGCACGCAGGAGAAAAACGATCTCTATGGTACACAGTCTCTCAGCATCAGT GTGGGAAGCAGTACGTATCAGAACAATTTTGTCCCCGTCGTTGGCGCCCGGCCACAAGTGAACGGACAAAGTGGTCGTATTGATTTTCATT GGACCATGGTCCAACCTGGTGATAATATAACATTTTCACACAATGGAGGACTGATAGCTCCTAACAGAGTATCCAAGCTGAAGGGAAGGGG GTTGGGGATTCAGTCCGGAGCATCAGTGGACAATGATTGCGAGAGCAAGTGCTTCTGGAAGGGAGGTTCCATCAACACGAAGCTGCCTTTC CAGAATCTGAGCCCACGAACAGTGGGACAGTGCCCGAAGTATGTGAATAAGAAGAGCCTTCTGCTTGCGACTGGCATGAGGAATGTTCCTG AAGTTGCGCAGGGAAGGGGGCTATTCGGGGCAATCGCCGGATTCATCGAGAACGGCTGGGAGGGCATGGTGGACGGGTGGTACGGGTTC AGGCACCAGAACGCCCAGGGTACAGGGCAGGCCGCTGATTACAAGTCAACACAGGCGGCGATAGACCAGATCACCGGCAAACTGAATAG GCTAATAGAGAAGACTAACACTGAATTTGAATCTATTGAGAGCGAGTTCTCGGAAATTGAACACCAGATCGGAAATGTGATCAATTGGACCA AGGATTCCATTACCGATATCTGGACTTATCAGGCAGAGCTGTTAGTGGCAATGGAAAACCAGCATACCATAGACATGGCTGATTCTGAGATG TTGAACCTCTACGAGCGGGTACGGAAACAACTGAGACAGAATGCAGAGGAAGACGGCAAGGGCTGTTTCGAAATCTACCACAAGTGCGAC GATAATTGTATGGAAAGTATACGCAACAATACCTATGATCACACACAATATAGGGAAGAGGCTCTTCTGAACCGACTGAACATCAATCCCGT GAAACTTAGCTCAGGATACAAGGACGTGATCCTTTGGTTTTCTTTCGGGGCTAGTTGCTTCGTGCTGCTTGCGGTTATCATGGGGCTTGTTT TCTTTTGTCTAAAGAATGGGAATATGAGATGCACAATCTGCATC

2949	ATGTACAAAATCGTGGTGATCATCGCCCTCCTGGGAGCTGTGAAGGGCCTTGACAAAATCTGCCTCGGCCACCACGCCGTCGCCAATGGAA CCATCGTGAAGACTCTGACCAATGAGCAGGAGGAAGTTACCAATGCAACTGAGACCGTGGAAAGTACCGGGATAAATCGCCTCTGCATGAA GGGAAGGAAACATAAAGACCTGGGGAACTGTCATCCGATTGGGATGCTGATTGGGACCCCTGCCTGTGATTTGCACCTTACCGGCACATGG GACACACTGATCGAGCGAGAAAATGCTATCGCATATTGTTATCCCGGGGCAACCGCAAACGTAGAGGCCCTACGGCAGAAAATTATGGAGT CCGGTGGCATAGACAAAATAAGCACAGGATTTACCTACGGCTCGAGCATCAACTCTGCTGGCACCACCAGAGCTTGCATGAGGAATGGCG GGAACTCCTTTTACGCTGAGTTGAAGTGGCTGGTGAGCAAGAGTAAAGGGCAGAACTTTCCTCAGACGACTAATACATATAGAAACACCGAC ACAGCAGAGCACCTCATAATGTGGGGAATCCATCACCCATCTTCAATTCAGGAGAAGAATGACCTTTATGGTACTCAGAGCCTCAGTATTTC CGTGGGATCTTCTACGTACAGAAACAACTTTGTGCCAGTTGTCGGCGCAAGACCTCAGGTCAACGGGCAGTCAGGGAGAATTGACTTTCAT TGGACACTGGTGCAACCTGGGGATAATATCACTTTTAGCCATAATGGAGGACTGATTGCTCCCAGTAGAGTCTCTAAACTGATTGGGAGGG GACTCGGTATCCAGTCGGATGCTCCTATCGATAATAACTGTGAGAGCAAGTGTTTTTGGCGAGGGGGATCTATCAATACACGGTTGCCATTT CAGAATCTATCGCCCCGCACGGTTGGTCAGTGCCCCAAGTACGTTAATCGTCGGTCGCTGATGCTGGCTACGGGAATGAGGAATGTTCCAG AACTGATTCAAGGACGCGGTTTGTTCGGCGCCATCGCAGGTTTCCTTGAAAACGGCTGGGAAGGTATGGTTGACGGTTGGTACGGATTCCG ACATCAGAACGCCCAGGGTACGGGACAGGCGGCTGACTACAAAAGCACCCAGGCAGCAATTGATCAGATCACGGGGAAGCTCAATAGACT GGTGGAAAAGACCAACACGGAATTTGAATCAATTGAGTCCGAATTTTCCGAGATTGAACATCAGATCGGCAATGTCATTAATTGGACAATGG ATAGCATCACTGACATATGGACTTATCAGGCCGAGCTGCTGGTTGCAATGGAGAATCAGCACACTATCGACATGGCTGATTCCGAGATGCT GAACCTTTACGAGAGAGTGCGCAAGCAGCTCCGGCAGAACGCCGAAGAAGATGGAAAGGGGTGCTTTGAGATTTATCACGCGTGCGACGA CAGTTGCATGGAATCCATCCGGAACAATACTTATGATCATTCTCAGTACCGCGAGGAAGCACTGCTCAACAGACTGAATATTAACCCAGTGA CCCTAAGCTCAGGGTACAAGGACATCATTCTGTGGTTCTCTTTCGGCCCGTCATGCTTCGTGCTGCTGGCAGTTGTGATGGGACTCGTGTT CTTTTGCCTCAAAAATGGAAACATGAGATGTACCATTTGTATC
2950	ATGTATAAAGTCGTAGTGATCATCGCACTGTTAGGAGCTGTAAAGGGACTGGACAAGATCTGCTTAGGACACCACGCAGTGGCAAACGGGA CCATAGTCAAGACATTGACTAATGAACAAGAGGAAGTCACCAACGCGACTGAAACAGTCGAATCTACAGGAATCAACAGACTTTGTATGAAA GGACGGAAACATAAAGATCTGGGCAACTGTCACCCCATAGGAATGCTGATTGGTACACCAGCTTGTGACTTGCATCTGACCGGCACATGGG ACACGCTGATCGAGAGAGAAAATGCCATTGCCTACTGTTACCCTGGGGCTACAGTAAACGTCGAGGCGCTGCGTCAGAAGATCATGGAATC TGGGGGCATCGATAAAATCAGCACGGGGTTCACATACGGATCCAGTATTAATTCTGCTGGCACTACCCGTGCTTGTATGCGGAACGGTGGA AACTCCTTCTATGCAGAGCTCAAATGGCTGGTTTCTAAGAACAAGGGGCAGAATTTTCCCCAGACGACAAATACATATAGAAATACAGACAC TGCTGAGCACCTAATCATGTGGGGCATCCACCACCCAAGTAGTATCCAGGAGAAAAACGACCTGTACGGCACACAAAGCCTGTCAATCTCA GTGGGCTCCTCAACTTACCGCAATAATTTTGTGCCAGTTGTGGGCGCCAGGCCCCAGGTTAATGGTCAGTCTGGGCGCATCGATTTTCATT GGACTCTGGTGCAACCTGGCGACAATATTACATTCTCTCACAACGGAGGACTCATCGCCCCATCCAGAGTGTCTAAGTTGATAGGACGAGG GTTAGGCATCCAAAGCGATGCCCCCATTGACAATAACTGCGAATCAAAGTGCTTTTGGCGGGGAGGATCCATTAACACACGGTTGCCTTTC CAGAATCTGTCTCCAAGGACCGTAGGGCAGTGCCCAAAATACGTGAACAGGCGGTCTCTTATGCTGGCCACGGGGATGCGCAACGTGCCA GAGCTCATCCAGGGGCGTGGATTATTTGGAGCTATTGCCGGATTTCTGGAAAATGGCTGGGAGGGGATGGTTGACGGTTGGTACGGTTTC AGGCACCAGAACGCACAGGGCACAGGACAGGCCGCCGACTATAAAAGTACCCAGGCAGCGATTGATCAGATAACCGGAAAGCTGAACAGG TTGGTGGAAAAAACCAATACTGAGTTCGAGTCTATCGAGAGCGAATTCTCGGAAATCGAACACCAGATTGGAAATGTAATCAACTGGACTAT GGATAGCATCACCGATATCTGGACCTACCAGGCTGAACTGTTGGTGGCTATGGAGAATCAACACACGATCGATATGGCTGACAGCGAGATG CTCAACCTGTATGAACGTGTTAGGAAACAGCTAAGGCAAAACGCTGAAGAAGACGGAAAGGGATGTTTTGAAATATATCATGCATGCGATGA CTCATGTATGGAGTCAATCCGGAACAATACATATGACCATTCTCAGTATAGGGACCAGCCCCTCCTGAACAGACTGAATATTAATCCCGTTA CCCTATCCAGCGGGTATAAGGACATCATCCTCTGGTTCTCTTTTGGTGCCAGTTGTTTCGTTTTGCTTGCTGTTGTGATGGGACTCGTCTTTT TCTGTCTGAAGAACGGGAACATGAGATGCACCATTTGCATC
2951	ATGTACAAAATTGTCGTCATCATAGCACTTCTAGGTGCCGTGAAAGGTTTGGATAAGATCTGTCTAGGCCATCATGCTGTCGCAAACGGCAC TATCGTGAAAACTTTGACTAACGAACAAGAGGAAGTGACTAACGCTACGGAGACAGTCGAATCGACCGGGATCAACAGACTGTGCATGAAA GGCCGCAAACACAAAGATTTGGGGAACTGCCATCCTATAGGCATGCTGATTGCGACACCAGCTTGCGACCTCCACCTGACGGGAATGTGG GACACCCTCATTGAGCGGGAGAATGCGATCGCGTACTGTTACCCGGGGGCTACCGTCAATGTGGAGGCACTCAGGCAAAAGATCATGGAG TCTGGCGGCATCAATAAGATTTCCACTGGGTTCACCTATGGATCTAGCATTAATTCTGCAGGTACAACTAGAGCTTGTATGAGAAATGGGGG GAATAGTTTCTACGCCGAACTTAAATGGCTGGTGAGCAAGTCAAAGGGACAAAACTTCCCCCAGACAACGAATACTTACAGGAATACGGATA CCGCAGAGCACTTGATCATGTGGGGGATTCATCATCCATCAAGCACACAGGAAAAAAACGATTTGTATGGGACTCAGTCACTGTCTATTAGC GTGGGATCTAGTACTTACAGAAACAACTTCGTACCTGTGGTGGGAGCCCGACCACAGGTGAATGGACAGAGTGGGAGGATTGACTTTCACT GGACCCTGGTGCAACCTGGTGATAATATTACATTTAGCCACAATGGCGGATTGATCGCACCATCCCGAGTCAGTAAGCTCATAGGCCGGGG ACTTGGCATACAGAGCGACGCTCCTATCGATAACAATTGCGAGTCCAAATGCTTCTGGCGGGGAGGAAGTATAAACACACGGCTGCCATTT CAGAACCTTTCGCCCCGCACGGTCGGCCAGTGCCCCAAATATGTGAATAGGCGTTCACTCATGCTCGCAACAGGAATGAGAAATGTGCCTG AGCTGATTCAGGGCAGGGGGCTTTTCGGTGCTATAGCGGGATTTTTAGAGAACGGCTGGGAAGGGATGGTTGATGGGTGGTACGGGTTCC GGCACCAAAATGCGCAGGGAACAGGACAGGCTGCCGACTACAAATCTACCCAGGCTGCAATTGATCAAATAACTGGCAAACTCAATAGGTT GGTGGAAAAGACGAACACCGAGTTCGAGTCAATCGAATCTGAATTCAGCGAGATTGAACACCAAATCGGAAATGTGATTAACTGGACCAAG GACTCCATCACCGACATCTGGACCTATCAGGCCGAACTGCTCGTCGCTATGGAGAACCAGCATACCATAGATATGGCCGACAGCGAGATGC TCAACCTTTACGAGCGCGTCAGGAAACAGCTTAGGCAGAATGCTGAAGAAGACGCGGAAGCGCATGTTTTGAGATCTATCACGCTTGCGATGA TTCCTGTATGGAGTCCATAAGGAACAACACCTATGACCATTCACAGTACCGTGAAGAGGCCTTGCTTAACCGCCTCAACATAAATCCCGTTA CTCTCAGCTCAGGTTATAAGGATATCATCCTGTGGTTTAGCTTTGGAGCCTCCTGCTTTGTGCTCCTCGCCGTCGTGATGGGCCTATTCTTTT TTTGTCTGAAAAACGGGAATATGCGATGCACTATATGTATA

2952	ATGTATAAGATCGTGGTGATCATTGCACTCCTGGGAGCCGTCAAGGGCCTGGATAAAATATGCCTGGGGCACCACGCCGTTGCCAACGGAA CTATAGTAAAGACCCTGACCAATGAACAGGAAGAAGTGACCAACGCAACCGAAACCGTGGAGAGTACAGGCATAAATCGCCTTTGCATGAA GGGTCGAAAGCACAAAGACCTGGGGAATTGCCACCCCATAGGGATGCTGATCGGCACCCCGGCTTGTGATTTGCACTTGACTGGGATGTG GGATACACTCATTGAGCGCGAAAACGCAATCGCATATTGCTACCCCGGGGCAACCGTGAACGTCGAGGCCCTGCGGCAGAAAATTATGGA GTCTGGCGGAATAAATAAGATTTCAACCGGTTTTACATACGGAAGTTCTATTAATAGCGCAGGCACAACAAGGGCGTGTATGAGGAACGGT GGTAATAGTTTCTATGCAGAACTAAAATGGTTGGTGTCTAAATCGAAAGGTCAAAATTTCCCCCAAACTACCAACACCTACCGCAATACTGAT ACTGCCGAACACTTGATTATGTGGGGTATTCACCACCCATCCTCAACACAGGAAAAGAACGATTTGTACGGGACACAGAGTCTGAGTATTAG TGTAGGCTCTAGCACCTACAGAAATAACTTTGTCCCAGTGGTGGGCGCTCGGCCTCAGGTGAATGGCCAATCTGGAAGAATTGACTTCCAT TGGACACTTGTGCAGCCAGGTGATAACATCACTTTTAGCCATAATGGTGGGCTGATTGCCCCTTCCAGGGTGAGTAAACTTATTGGTCGCG GGCTGGGGATACAATCGGATGCCCCTATCGACAATAATTGCGAATCCAAGTGTTTTTGGAGGGGCGGGTCAATCAATACCAGGCTACCTTT CCAGAATCTGAGTCCACGTACCGTCGGGCAGTGTCCTAAGTATGTCAATCGACGGTCACTGATGCTCGCTACCGGCATGCGGAATGTCCCA GAGCTGATCCAAGGGCGTGGTCTGTTCGGCGCAATTGCCGGCTTCCTCGAGAATGGATGGGAGGGCATGGTAGATGGATGGTATGGTTTT AGACACCAAAACGCCCAAGGAACTGGGCAGGCGGCCGATTATAAATCCACTCAGGCCGCAATTGATCAGATTACTGGCAAGCTAAACCGTT TAGTTGAGAAAACGAACACCGAATTCGAGAGCATAGAGAGTGAATTTAGTGAGATCGAGCATCAAATTGGGAATGTGATTAACTGGACCAAA GACTCAATAACCGATATCTGGACCTACCAGGCAGAGTTACTGGTAGCTATGGAGAACCAGCATACCATCGACATGGCCGACTCTGAGATGC TCAATTTGTATGAGAGGGTCAGGAAACAACTGCGGCAGAATGCTGAAGAAGATGGAAAAGGGTGCTTTGAAATATATCACGCATGCGACGA TAGTTGCATGGAGTCCATACGAAACAATACCTACGATCACTCACAGTATAGAGAGGAAGCCCTCTTAAATAGGCTAAATATAAACCCTGTCAC ACTGAGCTCTGGGTACAAAGATATTATCTTATGGTTCTCATTTGGGGCTAGTTGTTTCGTGCTCCTGGCGGTCGTGATCGCCCTGTTCTTCT TCTGTCTAAAGAACGGTAATATGAGATGTACTATCTGCATT
2953	ATGTATAAAGTGGTGGTTATTATAGCGTTGCTGGGCGCTGTTAAAGGTTTGGACAAGATCTGCTTGGGACATCATGCTGTCGCCAACGGGA CAATCGTCAAGACCCTGACAAACGAACAGGAAGAGGTCACAAACGCGACTGAAACCGTGGAGAGCACAGGAATTAATCGGCTCTGTATGAA AGGCCGCAAACACAAAGATCTTGGAAATTGCCACCCAATTGGGATGCTCATTGGCACACCAGCCTGTGATCTACATCTGACCGGCACATGG GACACTCTCATCGAACGCGAGAACGCTATCGCATATTGTTACCCCGGGGCGACCGTGAATGTTGAGGCCCTGCGCCAGAAGATTATGGAG AGTGGAGGCATCGATAAGATATCGACCGGCTTCACTTACGGAAGCTCCATTAACTCCGCTGGTACCACCCGGGCCTGCATGAGGAACGGC GGCAATTCCTTTTACGCCGAACTGAAGTGGTTAGTGTCTAAAAATAAGGGGCAGAATTTTCCCCAAACCACAAACACATACCGAAACACCGA CACGGCCGAGCATCTGATCATGTGGGGGATTCACCATCCTTCTTCCATCCAGGAAAAGAACGATCTGTACGGAACTCAATCTCTCTCCATCT CCGTTGGTTCCTCCACGTATAGGAACAATTTCGTACCCGTGGTCGGCGCCCGACCACAAGTAAACGGCCAGTCCGGCCGCATCGACTTCC ACTGGACACTTGTGCAGCCAGGAGACAACATTACATTCTCCCATAATGGCGGGCTGATTGCACCCAGCCGGGTCTCCAAGCTTATCGGCCG CGGCCTGGGGATCCAGTCAGACGCGCCCATCGATAACAACTGCGAAAGCAAGTGTTTTTGGCGGGGTGGATCTATTAACACCCGTCTGCCT TTCCAGAATCTTAGTCCCCGGACAGTGGGGCAGTGCCCCAAGTATGTGAATCGAAGGTCTCTGATGCTGGCCACAGGCATGAGAAATGTGC CAGAGTTGATTCAGGGCCGGGGTCTCTTTGGAGCCATTGCCGGTTTTCTTGAAAACGGTTGGGAGGGTATGGTCGACGGTTGGTACGGTTT CCGGCACCAGAACGCACAGGGCACTGGTCAGGCTGCAGATTATAAATCTACACAAGCCGCTATCGACCAGATCACGGGCAAACTAAATAGA CTGGTCGAGAAGACTAACACTGAATTTGAGAGCATTGAGTCGGAGTTTTCCGAGATTGAACACCAGATCGGAAATGTGATTAATTGGACAAT GGACTCAATTACTGACATCTGGACATACCAAGCCGAACTGCTGGTTGCGATGGAGAACCAGCACACCATCGATATGGCCGATAGCGAAATG CTAAATCTGTATGAGAGGGTAAGGAAGCAGCTGCGCCAGAATGCCGAAGAAGACGGGAAGGGTTGTTTTGAGATCTACCATGCTTGTGACG ACAGTTGCATGGAAAGCATTCGAAACAACACTTATGACCACTCGCAATATCGCGAAGACGCCCTTTTGAATCGGCTCAACATAAACCCAGTG ACACTTTCTTCCGGTTACAAGGATATTATCCTGTGGTTCTCTTTTGGAGCATCGTGTTTTGTACTGCTCGCTGTCGTTATGGGCTTGGTCTTC TTTTGTCTGAAAAATGGCAATATGCGGTGTACCATCTGCATT
2954	ATGTATAAGATTGTGGTGATCATTGCACTGCTGGGGGCTGTGAAGGGACTCGACAAGATTTGCCTGGGCCACCATGCCGTTGCAAACGGGA CCATCGTCAAAACATTGACAAACGAGCAGGAAGAGGTGACAAACGCCACTGAAACAGTCGAAAGTACGGGGATTAACCGGCTATGTATGAA GGGGAGGAAACATAAAGATCTAGGCAACTGCCACCCCATAGGCATGCTTATCGGCACCCCGGCCTGCGACCTTCACCTGACCGGAATGTG GGATACCTTGATCGAGAGGGAAAACGCCATCGCTTATTGCTACCCAGGAGCCACGGTGAACGTAGAGGCTCTACGACAAAAGATTATGGAG AGCGGCGGAATAAATAAAATATCCACGGGATTCACCTACGGTTCCTCGATAAATTCAGCGGGCACCACCAGAGCCTGCATGCGCAACGGCG GCAATTCCTTTTATGCCGAGCTTAAGTGGCTCGTTTCCAAAAGCAAGGGGCAGAACTTTCCTCAGACCACAAACACCTATAGGAATACCGAC ACCGCGGAGCACCTGATTATGTGGGGGATCCACCATCCCTCAAGTACACAGGAAAAGAACGATCTCTATGGAACCCAGAGCCTGAGTATCA GCGTGGGGTCCAGCACATATCGCAACAACTTTGTCCCAGTGGTCGGAGCCCGCCCACAGGTTAACGGCCAATCCGGTCGCATTGACTTTC ACTGGACCTTGGTGCAACCAGGGGATAATATTACCTTCAGCCACAATGGTGGACTGATTGCCCCAAGTCGTGTTTCCAAGCTGATCGGAAG AGGGTTAGGTATCCAGAGTGACGCCCCAATTGATAATAACTGCGAGAGCAAATGCTTCTGGCGGGGGGGATCGATTAATACGCGTCTCCCC TTTCAGAACCTCTCCCCACGTACGGTCGGGCAGTGTCCAAAGTATGTGAACAGACGTTCACTGATGCTGGCAACAGGGATGAGAAACGTAC CCGAGCTGATCCAGGGGAGAGGGCTGTTCGGTGCCATTGCCGGATTTCTCGAAAATGGTTGGGAGGGAATGGTGGACGGTTGGTACGGAT TCCGGCACCAGAACGCTCAGGGGACCGGCCAAGCAGCTGATTATAAATCTACCCAGGCCGCCATTGATCAGATCACCGGGAAGTTGAACC GCCTCGTGGAGAAGACTAACACAGAGTTCGAGTCTATCGAATCAGAGTTCAGCGAGATTGAACACCAGATTGGGAACGTGATTAACTGGAC CAAGGACTCCATCACCGACATTTGGACATATCAAGCCGAGCTGCTGGTAGCCATGGAGAACCAGCACACCATAGATATGGCAGATAGTGAG ATGCTGAACCTCTATGAACGCGTGCGAAAGCAACTGCGACAAAATGCTGAGGACGATGGGAAGGGATGCTTTGAAATCTACCACGCGTGC GACGATTCTTGTATGGAAAGTATCCGGAATAATACTTATGATCACTCTCAGTATAGGGAGGAGGCACTACTAAATCGTCTGAATATCAACCCA GTGACCTTATCCTCCGGATACAAGGACATTATCCTGTGGTTCAGTTTTGGCGCTTCCTGTTTCGTGCTCCTGGCTGTAGTGATGGGCTTATT TTTCTTCTGCCTGAAAAACGGAAATATGCGGTGCACCATTTGCATT

2955	ATGTATAAGATAGTCGTAATTATCGCACTCCTCGGGGCTGTTAAGGGCCTGGACAAAATCTGTCTGGGTCACCATGCAGTCGCCAACGGAA CAATAGTCAAGACTCTGACCAATGAACAGGAGGAGGTTACGAATGCCACGGAAACCGTGGAAAGTACAGGGATCAACCGCCTGTGTATGAA AGGGAGAAAGCATAAAGACTTGGGTAATTGCCATCCAATCGGGATGCTGATTGGAACACCCGCTTGTGACCTCCATTTGACCGGTATGTGG GACACACTGATCGAAAGAGAGAATGCCATCGCGTATTGTTACCCTGGTGCCACTGTGAACGTGGAGGCACTGCGCCAGAAGATCATGGAAT CCGGCGGAATCAATAACATCTCCACCGGATTTACCTACGGATCCAGTATAAATTCACCCGGACGACAAGAGCATGCATGAGAAACGGCGG CAACAGTTTTTACGCCGAGCTGAAGTGGCTTGTGTCCAAGTCTAAAGGCCAGAACTTTCCTCAGACAACCAACACCTACCGTAACACTGATA CAGCAGAGCATCTGATCATGTGGGGCATTCACCATCCATCTTCCACCCAAGAAAAGAATGATCTGTACGGAACACAGAGCCTGTCTATTAGT GTTGGATCGTCAACCTACCGGAATAACTTTGTTCCCGTAGTGGGCGCACGCCCCCAGGTGAATGGGCAGTCAGGTCGAATTGATTTTCACT GGACCCTCGTGCAGCCTGGCGACAATATTACATTCAGTCATAACGGGGGACTGATCGCCCCAAGCCGCGTTTCTAAGCTGATTGGGAGAG GTTTGGGAATTCAGAGCGACGCACCTATCGATAACAATTGCGAATCAAAGTGCTTTTGGCGGGGGGGGCAGCATAAATACCAGACTGCCCTT TCAGAACCTTAGCCCTAGGACAGTCGGGCAGTGTCCCAAATACGTCAACCGCCGGAGTCTGATGCTCGCTACAGGAATGAGAAATGTTCCA GAGCTAATCCAGGGACGAGGACTCTTCGGAGCCATCGCTGGATTTCTGGAGAATGGCTGGGAGGGGATGGTTGATGGGTGGTATGGCTTT AGACATCAGAACGCCCAGGGGACAGGTCAGGCCGCCGATTACAAGTCTACCCAGGCCGCGATTGACCAGATCACTGGCAAATTGAATAGG CTGGTTGAGAAAACTAACACTGAATTTGAGTCAATCGAGAGCGAGTTTTCTGAGATCGAACACCAGATCGGGAACGTGATTAATTGGACTAA AGACTCTATAACGGACATTTGGACCTATCAAGCCGAGTTATTAGTCGCGATGGAGAACCAGCACACAATAGATATGGCAGACTCTGAGATGC TGAACCTGTACGAGAGGGTGCGGAAGCAATTAAGGCAGAATGCCGAAGAGGATGGGAAGGGATGCTTCGAAATATACCATGCCTGCGACG ACTCGTGCATGGAGAGCATTAGAAACAATACATACGACCATAGCCAGTATCGCGAGGAAGCACTACTGAATCGGCTGAATATCAATCCAGTC ACACTCTCTTCTGGGTACAAAGATATCATTCTGTGGTTTAGCTTCGGCGCCTCCTGCTTCGTCCTTCTGGCCGTGGTTATGGGGCTGTTTTTT TTCTGTCTTAAGAACGGAAATATGAGGTGTACCATCTGTATT
2956	ATGTATAAGATTGTGGTGATCATTGCCCTCCTAGGCGCCGTAAAAGGGTTAGATAAAATCTGCCTGGGACACCACGCCGTGGCCAACGGCA CTATAGTCAAGACCCTCACTAACGAGCAGGAAAAAGTGACCAATGCTACGGAAACAGTGGAATCTACAGGCCTGAATCGCCTTTGCATGAAA GGCAGGAAACACAAAGACTTAGGGAATTGTCACCCCATTGGAATGTTAATTGGGACTCCAGCTTGCGATCTGCACCTCACTGGTACATGGG ATACAATTATTGAACGGGAGAATGCCATCGCATACTGCTACCCTGGGGCGACCGTCAATGAGGAAGCTCTGCGGCAGAAGATCATGGAGTC TGGTGGCATTGATAAAATCTCCACCGGGTTCACGTATGGGTCTTCAATTAATAGCGCTGGAACGACTAGAGCTTGTATGCGCAATGGTGGAA ATTCCTTTTACGCTGAGCTTAAATGGCTCGTCTCGAAATCCAAGGGACAGAACTTTCCGCAAACTACAAACACTTATAGAAACACAGACACC GCAGAACACCTCATCATGTGGGGGATTCACCACCCTTCTAGTACCCAAGAAAAGAATGACCTCTATGGCACACAGAGCCTGTCCATCTCGG TCGGCAGCAGCACCTACCGGAACAACTTTGTGCCCGTGGTGGGCGCGCGCCCCCAAGTTAACGGACAGTCCGGTCGGATAGATTTCCACT GGACACTGGTCCAGCCCGGGGATAACATAACGTTCTCTCACAACGGAGGGCTGATAGCTCCTTCTCGAGTGTCAAAATTGATTGGAAGGGG CTTGGGGATCCAGTCCGATGCCCCAATAGACAATAACTGCGAGTCTAAATGTTTTTGGAGGGCGCGGTTCTATAAATACCCGTCTGCCTTTTC AGAATCTTAGCCCCAGAACAGTGGGACAGTGTCCCAAGTACGTAAATAAAAGAAGTCTGATGCTTGCAACCGGCATGAGGAATGTCCCTGA GCTGATCCAGGGTCGTGGCTTGTTCGGTGCCATTGCGGGCTTCTTGGAAAATGGTTGGGAGGGAATGGTAGACGGATGGTATGGGTTCCG CCACCAAAATGCTCAGGGGACGGGCCAAGCTGCCGACTATAAGAGTACGCAAGCCGCTATTGACCAGATCACTGGCAAATTGAACAGACTT GTTGAAAAAACAAATACAGAATTCGAAAGCATCGAATCAGAATTTAGCGAGATTGAGCACCAAATCGGCAACGTGATCAATTGGACAAAAGA CTCCATAACCGATATTTGGACCTACCAGGCCGAGCTGCTCGTGGCTATGGAGAACCAGCACACCATCGACATGGCTGATAGCGAGATGCTG AATCTATATGAAAGGGTTCGGAAGCAGTTGCGTCAGAATGCGGAGGAGGATGGGAAGGGTTGCTTTGAAATCTATCACGCCTGCGATGACA GCTGTATGGAGAGCATACGAAATAATACCTATGACCATTCACAGTACCGCGAGGAGGCCCTGCTCAATCGGCTGAACATCAATCCGGTCAC ACTGAGCTCTGGCTACAAGGACATCATTCTGTGGTTCTCATTCGGGGCGAGCTGCTTTGTTTTGTTAGCCGTCGTGATGGGCCTCGTTTTCT TCTGCCTGAAGAATGGGAATATGCGCTGTACAATCTGCATC
2957	ATGTACAAGATTGTCGTGATTATCGCTTTGCTCGGCGCCGTGAAGGGACTGGATAAAATCTGTCTGGGGCATCACGCTGTGGCTAATGGAA CAATTGTGAAGACCCTCACCAATGAGCAGGAGGAAGTGACAAACGCTACAGAGACAGTCGAGTCCACCGGAATCAACAGGCTGTGCATGAA GGGCCGGAAACATAAGGACCTCGGGAACTGCCACCCTATTGGGATGCTGATAGGCACACCTGCCTGTGACTTACACCTTACCGGATGTG GGATACACTGATAGAACGGGAGAATGCCATAGCATACTGCTACCCAGGAGCTACGGTGAATGTGGAAGCTCTCCGCCAGAAAATTATGGAA TCAGGCGGGATTAATAAGATTTCTACGGGCTTCACTTATGGAAGTTCAATAAATTCAGCCGGAACAACCCGGGCTTGTATGAGAAATGGTGG AAATTCTTTCTACGCAGAACTCAAGTGGCTTGTTAGCAAGTCCAAGGGCCAAAACTTCCCTCAGACCACTAATACTTACAGGAATACTGACAC CGCCGAACACCTTATCATGTGGGGTATCCACCATCCATCTAGCACCCAGGAGAAAAATGATCTTTACGGTACCCAATCACTGAGCATCAGCG TGGGGAGCTCAACTTATCGCAACAATTTCGTGCCTGTGGTGGGTGCCCGACCACAGGTTAACGGACAAAGTGGACGAATTGACTTTCATTG GACCCTGGTGCAGCCTGGCGACAACATCACATTCTCGCATAATGGAGGTTTGATCGCTCCCTCCCGTGTGTCTAAGCTGATAGGGAGAGGC CTCGGTATACAGTCCGACGCTCCCATCGATAATAATTGTGAAAGCAAGTGCTTTTGGCGAGGTGGCAGCATTAACACGCGGTTACCCTTTCA GAACCTCAGCCCACGCACAGTCGGCCAATGCCCTAAGTATGTAAACCGCCGGAGCTTGATGCTGGCAACGGGCATGAGAAACGTCCCCGA GCTAATCCAGGGCAGAGGACTGTTCGGCGCCATTGCTGGCTTTCTTGAGAACGGATGGGAGGGAATGGTGGACGGCTGGTACGGATTCCG GCACCAGAACGCCCAGGGCACCGGCCAGGCCGCCGATTATAAGTCCACCCAAGCTGCCATCGACCAGATCACCGGCAAACTGAATCGGCT TGTAGAGAAAACCAACACCGAATTCGAGTCAATCGAGTCCGAATTTAGCGAGATCGAACACCAGATCGGCAATGTGATTAACTGGACAAAG GACAGTATCACTGATATCTGGACCTACCAAGCAGAATTACTGGTCGCTATGGAAAATCAGCACACCATTGATATGGCCGACAGCGAGATGCT CAACCTATATGAACGCGTCCGCAAGCAGCTGCGTCAGAACGCAGAGGAAGATGGAAAAGGCTGCTTCGAGATCTACCATGCTTGCGATGAC TCTTGTATGGAGTCCATCCGCAATAACACATACGACCACTCCCAGTACAGGGAGGAGGCGCTTTTGAACAGACTGAATATAAATCCTGTTAC TCTTTCATCGGGGTATAAAGACATCATCCTGTGGTTCTCCTTCGGCGCCTCCTGTTTCGTCCTGCTGGCGGTAGTTATGGGCCTATTTTTTTT TTGTCTGAAGAATGGGAACATGCGCTGTACAATTTGTATC

2958	ATGTACAAAATTGTTGTCATCATTGCTCTGCTGGGGGCTGTGAAAGGACTGGACAAAATCTGCTTAGGCCATCACGCTGTAGCCAATGGTAC CATTGTGAAGACACTGACCAATGAACAGGAAAAAGTAACAAATGCTACCGAGACGGTCGAGTCCACAGGGCTGAACCGATTGTGCATGAAG GGACGCAAGCACAAGGATCTGGGAAATTGCCACCCGATTGGTATGCTGATTGGGACTCCAGCCTGCGATCTGCATCTGACCGGGACTTGG GATACTATCATAGAACGGGAGAACGCCATCGCCTACTGCTACCCAGGTGCCACTGTAAACGAGGAAGCCTTGCGCCAGAAAATTATGGAGT CACGTGGCATTGACAAGATTTCCACTGGGTTCACTTATGGGAGCAGTATTAACTCAGCTGGCACAACCCGTGCTTGTATGCGGAACGGAGG AAACTCATTCTACGCAGAGCTGAAATGGCTGGTTAGCAAAAGCAAGGGCCAGAATTTCCCACAGACAACCAATACATATCGGAACACTGATA CCGCAGAGCACCTGATTATGTGGGGCATTCACCACCCATCCTCCACGCAAGAGAAGAATGATCTGTATGGAACTCAGAGCCTGTCAATCTC CGTGGGATCTAGCACATATAGAAATAATTTTGTGCCTGTGGTCGGCGCGCGACCACAGGTTAATGGACAATCAGGACGCATTGATTTCCACT GGACCTTGGTTCAGCCCGGGGACAATATAACATTCAGCCACAACGGTGGCCTGATCGCCCCCTCGCGGGTGAGTAAGCTGATTGGCAGAG GACTCGGTATTCAGAGCGATGCCCCTATTGATAATAACTGCGAAAGCAAATGTTTTTGGAGAGGCGGGTCGATCAACACCAGATTACCCTTT CAGAATCTCTCTCCTAGAACCGTTGGTCAGTGCCCGAAATATGTCAACAAGCGTAGTTTGATGCTAGCCACTGGAATGCGCAACGTACCAGA ACTCATCCAGGGAAGGGGTTTGTTCGGAGCGATTGCAGGTTTCCTGGAAAATGGGTGGGAGGGAATGGTGGACGGGTGGTATGGGTTTCG CCATCAGAACGCACAGGGAACAGGTCAGGCAGCCGACTATAAGTCTACACAAGCCGCCATAGATCAGATCACGGGAAAGCTAAACCGCTT GGTGGAGAAGACAAATACCGAGTTCGAGTCTATAGAGAGTGAGTTTTCCGAGATAGAACATCAAATCGGTAATGTGATCAACTGGACTAAAG ACTCGATAACAGACATCTGGACCTACCAGGCGGAACTGCTCGTCGCCATGGAGAACCAGCACACTATCGATATGGCTGATTCAGAAATGCT GAACCTGTATGAACGCGTACGGAAACAGCTGCGCCAGAATGCAGAGGAGGATGGGAAAGGCTGTTTTGAAATCTACCACGCCTGCGACGA TAGCTGCATGGAAAGCATCCGCAATAACACATATGACCACTCTCAGTATCGCGAGGAAGCTCTTCTGAACCGTCTCAACATCAATCCAGTGA CGTTGTCATCTGGATACAAAGACATCATATTATGGTTTTCCTTCGGCGCAAGTTGCTTCGTGCTCCTGGCAGTTGTCATGGGCCTGGTCTTC TTCTGTCTCAAGAATGGGAATATGAGATGCACGATCTGTATC
2959	ATGTACAAGATTGTTGTGATCATCGCTCTGCTCGGTGCCGTCAAAGGGTTGGATAAGATTTGCCTCGGCCACCATGCAGTAGCCAACGGCA CAATTGTGAAAACTCTCACCAACGAGCAGGAAGAGGTCACCAATGCCACCGAAACAGTCGAGTCAACTGGCATTAATCGATTATGTATGAAA GGAAGAAAGCATAAAGACTTGGGAAACTGTCATCCTATCGGGATGCTCATCGGAACACCAGCATGTGACCTCCATTTAACCGGAATGTGGG ACACATTAATCGAGCGTGAGAACGCAATTGCCTATTGCTACCCTGGCGCAACAGTTAACGTCGAAGCTTTGACACAGAAGATTATGGAGTCA GGAGGCATCAATAAAATCTCGACCGGGTTTACCTACGGCTCCTCCATCAACTCGGCTGGCACAACCAGAGCCTGCATGCGTAACGGAGGC AATAGTTTCTACGCTGAGCTTAAGTGGCTGGTAAGTAAATCAAAAGGTCAAAATTTCCCACAGACTACGAATACTTACAGAAATACAGACACA GCCGAACACCTTATCATGTGGGGGATCCACCACCCTAGCAGTACTCAGGAAAAAAACGATCTGTACGGCACCCAGAGCCTGTCGATCTCTG TGGGAAGTTCAACTTACCGAAATAATTTCGTCCCTGTCGTTGGTGCAAGGCCTCAGGTCAATGGCCAGTCAGGACGCATCGATTTCCATTGG ACTCTGGTGCAACCGGGCGACAACATTACCTTTTCACATAATGGCGGCCTTATCGCCCCCTCCCGAGTGAGTAAGCTGATCGGTCGGGGAT TGGGAATCCAGTCTGATGCACCCATTGATAATAACTGTGAATCTAAATGTTTCTGGAGAGGAGGTTCCATTAATACACGGCTGCCCTTTCAG AACCTGTCCCCTCGAACCGTCGGACAGTGCCCCAAGTACGTGAACAGGAGAAGTCTCATGCTGGCGACAGGGATGAGAAACGTTCCCGAA CTGATTCAGGGACGGGGCTTATTCGGGGCTATTGCAGGCTTTCTGGAAAACGGCTGGGAGGGGATGGTGGACGGCTGGTACGGATTCAG GCATCAGAACGCACAGGGCACCGGCCAAGCAGCCGACTACAAGAGCACTCAGGCAGCCATCGACCAGATTACAGGAAAACTTAATAGGCT TGTGGAGAAAACCAATACCGAATTCGAGTCAATTGAATCCGAGTTTAGCGAGATCGAACATCAAATCGGCAACGTGATTAACTGGACAAAGG ACTCCATCACGGACATATGGACTTACCAGGCCGAACTGTTGGTGGCCATGGAAAATCAACATACAATTGATATGGCAGACAGTGAGATGCT GAACCTGTACGAGAGGGTCAGAAAGCAGCTACGTCAGAACGCTGAAGAGGACGGGAAGGGCTGTTTCGAAATTTATCACGCCTGTGACGA CTCCTGCATGGAATCTATCCGCAATAATACTTATGACCACAGTCAGTATCGCGAAGAGGCCCTCTTAAACAGATTAAACATCAATCCGGTGA CTCTTAGCAGTGGCTACAAGGACATTATTTTATGGTTTTCGTTCGGAGCTAGCTGTTTCGTCCTCCTCGCCGTGGTGATGGGTCTGTTCTTCT TTTGCCTCAAGAATGGAAATATGAGATGTACAATCTGTATA
2960	ATGTATAAGATAGTCGTGATCATTGCGCTTCTGGGAGCAGTAAAGGGTCTTGACAAGATTTGTCTGGGCCACCATGCTGTGGCCAACGGGA CTATTGTTAAGACTCTGACTAATGAACAGGAAGAAGTTACAAACGCTACTGAGACGGTGGAGAGCACTGGCATAAATAGATTGTGTATGAAA GGGAGGAAACACAAAGACTTGGGTAATTGCCCACCCGATTGGGATGCTAATTGGGACTCCCGCTTGTGACCTGCACCTTACTGGCATGTGGG ATACCCTAATAGAACGTGAGAACGCCATTGCCTATTGTTACCCAGGCGCTACTGTCAATGTGGAGGCACTTAGGCAGAAGATCATGGAAAG CGGCGGCATCAACAAGATCTCCACCGGATTTACCTATGGTTCAAGTATCAACAGTGCCGGTACCACCCGCGCTTGTATGCGCAACGGAGGA AACAGTTTTTATGCTGAGCTCAAATGGCTAGTAAGTAAGTCTAAGGGACAAAATTTCCCTCAGACCACCAACACTTATCGTAACACAGACACC GCAGAGCATCTCATTATGTGGGGGATACACCATCCCAGCAGCACACAGGAGAAGAACGATCTGTATGGCACCCAATCGCTCTCCATCTCAG TGGGTTCTTCTACCTATAGAAATAACTTCGTGCCTGTCGTGGGCGCACGGCCACAGGTGAACGGGCAGAGTGGCAGAATCGATTTTCATTG GACGCTGGTCCAGCCCGGCGATAACATTACCTTCTCACACAATGGGGGATTGATCGCCCCTTCTACGGTATCAAAATTGATCGGCAGGGGA CTTGGGATCCAAAGCGATGCCCCTATTGACAACAACTGCGAATCGAAATGCTTCTGGCGGGGCGGCAGCATTAATACACGGCTGCCCTTCC AAAATTTATCCCCAAGAACAGTTGGCCAGTGCCCTAAATACGTCAACCGTCGCAGCCTTATGTTAGCCACAGGAATGCGGAACGTCCCGGA GCTTATTCAGGGCAGGGGCCTGTTCGGCGCAATCGCGGGCTTTCTGGAAAATGGTTGGGAAGGGATGGTTGACGGCTGGTACGGGTTCAG GCACCAGAATGCCCAGGGAACAGGTCAGGCCGCTGACTATAAGTCAACACAAGCAGCCATCGACCAAATTACCGGCAAGCTGAATAGACTT GTGGAAAAAACGAATACAGAATTCGAGAGTATTGAAAGTGAATTTAGTGAGATTGAACACCAAATTGGCAATGTCATTAACTGGACGAAAGAT TCAATCACCGACATCTGGACCTATCAGGCTGAACTCTTGGTCGCAATGGAAAACCAACATACCATCGACATGGCTGATTCTGAGATGCTCAA CTTATACGAGCGAGTTCGAAAGCAACTGCGGCAGAATGCCGAGGAGGATGGGAAGGGGTGTTTCGAAATTTATCATGCCTGCGACGACAG TTGTATGGAGAGCATCCGTAATAATACTTATGACCATAGTCAATACAGAGAGGAGGCTCTTCTGAACAGGCTGAACATCAACCCTGTGACCT TAAGCTCCGGATACAAGGACATTATCTTGTGGTTCAGTTTTGGAGCCTCATGTTTCGTTCTGTTGGCAGTTGTGATGGGCCTCTTCTTCTTCT GCCTGAAAAACGGAAACATGCGTTGCACTATATGCATC

2961	ATGTACAAGATTGTGGTCATCATTGCCCTGCTTGGGGCCGTCAAGGGCCTGGACAAAATTTGCTTGGGGCACCATGCCGTGGTTAATGGCA CCATCGTCAAAACCCTTACGAACGAACAAGAAGAGGTCACCAACGCCACTGAGACAGTTGAAAGTACAGGCCTCAATCGATTGTGCATGAA AGGTAGGAATCACAAAGATCTGGGGAACTGTCACCCCATAGGGATGCTGATTGGCACACCAGCCTGTGATCTGCATTTGACAGGGACCTGG GATACTCTAATTGAGAGAGAGAACGCTATTGCCTACTGCTACCCCGGCGCCACAGTTAACGAAGAGGCTCTGCGGCAGAAAATAATGGAAT CCGGTGGTATTAACAAGATTAGTACAGGGTTTACATACGGATCGTCTATTAATTCAGCTGGTACAACTCGGGCTTGCATGAGGAACGGTGGC AACTCATTCTATGCAGAACTGAAGTGGTTGGTAAGTAAGTCGAAAGGTCAAAATTTTCCCCAGACAACAAATACATATCGCAATACCGACACC GCGGAACATTTGATCATGTGGGGCATCCACCATCCATCAAGTACACAGGAGAAGAATGACCTTTACGGTACTCAGAGTCTGAGCATTTCAGT TGGATCCAGTACATACCAAAATAACTTCGTTCCTGTCGTCGGAGCCCGCCCGCAGGTAAATGGTCAGTCCGGGCGCATTGACTTTCATTGG ACCCTCGTGCAGCCCGGCGATAACATAACCTTCTCACACAATGGAGGACTGATCGCACCCAGCAGAGTGAGCAAACTTATCGGGAGAGGT CTGGGGATCCAAAGTGACGCCCCCATTGATAACAACTGTGAGTCCAAATGTTTCTGGCGAGGGGGTTCAATTAACACCGCTTGCCTTTCC AAAACCTGAGCCCTCGGACGGTGGGCCAGTGTCCAAAGTATGTGAATAAACGGTCTCTGATGTTAGCTACTGGCATGAGAAACGTTCCTGA GTTGATGCAGGGAAGGGGTCTCTTCGGCGCAATCGCTGGCTTTATTGAGAACGGGTGGGAGGGAATGGTGGACGGCTGGTATGGTTTTAG GCATCAGAATGCCCAGGGCACCGGACAGGCGGCTGATTATAAATCAACTCAAGCCGCCATCGACCAAATCACCGGGAAACTAAACCGTCTG ATCGAAAAAACCAATACTGAGTTCGAGTCTATCGAGAGCGAGTTTAGTGAAATCGAACACCAGATTGGAAACGTGATCAATTGGACAAAAGA TAGCATTACCGATATTTGGACCTACCAGGCCGAATTGCTGGTGGCGATGGAAAATCAGCACACAATCGACATGGCAGACTCGGAGATGTTG AATCTGTATGAGAGAGTGAGGAAGCAGCTGCGCCAGAATGCCGAAGAGGATGGAAAGGGATGCTTCGAGATCTACCATGCATGTGACGAC TCCTGTATGGAAAGCATCAGAAATAATACATACGATCACTCTCAATATCGAGAGGAGGCGCTTCTTAATCGTTTAAATATCAATCCCGTCACA CTGTCCTCTGGTTATAAGGACATAATTTTGTGGTTCTCGTTCGGTGCCAGTTGTTTCGTCCTGCTAGCGGTCGTGATGGCCCTGGTGTTTTT TTGCCTTAAAAACGGGAACATGCGGTGCACCATCTGCATC
2962	ATGTACAAGATTGTCGTCATTATCGCGCTGTTGGGGGCCGTCAAGGGACTGGATAAGATTTGTCTGGGCCATCACGCCGTGGCCAACGGCA CAATCGTCAAGACTCTGACCAACGAACAGGAAGAGGTCACCAACGCCACAGAAACCGTGGAGAGTACGGGTATTAACCGCCTCTGCATGAA GGGTCGGAAGCATAAGGACCTCGGTAACTGTCATCCAATCGGCATGTTGATTGGCACACCCGCCTGCGACCTGCACCTGACTGGCATGTG GGACACACTGATTGAAAGGGAAAACGCAATCGCGTACTGTTACCCAGGTGCAACCGTGAACGTTGAAGCACTGAGGCAGAAGATCATGGA GAGCGGGGGCATCAACAAAATTAGCACGGGCTTCACATACGGTAGCAGTATCAACAGCGCTGGCACCACACGGGCCTGCATGCGGAATGG AGGGAACAGTTTCTACGCTGAACTGAAGTGGTTAGTGAGCAAATCTAAGGGTCAGAATTTCCCCCAAACCACTAATACCTACCGCAACACTG ACACCGCCGAGCACTTAATAATGTGGGGCATCCATCATCCTTCAAGCACGCAGGAGAAGAACGACCTGTACGGCACCCAGTCTCTGAGTAT CTCGGTGGGTTCGTCCACCTATAGAAACAATTTTGTGCCAGTGGTGGGGGCCAGGCCACAGGTCAATGGACAATCGGGACGTATCGACTTT CATTGGACCCTCGTGCAGCCTGGTGATAACATTACATTCAGCCATAATGGGGGCCTCATCGCCCCCAGTAGGGTCTCCAAGCTCATTGGGA GAGGACTCGGAATCCAGAGTGATGCACCTATTGACAACAACTGTGAGAGCAAATGTTTTTGGAGAGGCGGGTCTATTAACACTCGATTGCCT TTCCAAAATCTCTCACCTAGGACCGTCGGTCAGTGTCCTAAGTACGTGAATAGGAGAAGCCTCATGCTCGCGACAGGCATGAGAAACGTGC CCGAACTGATCCAGGGCAGGGGACTCTTCGGGGCTATCGCAGGATTCTTAGAGAATGGGTGGGAAGGAATGGTGGATGGATGGTATGGCT TTAGGCATCAGAACGCTCAGGGCACTGGCCAAGCTGCGGACTACAAGAGCACTCAGGCCGCAATAGACCAGATTACCGGCAAGTTGAATC GCCTAGTTGAAAAGACTAATACAGAGTTTGAGAGCATTGAAAGCGAGTTTAGCGAAATTGAACATCAGATCGGCAATGTTATAAACTGGACA AAGGACTCCATAACTGATATATGGACTTACCAGGCAGAGCTACTAGTCGCCATGGAGAACCAGCATACTATCGACATGGCCGACTCAGAGA TGCTGAACCTATACGAAAGAGTTCGGAAGCAGTTAAGACAAAACGCCGAGGAGGACGGAAAGGGTTGCTTCGAAATCTACCACGCATGTGA CGACTCATGCATGGAATCTATCAGGAATAACACATACGATCATTCACAGTATAGGGAGGAGGCTTTGCTGAATCGCCTCAATATCAATCCCG TCACTTTATCTAGCGGCTACAAGGACATTATCTTGTGGTTCAGTTTTGGTGCCTCTTGCTTTGTGCTCCTGGCAGTGGTGATGGGACTCTTTT TTTTCTGTTTGAAGAACGGTAATATGCGTTGTACAATTTGCATC
2963	ATGTACAAGATTGTGGTGATCATCGCACTGCTGGGTGCTGTGAAAGGCCTGGACAAAATATGTTTGGGTCACCACGCCGTGGCAAATGGCA CCATAGTCAAAACCCTCACCAACGAACAAGAGGAAGTGACGAATGCGACTGAAACAGTGGAGTCGACCGGCATCAACCGTCTGTGTATGAA AGGCAGAAAACATAAGGATCTCGGCAATTGCCATCCCATTGGAATGTTAATAGGGACGCCTGCGTGCGATTTGCATCTGACTGGGATGTGG GATACCCTAATAGAGCGGGAGAACGCCATAGCCTATTGCTACCCAGGAGCTACCGTCAATGTGGAAGCCCTACGGCAGAAAATTATGGAAA GCGGTGGTATCAATAAGATTTCCACCGGATTTACATACGGGTCCTCCATCAATTCAGCCGGGACGACTCGTGCTTGCATGCGCAACGGAGG TAACTCGTTCTATGCCGAACTCAAATGGCTGGTGAGCAAGTCTAAGGGACAGAATTTTCCACAAACAACCAACACCTACCGTAATACCGATA CCGCTGAACACCTTATCATGTGGGGCATTCACCACCCTTCCAGCACACAGGAGAAAAACGACTTGTACGGCACACAGTCCCTCAGTATTTC CGTTGGTAGCTCTACTTACAGAAACAATTTTGTTCCCGTCGTCGGTGCCCGTCCTCAAGTCAACGGACAGTCCGGCAGAATTGACTTTCATT GGACGTTAGTGCAGCCTGGGGACAACATTACCTTCAGCCACAATGGGGGTCTGATTGCGCCGTCCCGCGTTAGCAAGCTGATAGGCCGAG GGCTGGGCATTCAGAGCGATGCACCCATCGACAATAATTGCGAATCCAAATGCTTTTGGCGGGGTGGGAGTATTAATACTCGCCTCCCCTT TCAAAATCTCTCACCACGCACTGTTGGGCAGTGTCCCAAATACGTGAATCGGCGTAGCTTAATGCTGGCTACAGGGATGAGGAACGTGCCC GAACTGATTCAAGGCCGTGGTCTTTTTGGAGCCATTGCCGGCTTTTTAGAGAATGGCTGGGAGGGCATGGTTGACGGTTGGTATGGGTTTA GGCACCAGAACGCCCAGGGCACTGGACAGGCAGCCGATTATAAGTCCACACAGGCCGCTATTGATCAGATTACTGGAAAATTGAATAGACT CGTGGAGAAAACTAACACCGAGTTCGAGTCTATTGAGTCAGAATTTTCTGAGATTGAGCACCAGATCGGTAATGTTATCAATTGGACAAAGG ACAGCATCACTGACATTTGGACGTACCAGGCCGAGCTGCTGGTGGCAATGGAGAACCAGCACACCATCGACATGGCGGACTCAGAGATGC TGAACCTATATGAAAGGGTGCGTAAACAGCTTAGGCAAAATGCTGAGGAGGACGCGCAAAGGATGTTTCGAGATTTATCATGCCTGCGATGA CTCCTGCATGGAATCAATTCGCAATAATACTTATGACCATTCACAGTATCGCGAAGAAGCGCTTCTGAATCGGCTAAATATTAACCCCGTAAC TCTGTCTTCTGGATACAAGGATATTATCCTGTGGTTTAGTTTTGGAGCTTCATGTTTCGTATTGTTGGCCGTGGTGATGGGGCTGTTCTTTTT CTGTCTTAAGAACGGCAACATGAGGTGTACAATTTGCATT

2964	ATGTACAAGATCGTCGTGATCATTGCCCTGCTCGGGGCCGTGAAGGGCCTTGACAAGATTTGTCTAGGCCACCATGCAGTTGCTAACGGAA CTATTGTCAAGACATTGACCAACGAACAGGAGGAAGTGACAAATGCCACAGAGACAGTTGAGTCTACTGGGATAAACAGATTGTGTATGAAA GGCAGGAAACACAAAGATCTCGGAAATTGTCATCCTATCGGAATGCTGATAGGAACTCCCGCTTGCGATTTGCATTTGACAGGAATGTGGG ACACACTGATCGAACGAGAAAACGCCATCGCGTACTGCTACCCCGGTGCCACTGTGAACGTGGAGGCCCTTCGGCAAAAGATCATGGAAT CACGAGCAATCAATAAGATCTCTACCGGGTTCACTTATGGCAGCTCTATCAACTCCGCCGGTACTACTAGGGCCTGCATGAGAAATGGGGC AAATAGCTTCTACGCAGAACTGAAGTGGCTGGTGAGCAAGTCGAAGGGGCAGAACTTCCCTCAGACTACGAATACCTATCGCAATACCGAT ACTGCTGAGCATTTGATTATGTGGGGCATCCATCATCCGTCGTCGACGCAAGAGAAAAATGACCTTTATGGCACACAGTCCTTGTCTATTTC TGTAGGGTCTTCTACCTATCGCAATAACTTTGTGCCTGTTGTCGGCGCCAGGCCACAGGTAAATGGACAGTCTGGCAGAATTGATTTCCACT GGACGCTCGTACAGCCTGGGGATAATATTACATTCTCACACAACGGCGGACTTATCGCCCCGAGCCGTGTCTCAAAGCTGATTGGAAGGGG ACTCGGCATTCAGTCCGATGCGCCCATTGACAACAATTGCGAGAGCAAGTGTTTTTGGCGCGGTGGATCCATCAATACCCGCCTTCCTTTTC AGAATTTATCACCACGTACCGTCGGCCAGTGTCCCAAATATGTGAATCGAAGATCTCTGATGCTGGCCACAGGCATGCGGAACGTCCCTGA GCTGATCCAGGGAAGGGGGCTTTTCGGAGCGATTGCCGGGTTTCTGGAAAATGGATGGGAAGGAATGGTGGACGGCTGGTACGGCTTCA GACATCAGAATGCCCAGGGAACCGGGCAGGCTGCGGATTACAAATCTACCCAAGCTGCTATCGACCAGATCACAGGTAAGCTGAATCGGC TCGTGGAGAAAACTAACACTGAATTTGAGTCCATAGAAAGCGAATTTAGTGAAATCGAACACCAAATCGGAAATGTGATTAACTGGACAAAA GATAGCATTACCGATATCTGGACCTACCAAGCTGAATTGCTGGTCGCCATGGAGAACCAACACACTATTGACATGGCAGACTCCGAAATGCT AAATTTGTACGAGCGCGTCCGCAAACAGTTGCGGCAAAACGCAGAAGAAGATGGAAAGGGCTGCTTTGAGATCTACCACGCCTGTGACGAT TCCTGCATGGAATCGATACGAAACAATACTTATGACCATAGCCAGTATCGTGAAGAAGCTCTGCTAAATCGCCTCAACATTAACCCTGTGAC ATTGAGCAGTGGGTACAAGGATATTATTCTCTGGTTTTCTTTCGGTGCCTCGTGCTTTGTTCTGCTGGCGGTGGTGATGGCACTCTTCTTCTT CTGCCTCAAAAATGGGAATATGCGGTGTACAATCTGTATC
2965	ATGTACAAGATCGTCGTTATCATTGCCCTGCTGGGAGCAGTAAAAGGACTGGACAAAATTTGCTTAGGCCATCACGCGGTGGCAAATGGGA CCATCGTAAAGACACTGACCAACGAGCAGGAGGAGGTGACAAACGCTACAGAAACAGTAGAATCGACCGGGATCAACAGGCTCTGTATGAA GGGCCGGAAACATAAGGATCTGGGGAATTGCCATCCTATCGGAATGCTCATTGGTACGCCTGCTTGTGACCTCCACCTCACTGGAATGTGG GACACACTCATCGAGCGCCAGAACGCAATAGCATATTGTTATCCGGGCGCAACTGTCAACGTTGACGGCACTGAGGCAGAAGATTATGGAGA GTGGGGGCATCAACAAGATATCAACAGGCTTTACATATGGATCGAGCATCAATTCAGCAGGGACGACCCGGGCGTGCATGAGGAATGGAG GGAATAGCTTTTACGCCGAATTGAAATGGCTCGTGTCTAAGAGCAAAGGCCAAAATTTTCCACAGACAACTAATACCTATCGCAACACCGAT ACGGCCGAACATCTAATCATGTGGGGAATTCACCACCCAAGTTCGACCCAGGAAAAGAACGATCTCTACGGCACCCAGAGTCTGAGCATCT CGGTGGGATCCTCGACGTACCGGAACAATTTCGTGCCTGTCGTGGGAGCGCGACCGCAGGTTAATGGGCAATCCGGGAGGATCGATTTTC ACTGGACGCTAGTCCAACCCGGGGACAACATCACTTTCAGCCACAACGGTGGACTTATTGCTCCTTCCCGAGTCAGTAAACTCATCGGCCG GGGACTTGGGATTCAAAGCGATGCACCAATAGATAATAACTGTGAGTCTAAGTGCTTCTGGCGGGGAGGTAGTATTAACACACGACTCCCA TTTCAGAACCTGTCCCCAAGGACCGTGGGTCAGTGCCCTAAGTACGTGAACAGACGAAGCCTGATGTTGGCAACCGGAATGAGAAACGTCC CCGAGCTGATCCAGGGGAGGGGACTGTTTGGGGCAATAGCGGGATTTCTGGAGAACGGATGGGAAGGGATGGTTGACGGATGGTACGGG TTTAGGCACCAGAACGCCCAAGGGACTGGACAGGCTGCAGATTATAAAAGTACACAAGCGGCCATCGATCAGATCACCGGTAAACTGAATA GACTTGTCGAGAAAACCAATACAGAGTTCGAGTCGATAGAATCGGAATTCAGCGAGATCGAACACCAGATCGGCAATGTGATCAATTGGACT AAAGACAGCATCACCGATATCTGGACCTATCAAGCTGAGCTCCTAGTAGCTATGGAGAATCAACACACAATAGACATGGCTGACTCGGAAAT GTTGAACCTGTATGAACGTGTCCGCAAGCAGCTGAGGCAGAACGCCGAGGAGGACGGTAAGGGATGCTTCGAGATCTACCATGCTTGTGA CGATTCTTGTATGGAAAGTATCAGAAACAACACCTACGACCATAGCCAGTACAGGGAAGAGGCCCTGCTGAATACACTGAATATCAACCCAG TGACCCTCAGCTCTGGGTATAAAGATATCATTCTGTGGTTCTCATTCGGTGCATCGTGCTTCGTACTGCTGGCCGTTGTCATGGGACTCTTT TTCTTTTGTCTTAAAAATGGCAATATGCGATGCACAATTTGCATT
2966	ATGTATAAAATAGTTGTCATCATCGCCCTCCTAGGTGCAGTTAAGGGGCTCGACAAAATTTGTTTAGGCCACCACGCCGTGGCAAACGGCAC CATCGTCAAGACCCTTACTAATGAACAAGAGGAGGTGACTAATGCTACAGAGACCGTGGAGTCTACCGGCATAAATCGACTATGTATGAAAG GCCGAAAGCACAAAGATTTGGGGAACTGTCATCCTATCGCAATGCTGATTGCAACCCCAGCCTGCGATCTGCACCTCACCGCATGTGGGA TACCCTGATCGAACGTGAAAATGCTATCGCATACTGTTACCCTGGAGCCACAGTCAACGTTGAGGCATTGAGGCAAAAGATCATGGAGAGC GGTGGAATCAATAAGATCTCAACAGGGTTTACCTACGGCAGCTCTATAAACTCCGCTGGCACAACAAGAGCCTGTATGCGCAATGGGGGAA ATTCCTTTTATGCCGAACTGAAATGGCTTGTATCAAAGAGCAAGGGCCAGAACTTCCCCCAAACGACTAATACTTATCGCAACACAGATACA GCTGAGCACCTGATAATGTGGGGTATCCACCATCCATCATCAACGCAAGAAAAGAACGATCTGTACGGGACTCAAAGCCTGTCTATCTCCGT CGGCAGCAGCACATATAGGAATAACTTTGTGCCCGTCGTCGGCGCTAGACCGCAGGTTAACGGACAGAGTGGCCGGATAGACTTCCACTG GACTCTGGTCCAGCCTGGCGATAACATCACCTTCAGCCACAATGGCGGTTTGATAGCACCCTCTCGTGTGTCTAAGCTGATCGGAAGAGGC CTCGGGATTCAGTCCGACGCCCCGATCGACAATAACTGCGAGAGTAAATGTTTTTGGAGAGGAGGATCCATTAACACCCGGCTGCCGTTCC AGAATTTGAGCCCTAGGACAGTGGGGCAGTGTCCCAAATATGTCAACCGCCGAAGTCTGATGCTAGCGACTGGCATGAGGAACGTTCCCG AGCTCATCCAGGGACGCGGGCTGTTTGGCGCCATTGCCGGTTTTCTGGAAAACGGATGGGAAGGTATGGTAGATGGCTGGTATGGATTTC GCCACCAGAACGCGCAGGGCACTGGGCAGGCGGCGGATTACAAGTCCACACAGGCAGCGATTGATCAGATCACCGGCAAACTGAATCGC CTGGTGGAGAAAACAAATACAGAGTTTGAGTCCATCGAAAGCGAGTTTAGCGAAATTGAGCATCAAATTGGCAACGTCATTAATTGGACAAA AGACAGCATTACCGATATCTGGACATACCAGGCGGAGCTACTGGTCGCAATGGAGAACCAGCATACCATCGATATGGCCGACAGTGAAATG CTGAACCTCTATGAGAGAGTGCGAAAGCAGTTAAGACAGAATGCGGAAGAGGATGGGAAAGGTTGTTTTGAGATCTATCACGCCTGTGACG ACTCTTGCATGGAAAGCATTCGAAATAATACCTACGACCATAGTCAGTATCGAGAAGAGGCTCTGTTAAACCGATTAAATATTAACCCGGTAA CTTTGTCGAGTGGGTACAAGGATATTATCCTTTGGTTTTCCTTCGGGGCATCCTGTTTTGTGTTGCTGGCTGTGGTAATGGGACTGTTCTTTT TTTGCTTAAAGAACGGTAATATGAGATGCACAATCTGTATA

2967	ATGTATAAAGTAGTTGTGATCATCGCTTTACTGGGCGCAGTCAAAGGTTTGGACAAGATTTGTCTGGGTCACCATGCCGTTGCCAATGGGAC CATAGTTAAAACCTTAACAAATGAGCAGGAAGAGGTCACAAACGCTACAGAGACTGTGGAGTCCACTGGGATAAACAGACTGTGTATGAAA GGACGCAAGCACAAGGACCTCGGAAATTGTCACCCAATTGGAATGCTGATTGGCACTCCTGCCTGCGATTTACATCTAACCGGCACATGGG ACACCCTGATCGAGCGGGAGAACGCGATCGCCTACTGCTACCCCGGAGCCACCGTCAACGTAGAGGCATTACGGCAGAAGATCATGGAGA GTCGAGGCATAGATAAGATCTCCACCGGATTCACCTATGGTTCTAGCATTAACTCCGCAGGTACAACAAGAGCTTGCATGCGAAATGGAGG GAATAGCTTTTACGCTGAGCTCAAATGGCTTGTGAGCAAAAACAAGGGACAGAATTTTCCCCAGACCACTAATACTTATAGAAACACTGACA CCGCTGAGCACTTAATCATGTGGGGAATCCACCACCCCTCCAGTATCCAAGAAAAGAACGACCTGTATGGAACACAGAGCTTAAGTATCAG CGTGGGGTCATCTACCTACCGGAACAACTTCGTTCCTGTGGTTGGGGCTAGACCACAGGTGAATGGTCAGAGCGGCAGAATCGATTTCCAC TGGACCCTCGTCCAGCCAGGAGACAACATCACCTTCAGCCATAATGGCGGCCTGATTGCCCCTTCTAGGGTTTCCAAATTGATTGGACGCG GCCTCGGGATTCAATCAGACGCTCCCATTGATAATAATTGTGAGAGCAAATGCTTCTGGAGGGGAGGATCCATTAACACCCGATTGCCATTC CAGAATTTATCCCCACGGACCGTGGGCCAGTGCCCTAAATACGTGAATAGGCGATCCCTGATGTTGGCAACAGGCATGAGAAACGTCCCTG AGCTGATCCAAGGGCGCGGACTATTTGGAGCAATAGCCGGCTTCTTAGAGAACGGATGGGAAGGAATGGTGGATGGTTGGTACGGTTTCC GACACCAGAACGCCCAAGGAACTGGACAGGCTGCCGATTACAAATCTACTCAGGCCGCTATAGATCAGATTACAGGAAAACTGAATAGGCT GGTTGAGAAGACGAATACGGAATTTGAATCCATTGAGAGTGAATTCAGTGAGATCGAACACCAGATTGGCAATGTGATTAACTGGACAATGG ACTCCATCACTGATATTTGGACTTACCAGGCCGAGCTGCTTGTAGCGATGGAGAATCAACATACTATCGACATGGCAGACTCCGAGATGCTG AACCTGTATGAAAGAGTGCGCAAGCAGCTGAGACAGAACGCCGAAGAAGACGGCAAAGGCTGTTTTGAAATTTACCATGCCTGCGACGACT CCTGTATGGAGTCCATTCGCAACAATACATACGATCACAGCCAGTACAGGGAGGAAGCGTTGCTGAACAGGTTGAACATAAATCCCGTGAC ACTTAGCTCTGGTTACAAGGATATTATACTCTGGTTTTCCTTTGGAGCTTCTTGCTTCGTGCTGCTCGCAGTCGTGATGGGTCTGGTTTTTTT TTGCCTCAAGAATGGCAACATGCGTTGCACTATATGTATC
2968	ATGTACAAGATCATCGTCATCATCGCTCTCCTCGGTGCGGTGAAGGGTCTGGATAAGATTTGCTTGGGTCACCACGCTGTTGCCAATGGCA CGATTGTAAAGACCCTGACTAACGAGCAAGAAGAAGTTACTAATGCCACAGAAACCGTGGAGTCTACCGGCATTAACAGATTATGCATGAAA GGTCGGAAACATAAAGATTTGGGGAATTGCCACCCTATAGGGATGCTCATCGGGACTCCTGCGTGCGACCTCCACCTTACCGGAACCTGG GACACTCTGATTGAGCGCGAAAATGCAATTGCCTATTGTTATCCCGGCGCAACAGTGAACGTCCAGGCCTTGCGGCAGAAGATAATGGAAT CCGGCGGAATCGACAAGATTAGCACCGGCTTCACCTACGGCTCCTCTATTAATTCTGCTGGAACAACTAGGGCTTGCATGAGAAACGGCGG CAACTCATTCTACGCCGAGCTCAAGTGGCTTGTGTCTAAGTCTAAAGGACAAAATTTCCCACAGACCACCAATACGTATCGCAATACAGACA CTGCGGAACATCTAATCATGTGGGGCATCCATCACCCGAGTAGTACCCAGGAAAAGAACGACCTCTACGGCACTCAATCACTGTCCATCAG CGTTGGCTCTAGCACATACCGGAACAATTTCGTCCCTGTGGTAGGAGCTAGACCCCAGGTTAACGGACAGTCAGGCCGTATCGACTTTCAT TGGACCTTAGTACAACCAGGGGATAACATTACCTTTTCCCATAACGGGGGACTGATCGCCCCTAGCCGCGTCAGTAAGCTCATCGGACGTG GACTCGGAATCCAGTCAGACGCCCCCATCGATAACAACTGCGAGTCAAAATGCTTTTGGCGAGGAGGATCTATCAACACTAGGCTGCCCTT TCAAAATCTGTCCCCTCGCACAGTGGGTCAGTGTCCTAAATACGTGAACCGCAGATCACTCATGCTCGCCACCGGAATGAGGAATGTGCCC GAACTCATCCAAGGGAGGGGCCTCTTCGGTGCCATCGCGGGATTCCTCGAAAATGGATGGGAGGGGATGGTGGATGGATGGTACGGCTTT CGGCACCAAAACGCTCAGGGCACCGGGCAGGCCGCCGATTATAAATCTACGCAGGCAGCTATAGACCAAATAACCGGCAAATTAAATCGG CTGGTAGAAAAGACCAACACCGAATTCGAGAGCATAGAGAGCGAATTTTCAGAAATAGAGCACCAGATTGGGAACGTAATTAACTGGACAAA GGACTCTATAACCGATATCTGGACATACCAGGCGGAACTCTTGGTGGCCATGGAGAATCAGCACACCATTGACATGGCCGATAGTGAGATG CTGAATCTCTACGAGAGGGTGCGAAAACAGCTAAGGCAAAATGCCGAAGAGGACGGCAAGGGTTGCTTTGAGATCTACCACGCCTGCGAC GACTCTTGTATCGAGTCCATTCGAAACAACACCTACGACCATTCTCAATACAGGGAGGAAGCTTTGCTCAATCGGCTCAACATCAATCCACT CACGCTGTCTTCCGGTTATAAGGACATCATCCTCTGGTTTAGTTTCGGAGCTTCTTGTTTTGTACTCTTAGCCGTGGTGATGGGCCTGTTCTT CTTCTGCCTAAAAAACGGCAACATGCGGTGCACAATCTGTATT
2969	ATGTACAAAATAGTGGTCATCATAGCCCTTCTGGGTGCCGTGAAGGGGCTAGACAAAATCTGTTTGGGACATCACGCAGTAGCGAACGGCA CCATCGTCAAAACGCTGACGAACGAGCAAGAAGAGGTTACCAATGCTACAGAGACTGTGGAGTCTACTGGCATCAATAGGCTCTGCATGAA AGGACCGAAACACAAGGACCTGGGCAATTGTCACCCTATCGGCATGTTGATCGGAACTCCCGCCTGTGATCTGCACCTTACAGGGATGTGG GATACGCTGATCGAGCGGGAAAATGCCATAGCTTACTGCTACCCCGGAGCGACCGTTAACGTGGAGGCCTTAAGACAGAAGATCATGGAG TCCGGCGGGATCAATAAGATTTCCACTGGCTTTACGTACGGTTCCAGTATCAACTCCGCAGGGACGACGCGCGCGTGCATGAGAAACGGC GGCAACAGCTTTTACGCTGAACTCAAGTGGCTGGTTAGCAAGTCAAAGGGCCAGAATTTCCCTCAGACCACTAACACATACAGGAATACTGA CACCGCCGAGCACCTCATCATGTGGGGAATTCACCACCCTTCCTCCACCCAGGAAAAGAACGATCTATACGGGACCCAGTCTTTGTCCATA TCTGTGGGCAGTTCAACATACCGGAACAACTTTGTGCCTGTGGTGGGTGCTAGGCCACAGGTCAACGGGCAATCCGGTAGGATTGATTTTC ACTGGACTCTGGTCCAGCCCGGGGACAATATAACTTTTTCGCATAATGGCGGGCTGATCGCTCCCTCCCGGGTGTCAAAGCTGATTGGACG GGGCCTTGGTATACAGTCTGACGCGCCCATTGATAACAACTGCGAGTCTAAGTGTTTTTGGCGGGGAGGCAGCATCAATACTCGCCTCCCT TTTCAAAACCTCTCCCCTAGAACAGTGGGGCAATGCCCAAAATACGTCAATAGGAGGTCCCTCATGCTAGCAACCGGTATGCGCAACGTTC CCGAACTGATCCAGGGCCGAGGGCTGTTCGGAGCTATCGCAGGGTTTCTCGAAAACGGCTGGGAAGGAATGGTCGATGGCTGGTATGGGT TCAGACATCAGAATGCCCAGGGCACAGGACAGGCAGCAGATTACAAAAGTACTCAGGCCGCCATCGACCAGATCACCGGCAAACTTAATAG GCTGGTGGAAAAAACCAACACCGAGTTTGAGAGTATAGAGAGCGAGTTTTCAGAGATCGAGCATCAGATTGGGAATGTTATCAACTGGACC AAGGATTCCATTACTGATATTTGGACTTACCAGGCGGAACTGCTGGTTGCCATGGAGAATCAGCACACGATCGATATGGCCGACTCCGAAAT GTTGAATTTATATGAGCGGGTGAGAAAGCAGCTCAGGCAGAACGCAGAAGAGGACGGAAACGGGTGCTTCGAGATCTACCATGCTTGTGA CGATTCCTGTATGGAGTCAATTAGAAACAATACCTACGACCACTCCCAGTATAGAGAGGAGGCTCTGCTAAACCGCCTTAATATCAACCCGG TTACCCTGTCAAGCGGATACAAGGATATTATATTGTGGTTTTCTTTTGGCGCATCTTGCTTCGTGTTACTCGCTGTGGTGATGGGTTTATTCT TCTTTTGTCTGAAGAATGGGAACATGCGGTGTACCATTTGCATA

2970	ATGTACAAGGTGGTGGTCATCATTGCCCTGCTGGGGGCAGTCAGGGGCTTAGATAAGATTTGTCTTGGGCATCACGCCGTGGCCAACGGA ACGACCGTGAAAACCCTCACAAATGAGCAGGAGGAAGTTACGAATGCCACCGAGACTGTGGAGTCTACTTCTCTGAACAAACTGTGCATGA AAGGCCGGCGTTATAAAGATCTGGGAAATTGTCATCCTATCGGCATGCTGATTGGTACTCCTGTATGTGACCTGCACCTCACGGGCACTTG GGATACACTGATCGAGAGGGAAAATGCCACAGCTTACTGCTACCCCGGAGTCACAATTAACGAGGAGGCACTCAGACAGAAAATCATGGAG AGCGGTGCTATCAGCAAAATGCGCACCGGCTTTACTTATGGCCCAAGCATCAACTCTGCCGGTACAACACGAAGTTGCATGAGGAACGGCG GCAATAGCTTCTACGCTGAACTCAAGTGGTTAGTCTCCGGTACGAAGGGGCAGAATTTTCCGCAGACTACAAATACATACCGAAATACAGAT ACAGCTGAGCACCTGATCATCTGGGGCATCCACCACCCTAGTTCAACCCAGGAGAAAAACGATTTGTATGGAACGCAATCTCTCTCCATTTC CGTAGGGTCCTCCACCTATCAAAACAACTTTGTCCCGGTTATTGGGGCTAGACCTCAGGTCAATGGCCAGAGCGGTAGGATTGAGTTTCATT GGACTCTCGTCAGACCAGGGGACAACATCACGTTTTCCCATAATGGCGGACTCATCGCTCCAGACAGGGTTTCCAAGCTCATCGGAAAGGG GATTGGGATTCAGTCAGGGGCAGTAATAGATAAGGACTGTGAGTCTAAATGTTTTTGGCGGGGAGGTTCCATAATTACCGAACTGCCATTTC AGAATCTCTCCCCTAGAACAGTGGGGCAGTGTCCCAAATACGTTAAAAAAAGGAGTTTGCTGCTGGCAACAGGCATGCGGAACGTCCCTGA GGTAGTACAGGGGCGGGGATTATTCGGAGCAATAGCCGGGTTTATCGAAAACGGGTGGGAAGGTATGGTGGATGGATGGTACGGCTTCCG TCACCAAAATGCTCAGGGGATCGGACAGGCCGCCGACTACAAATCTACTCAGACTGCCATAGATCAGATTACCGGTAAACTCAACAGGCTC ATCGAAAAAACAAATACTGAGTTTGAAAGCATTGAGTCTGAGTTTTCTGAGATTGAGCATCAGATTGGCAACGTCATCAACTGGACCAAGGA TTCCATTACAGATATCTGGACGTACCAAGCCGAGCTGCTCGTGGCAATGGAGAACCAGCATACAATAGACATGGCTGATTCTGAAATGCTGA ATCTGTACGAACGTGTGAGAAAGCAACTGAGGCAGAACGCTGAGGAGGACGGCAAGGGCTGTTTCGAGATCTATCATACCTGCGATAACTC ATGTATGGAATCCATCCGGAATAATACTTATGATCACAGTCAGTATCGCGAGGAAGCCCTGCTGAATAGATTAAATATCAATCCTGTGAAACT GAGTTCTGGGTACAAAGATATCATCCTGTGGTTCAGCTTTGGGGCCAGCTGTTTCGTACTGCTGGCCGTAATCATGGGGCTAGGATTCTTTT GCCTGAAGAACGGCAATATGCGCTGCACGATTTGCATT
2971	ATGTACAAAATTGTGGTGATAATTGCCCTTTTAGGCGCCGTGAAGGGCCTCGATAAAATTTGCCTTGGCCATCATGCAGTCGCTAACGGGAC TATCGTAAAGACCCTGACCAACGAACAGGAAGAAGTTACCAATGCCACTGAAACTGTGGAGAGTACAGGCATTAACAGGCTGTGCATGAAG GGACGCAAGCACAAGGATTTAGGGAATTGCCATCCGATCGGAATGCTGATCGGGACCCCCGCGTGCGACCTGCACCTGACAGGAATGTGG GACACCCTTATTGAGAGAGAAAATGCTATCGCTTATTGCTACCCTGGTGCCACTGTCAACGTGGAGGCCTTACGTCAGAAGATTATGGAATC CGGAGGCATAAATAAGATTAGTACCGGGTTCACATATGGCTCTTCCATAAACTCTGCGGGCACCACTAGAGCATGTATGAGGAACGGTGGG AATAGCTTCTATGCTGAGCTCAAGTGGTTGGTCAGCAAGTCCAAAGGTCAGAACTTTCCGCAGACAACTAATACCTACCGAAATACAGACAC CGCTGAGCACCTCATTATGTGGGGCATTCACCACCCGAGCTCAACTCAAGAGAAGAATGACTTGTACGGAACACAATCTCTGTCCATCTCTG TGGGCAGCAGTACCTATCGGAATAATTTCGTCCCAGTAGTGGGCGCAGGACCTCAGGTCAATGGTCAGTCCGGTCGCATCGACTTCCATTG GACCCTGGTGCAACCAGGCGACAACATCACCTTTTCTCACAACGGCGGGCTGATCGCACCCTCGCGTGTGTCTAAGCTAATCGGCCGTGG CCTGGGGATCCAGTCTGATGCACCCATTGACAATAACTGCGAGTCTAAATGTTTCTGGAGAGGCGGATCCATTAACACGCGGCTGCCATTT CAGAACCTGAGTCCAAGAACGGTAGGCCAGTGTCCGAAGTACGTCAATCGGAGGTCATTGATGCTGGCAACCGGCATGAGAAATGTCCCG GAGCTGATACAGGGAAGGGGACTATTTGGAGCTATCGCTGGCTTCCTAGAAAATGGATGGGAGGGTATGGTCGATGGATGGTATGGGTTT CGCCACCAAAACGCACAGGGGACTGGGCAGGCCGCAGATTATAAGAGCACTCAAGCTGCAATCGACCAGATCACTGGCAAACTGAATCGG CTTGTGGAGAAAACCAACACAGAATTCGAGTCAATTGAGTCCGAGTTTAGCGAAATTGAACACCAGATCGGAAATGTTATTAACTGGACTAA GGATTCCATCACAGATATCTGGACCTACCAGGCAGAGCTATTGGTAGCCATGGAGAACCAACATACAATCGACATGGCGGATTCAGAAATG CTGAACCTGTATGAACGCGTTAGGAAGCAGCTCAGGCAGAACGCCGAGGAGGATGGGAAAGGTTGCTTCGAGATATATCATGCCTGCGAC GACAGCTGTATGGAGTCCATACGAAATAACACGTATGACCACAGCCAGTACCGAGAAGAGGCTCTTCTTAATCGGCTCAATATAAACCCAGT GACACTGTCATCCGGATACAAGGACATCATACTATGGTTCAGCTTTGGTGCCTCTTGTTTCGTGCTACTTGCTGTGGTGATGGGCCTTTTTTT TTTCTGCCTCAAGAACGGGAACATGAGGTGTACGATATGCATC
2972	ATGTACAAGATAGTCGTGATCATTGCCCTCCTTGGTGCAGTAAAGGGATTAGACAAGATTTGCTTGGGACACCACGCGGTTGCTAACGGAA CAATAGTTAAGACACTCACTAATGAGCAAGAGGAAGTCACAAATGCTACCGAAACCGTAGAGTCTACGGGCATCAACAGGCTTTGTATGAAA GGCAGGAAGCACAAGGACCTTGGGAATTGTCACCCTATTGGTATGCTTATTGGCACCCCAGCCTGCGATCTACATCTGACTGGGATGTGGG ACACCCTCATCGAACGAGAGAATGCCATAGCCTATTGCTACCCTGGCGCAACTGTGAATGTGGAAGCTTTGAGGCAGAAGATAATGGAATC CGGCGGAATCAACAAGATATCAACAGGCTTTACTTACGGCTCGAGTATCAATTCAGCCGGCACAACTCGGGCCTGCATGCGGAATGGGGG AAACTCTTTCTACGCAGAACTGAAATGGCTGGTATCCAAGTCTAAGGGCCAGAATTTCCCGCAGACCACCAACACCTATCGCAACACTGACA CAGCAGAACATTTGATAATGTGGGGTATCCACCACCCGTCTAGTACCCAGGAGAAGAACGACTTGTATGGCACGCAGTCATTATCTATCAGC GTGGGCTCCAGCACGTACCGGAACAACTTTGTCCCCGTGGTTGGCGCACGACCACAAGTGAACGGACAATCCGGCCGGATCGATTTTCAT TGGACACTCGTTCAGCCCGGCGACAATATCACCTTTTCTCACAATGGGGGCCTGATCGCGCCAAGCCGGGTTTCCAAGCTGATCGGCAGA GGCCTCGGCATACAGAGCGATGCCCCGATCGATAACAATTGTGAGTCGAAATGTTTTTGGAGAGGCGGCTCTATTAACACGCGACTGCCCT TTCAGAATCTGAGCCCGCGGACAGTGGGACAATGCCCAAAATACGTAAACAGGAGGAGTCTGATGCTCGCGACTGGCATGAGGAACGTTC CTGAGTTGATCCAGGGCCGGGGCCTATTTGGCGCCATCGCAGGGTTCCTGGAGAATGGTTGGGAGGGTATGGTAGATGGATGGTACGGGT TCCGCCACCAGAATGCCCAGGGAACAGGCCAGGCTGCGGACTACAAATCCACTCAGGCCGCCATCGATCAAATTACCGGAAAGCTTAACA GACTCGTCGAGAAAACGAATACAGAGTTTGAGTCAATTGAGTCCGAGTTCAGTGAAATCGAGCACCAAATAGGAAACGTGATCAATTGGACA AAGGACAGTATCACCGATATCTGGACTTACCAGGCTGAACTGCTCGTAGCTATGGAGAATCAGCACACCATAGATATGGCAGACTCCGAGA TGTTGAATTTATACGAGAGAGTCCGGAAACAGCTGCGCCAGAACGCCGAAGAGGATGGGAAGGGCTGCTTCGAGATATACCACGCCTGTG ACGACTCATGCATGGAGTCTATCCGAAACAATACCTACGATCATTCCCAGTATAGGGAAGAGGCCTTGCTGAACCGGTTGAATATTAATCCT GTCACATTGTCTAGCGGCTACAAAGACATCATCTTATGGTTTTCGTTTGGTGCTTCATGTTTCGTGCTTCTGGCTGTGGTGATGGGACTCTTT TTCTTTTGCCTTAAGAATGGCAATATGAGATGCACCATTTGTATT

2973	ATGTATAAGATCGTGGTGATCATTGCCCTACTGGGTGCCGTGAAAGGGCTGGACAAAATTTGCCTCGGCCACCATGCTGTCGCCAACGGCA CCATCGTCAAAACGCTTACGAATGAGAAGGAGGAAGTGACCAATGCTACCGAGACTGTGGAAAGTACAGGCCTCAACAGGCTCTGTATGAA GGGCAGGAAACACAAAGACTTGGGTAACTGCCATCCTATAGGAATGCTCATTGGCAGCCCGGCTTGTGACCTGCACCTTACTGGGACCTGG GATACTCTGATCGAGAGGGAGAATGCCATTGCGTATTGTTATCCAGGAGCCACCGTAAACGGCGAGGCACTAAGGCAAAAGATTATGGAAA GCGGGGGTATTGACAAAATTTCAACAGGGTTCACATATGAGTCTAGCATCAACTCAGCCGGCACAACGAGAGCATGCATGCGGAACGGGG GCAATTCCTTTTACGCTGAGTTGAAGTGGCTGGTGTCCAAGAGCAAAGGGCAGAATTTCCCACAGACTACCAACACCTATAGAAATACGGAT ACAGCAGAGCACCTGATTATGTGGGGTATCCACCACCCAAGCTCCACGCAAGAAAAGAATGATCTGTATGGAACTCAAAGCCTCTCTATTAG TGTGGGGTCTAGCACTTACCGGAACAACTTCGTCCCTGTGGTGGGCGCTCGCCCTCAGGTTAATGGCCAGTCCGGAAGAATTGACTTTCAT TGGACTCTCGTCCAGCCAGGGGATAATATAACATTCTCTCACAATGGAGGACTGATTGCACCTTCTCGGGTGAGCAAGCTGATAGGCCGCG GGCTGGGGATCCAGAGCGATGCTCCAATAGATAATAATTGCGAATCAAAGTGCTTTTGGAGGGGGGGATCTATTAATACGCGGCTTCCATT CCAGAACCTCAGTCCTCGGACTGTTGGACAATGCCCCAAATATGTGAACAAGCGCAGCCTTATGCTCGCTACCGGGATGAGGAATGTCCCT GAATTAATGCAGGGACGCGGCCTATTCGGAGCAATTGCCGGATTTCTCGAGAACGGTTGGGAAGGCATGGTTGACGGGTGGTATGGTTTC AGACACCAGAACGCCCAGGGGACAGGGCAGGCCGCCGATTATAAATCCACCCAGGCCGCCATCGACCAGATTACCGGTAAATTAAACAGG CTAGTTGAAAAAACCAATACCGAGTTTGAATCCATAGAATCTGAATTCTCCGAAATTGAACACCAGATCGGTAATGTTATCAACTGGACCAAA GACAGCATCACCGACATATGGACTTATCAGGCTGAGCTGCTTGTCGCTATGGAGAACCAGCACACCATCGACATGGCCGATTCTGAAATGC TGAACCTGTACGAAAGGGTGCGGAAGCAGCTACGGCAGAACGCTGAGGAGGACGGCAAGGGGTGCTTTGAAATCTACCACGCCTGCGAT GATTCCTGTATGGAGTCCATCCGTAACAACACGTACGACCACTCACAGTATAGAGAGGAAGCCCTCCTAAACCGCCTGAATATCAACCCCGT AACCCTCTCATCTGGTTATAAAGACATCATCCTCTGGTTTAGTTTCGGCGCCTCATGTTTCGTACTGCTCGCAGTGGTGATGGGCCTGGTAT TCTTCTGTCTAAAGAACGGGAATATGAGATGCACCATCTGCATT
2974	ATGTATAAGATCGTGGTGATCATTGCACTCCTTGGGGCAGTCAAAGGCTTGGACAAAATTTGTTTGGGACACCACGCAGTCGCTAACGGTAC AATAGTCAAAACACTGACAAATGAGCAGGAAGAAGTGACTAACGCCACAGAAACAGTGGAGTCCACAGGCATTAATCGCCTTTGTATGAAG GGTCGCAAGCATAAGGATCTGGGGAACTGTCACCCTATAGGTATGCTGATAGGGACTCCAGCCTGTGATTTACACCTGACAGGGATGTGGG ACACGCTCATCGAGAGGGAAAACGCCATCGCCTACTGCTATCCTGGGGCTACAGTCAACGTCGAGGCCCTACGACAAAAAATTATGGAGAG CGGCGGGATCAATAAGATTTCCACTGGCTTTACTTATGGCTCATCTATAAACTCCGCAGGCACAACACGCGCCTGTATGCGCAACGGCGGA AACTCATTTTACGCGGAACTGAAGTGGCTGGTTTCTAAAAGCAAGGGCCAGAATTTCCCACAGACTACAAATACCTATCGCAACACTGACAC CGCAGAACACTTAATAATGTGGGGGATTCACCACCCCTCGAGCACCCAAGAGAAGAACGACCTTTATGGAACGCAGTCTCTCAGCATCAGC GTGGGATCATCCACATACCGGAACAACTTTGTGCCTGTGGTGGGAGCTCGACCACAAGTCAACGGCCAGTCTGGTAGAATTGATTTTCACT GGACCCTTGTGCAACCTGGGGATAATATAACTTTCTCGCACAATGGTGGTTTAATAGCCCCATCTCGCGTCAGCAAACTGATCGGTAGAGG CCTCGGAATACAGTCCGATGCACCTATCGATAACAATTGTGAATCCAAGTGCTTCTGGAGGGGGGGATCCATTAATACTCGGCTGCCTTTCC AAAATTTGTCACCCCGCACTGTGGGCCAGTGTCCAAAGTACGTGAACAGAAGGAGCCTGATGTTGGCAACCGGGATGCGCAACGTTCCCG AACTCATCCAGGGGAGAGGTTTGTTTGGAGCTATCGCGGGCTTTCTCGAAAACGGCTGGGAGGGGATGGTGGATGGATGGTACGGCTTTC GTCACCAGAACGCCCAGGGAACGGGGCAAGCAGCCGATTACAAGAGCACCCAGGCCGCTATCGACCAGATTACGGGGAAGCTGAACCGC CTAGTAGAGAAGACCAACACAGAATTCGAATCTATTGAATCAGAGTTTAGCGAAATAGAGCATCAGATCGGAAACGTGATCAATTGGACTAA AGATTCTATCACAGATATCTGGACTTACCAGGCTGAACTGCTTGTAGCAATGGAGAATCAACACACAATAGACATGGCCGATTCTGAGATGC TGAACCTGTACGAGCGTGTACGCAAACAGCTGAGACAGAACGCCGAAGAGGATGGTAAAGGATGTTTCGAAATCTACCATGCTTGCGACGA CTCTTGCATGGAAAGTATCCGGAACAATACATACGATCACTCCCAATATCGCGAGGAGGCGCTGCTGAACAGACTCAACATTAACCCCGTG ACTCTATCCTCAGGGTACAAGGATATCATATTATGGTTCTCTTTCGGCGCCTCCTGTTTCGTCCTGCTGGCCGTCGTAATGGGCCTTTTCTTT TTCTGTTTGAAGAATGGAAATATGCGGTGCACCATCTGTATA
2975	ATGTACAAAATCGTGGTAATTATTGCTCTTCTTGGGGCTGTTAAAGGGTTGGACAAGATTTGTCTCGGGCATCACGCCGTGGCCAATGGAAC GATTGTGAAGACACTGACCAACGAACAGGAGGAGGTAACTAACGCCACGGAAACTGTCGAGTCCACTGGCATCAATCGGCTCTGTATGAAG GGCCGCAAACACAAGGACTTGGGGAACTGCCACCCAATCGGCATGCTTATCGGCACACCCGCTTGTGACTTACATCTGACTGCGATGTGG GACACGCTTATAGAACGGGAAAACGCCATTGCCTACTGCTACCCAGGTGCAACTGTTAACGTCGAAGCTCTGCGCCAAAAGATCATGGAAA GCGGCGGTATCAACAAAATTAGTACGGGCTTTACCTATGGCTCAAGCATTAACTCAGCTGGCACCACCCGTGCCTGTATGAGGAATGGAGG CAACTCCTTTTACGCTGAGCTCAAGTGGCTCGTGTCCAAGTCCAAGGGACAGAATTTCCCCCAGACTACCAATACCTATAGAAATACTGACA CCGCTGAACACTTGATTATGTGGGGTATCCATCACCCTAGCTCCACTCAGGAAAAAAACGATCTATACGGCACCCAGTCGTTGTCCATCTCA GTCGGCAGTTCCACTTATCGCAACAACTTTGTGCCCGTGGTCGGAGCACGTCCTCAGGTTAATGGCCAGTCTGGCCGTATCGATTTTCACT GGACACTGGTACAGCCTGGCGATAATATTACTTTTAGCCACAACGGGGGCCTGATCGCCCCGAGTCGTGTGTCCAAGCTTATCGGACGAG GTCTTGGGATTCAATCTGATGCCCCAATTGACAATAACTGTGAAAGCAAATGCTTTTGGCGCGGGGGATCCATTAACACGCGGCTTCCTTTC CAGAATCTATCCCCTCGTACTGTCGGGCAATGCCCTAAGTACGTCAACCGGAGGTCCCTGATGCTGGCCACTGGCATGCGTAACGTCCCAG AGCTCATTCAGGGCAGAGGTTTATTCGGAGCAATTGCAGGGTTCCTGGAGAACGGCTGGGAGGGAATGGTCGATGGCTGGTATGGTTTCC GCCATCAAAACGCTCAAGGCACAGGCCAGGCGGCCGATTACAAATCGACTCAGGCGGCCATTGACCAAATCACAGGCAAACTGAATAGACT GGTGGAAAAAACAAACACCGAGTTTGAGTCTATTGAATCTGAGTTTAGCGAAATCGAGCACCAAATCGGCAATGTTATTAACTGGACTAAGG ATTCAATTACGGATATTTGGACATACCAGGCGGAGTTATTAGTTGCCATGGAGAATCAGCACACTATTGATATGGCTGATAGTGAGATGTTAA ATTTGTACGAGACGGTTAGGAAGCAGTTGAGGCAGAACGCGGAGGAAGACCGCGAAAGGCTGCTTCGAGATTTACCACGCCTGCGACGATT CCTGCATGGAGAGCATCCGGAATAATACCTACGACCATTCACAGTATCGAGAAGAAGCTCTATTAAACCGTTTGAACATCAACCCAGTGACC TTAAGTAGCGGATACAAAGACATTATTCTATGGTTCTCCTTCGGCGCCTCCTGTTTCGTGTTGCTCGCCGTTGTAATGGGATTGTTTTTCTTC TGTCTGAAGAATGGAAACATGAGATGCACAATTTGTATC

2976	ATGTACAAGATTGTTGTGATCATCGCCCTTCTGGGGGCCGTGAAGGGACTTGACAAAATATGCCTCGGTCATCACGCTGTTGTGAATGGCA CAATAGTTAAAACGCTGACCAACGAGCAAGAAGAGGTGACCAACGCAACCGAGACAGTGGAGAGTACAGGTCTGAACCGACTGTGCATGA AGGGCCGAAACCACAAGGACCTCGGCAATTGTCACCCCATCGGAATGTTAATTGGTACCCCCGCCTGCGATCTGCATCTGACCGGTACTTG GGATACCTTAATTGAAAGGGAAAACGCCATCGCGTACTGTTATCCCGGAGCAACCGTGAACGAGGAAGCCCTCAGACAGAAGATCATGGAG TCGGCAGCCATCAACAAGATCTCCACAGGCTTCACGTACGGCTCCTCTATTAATTCTCCCGGAACAACTCGGGCCTGTATGCGCAACGGAG GCAACTCCTTCTACGCAGAACTGAAATGGCTTGTCAGCAAGAGCAAGGGTCAGAATTTCCCCCAAACTACCAATACTTATAGGAATACGGAC ACCGCAGAACACCTGATAATGTGGGGCATTCATCATCCCTCTTCAACTCAGGAGAAAAATGATTTGTATGGGACTCAATCCTTATCAATCTCC GTGGGTAGCAGCACCTATCAGAATAACTTTGTTCCAGTCGTAGGGGCCCGGCCTCAGGTGAACGGGCAGTCAGGCAGAATTGATTTCCACT GGACCCTTGTGCAACCCGGAGACAACATTACCTTTAGTCACAATGGCGGTCTCATCGCCCCTTCCAGAGTGTCTAAACTGATCGGTCGGGG CCTGGGCATCCAGAGCGACGCACCAATCGACAATAATTGCGAGAGTAAGTGCTTCTGGAGAGGTGGTAGTATCAATACTAGGCTCCCATTC CAAAATTTATCGCCTAGGACCGTCGGACAGTGTCCCAAATATGTAAATAAAAGGTCCTTAATGCTTGCGACGGGAATGCGTAACGTTCCTGA ACTGATGCAGGGCAGAGGCCTATTTGGCGCAATAGCCGGTTTTATCGAGAACGGCTGGGAGGGGATGGTGGACGGTTGGTATGGATTTAG GCACCAAAATGCCCAGGGAACTGGGCAGGCGGCGGATTACAAGAGCACACAAGCTGCTATCGACCAGATCACGGGCAAACTGAATCGATT GATTGAAAAGACCAACACTGAGTTTGAATCAATTGAGAGTGAGTTTTCAGAGATTGAACACCAGATCGGGAACGTGATAAACTGGACTAAAG ATTCCATCACTGACATCTGGACATATCAGGCGGAGCTCCTGGTGGCCATGGAGAACCAACACACCATCGACATGGCAGACTCTGAAATGTT AAATCTGTACGAGAGAGTCAGGAAACAGCTGCGCCAGAATGCCGAGGAAGATGGCAAGGGTTGCTTCGAGATCTACCACGCATGTGACGA CAGCTGCATGGAGTCAATTCGCAACAACACCTACGACCACTCCCAATACAGGGAGGAGGCACTGCTTAACAGGCTGAACATTAATCCGGTC ACCCTCTCTAGCGGCTATAAGGATATTATTCTGTGGTTCAGTTTTGGCGCTTCTTGTTTTGTCCTGCTGGCAGTGGTGATGGGCCTGGTCTT CTTTTGCCTAAAAAATGGGAATATGAGGTGCACCATCTGTATT
2977	ATGTATAAAATTGTGGTCATCATTGCTCTGCTGGGTGCAGTAAAGGGGCTGGACAAAATTTGTCTGGGACACCATGCCGTTGCCAACGGAA CAATCGTGAAGACGCTTACTAACGAACAGGAAGAAGTTACCAACGCTACCGAAACCGTCGAATCCACGGGTATCAACAGGCTGTGTATGAA AGGGAGGAAGCACAAAGATTTGGGGAACTGCCACCCAATCGGCATGTTGATTGGCACTCCCGCCTGTGATCTGCATTTAACTGGTATGTGG GATACCCTGATCGAGCGAGAAAATGCTATTGCATACTGTTATCCAGGCGCGACTGTGAATGTCGAAGCTCTGCGACAGAAAATCATGGAATC CGGCGGTATCAATAAAATTTCGACAGGCTTCACATATGGGAGTTCTATTAACTCTGCCGGGACCACCAGAGCCTGCATGCGCAACGGAGGA AATAGCTTCTACGCTGAGTTAAAGTGGCTAGTCTCTAAATCTAAAGGACAAAACTTCCCACAGACCACTAACACCTACCGGAATACCGATAC CGCGGAACACTTAATTATGTGGGGGATCCACCACCCAAGCTCTACGCAGGAAAAGAACGATCTCTACGGGACTCAGTCTTTATCAATTTCCG TGGGCTCCTCTACATACCGCAATAATTTCGTGCCCGTGGTTGGCGCTAGGCCTCAGGTAAACGGGCAGAGTGGAAGGATAGATTTTCACTG GACCCTGGTTCAGCCGGGCGACAACATCACTTTCAGTCACAATGGCGGCCTTATCGCTCCCAGCCGGGTGTCTAAATTGATCGGCAGGGG ACTGGGTATCCAGTCGGACGCTCCAATTGACAACAATTGTGAGTCAAAGTGCTTTTGGCGAGGGGGGAGCATCAACACTCGCCTCCCTTTC CAAAACCTGTCTCCTCGGACGGTCGGACAATGCCCTAAGTACGTAAACAGGCGGTCGCTGATGCTGGCCACTGGGATGAGAAACGTGCCA GAACTTATTCAGGGCCGTGGCTTGTTCGGCGCAATCGCTGGGTTTCTTGAAAATGGCTGGGAAGGGATGGTGGATGGTTGGTATGGGTTCC GACATCAGAACGCTCAGGGCACTGGGCAGGCCGCTGATTATAAGTCCACCCAGGCAGCTATAGACCAAATAACAGGGAAACTAAATCGCCT GGTCGAAAAGACAAATACAGAGTTCGAATCTATCGAGAGCGAATTCTCAGAAATCGAGCACCAGATTGGGAATGTGATTAATTGGACCAAGG ATAGTATCACAGATATCTGGACATACCAGGCAGAACTGTTGGTTGCCATGGAAAACCAGCACACCATTGATATGGCAGATTCAGAGATGCTT AACCTGTATGAGAGAGTGCGAAAGCAGCTGCGCCAGAACGCAGAAGAGGATGGAAAGGGGTGCTTCGAAATTTACCACGCTTGCGATGAT AGCTGTATGGAGTCAATAAGAAACAACACCTATCACCACAGCCAGTACAGGCAGGAGGCACTACTTAATAGGCTGAATATCAATCCAGTTAC TTTGTCCAGCGGCTATAAGGACATTATACTGTGGTTTAGCTTTGGGGCGTCCTGCTTTGTGCTCCTGGCTGTAGTGATGGGGCTATTTTTCT TCTGTTTAAAGAACGGTAATATGCGGTGCACAATCTGTATC
2978	ATGTACAAAATAGTGCTGGTGCTCGCCCTCCTCGGTGCTGTGCATGGTCTTGACAAGATATGTCTCGGACATCACGCCGTCCCGAACGGAA CTATTGTGAAAACCCTAACCAACGAGAAGGAAGAGGTGACTAACGCGACCGAAACTGTGGAAAGCAAGAGTCTGGATAAACTGTGTATGAA GAATAGGAATTACAAGGACCTGGGCAATTGTCATCCTATTGGGATGGTGGTCGGTACCCCCGCTTGTGATCTTCATCTGACCGGAACCTGG GATACCCTGATAGAGCGCGACAATTCAATTGCGTATTGCTACCCCGGTGCCACAGTTAGCGAGGAAGCCTTGCGCCAGAAGATCATGGAGA GTGGCGGCATAGATAAAATCAGCACCGGATTCACGTACGGGAGTAGTATCAATTCAGCCGGCACAACAAAAGCCTGCATGAGAAATGGCGG AAACTCCTTTTATTCTGAACTGAAATGGCTCGTGTCCAAGAACAAGGGCCAGAACTTTCCTCAGACGACCAATACTTATCGCAACACAGATTC CGTCGAACATTTGATCATTTGGGGCATTCACCATCCTAGCTCTACCCAGGAGAAAAACGATCTGTACGGCACCCAAAGCCTGTCCATTAGCG TCGGGTCCAGTACATACCAGAACAACTTCGTGCCAGTCGTCGGGGCGCGGCCGCAGGTCAACGGCCAGTCAGGGAGAATAGACTTTCATT GGACCATGGTGCAGCCCGGGGACAACATTACTTTTTCCCATAATGGAGGCCTGATCGCACCAAATCGGGTCTCTAAGCTGAAAGGGAGAG GTCTCGGGATCCAGTCCGGAGCATCTGTGGACAACGACTGTGAATCCAAATGCTTCTGGAAGGGTGGGTCTATCAATACCAAGCTCCCCTT TCAGAATCTCTCGCCTAGGACCGTGGGACAGTGCCCAAAATACGTGAACAAAAAAAGTCTCCTCCTGGCTACTGGTATGCGGAATGTCCCA GAGGTTGCACAGGGGCGAGGCCTGTTCGGTGCCATAGCGGGTTTTATTGAGAACGGTTGGGAGGGAATGGTGGATGGCTGGTATGGGTTT CGGCACCAAAATGCACAGGGCACAGGTCAGGCCGCCGATTATAAGAGCACACAGGCAGCCATTGATCAGATTACAGGGAAGCTGAATCGA CTGATAGAAAAAACAAATACCGAGTTTGAGTCCATTGAAAGCGAATTTAGCGAGATTGAGCACCAGATAGGGAATGTCATCAATTGGACCAA AGATTCAATCACTGATATTTGGACTTACCAGGCTGAGCTGCTGGTTGCAATGGAAAATCAGCACACAATTGACATGGCCGACAGCGAGATGC TGAACCTCTACGAGAGAGTGCGTAAACAGCTGCGCCAGAACGCAGAAGAGGATGGGAAGGGATGTTTCGAAATTTATCACAAGTGTGATGA CAACTGTATGGAGAGCATTCGAAACAACACGTACGACCATACTCAGTATCGGGAGGAAGCCCTGTTGAATCGCCTCAATATTAATCCCGTGA AACTCTCATCCGGTTATAAAGACGTCATACTGTGGTTTTCCTTCGGGGCCAGCTGCTTCGTACTGCTGGCCGTCATTATGGGCCTTGTTTTTT TCTGTCTGAAAAATGGAAACATGAGGTGTACCATCTGTATT

2979	ATGTATAAAATCGTGGTGATCATCGCACTGCTGGGTGCGGTTAAGGGGCTCGACAAGATTTGCCTGGGACACCATGCAGTGGCAAATGGCA CTATTGTCAAAACACTTACTAACGAGCAGGAGGAAGTGACAAACGCCACGGAGACCGTTGAGAGCACGGGCATCAATCGGCTGTGCATGAA AGGCAGGAAGCATAAGGACCTCGGCAACTGCCATCCTATTGGTATGCTGATTGGCACGCCCGCTTGCGATCTGCATCTGACCGGAATGTG GGATACGCTTATAGAGAGAGAGAACGCGATTGCCTACTGCTACCCCGGCGCTACAGTCAATGTCGAAGCTCTGCGCCAAAAAATTATGGAA TCTGGCGGGATTAACAAGATCTCCACTGGCTTCACGTATGGATCTTCAATAAACTCAGCCGGCACCACACGGGCCTGCATGCGCAATGGGG GAAACTCATTCTATGCTGAACTGAAATGGCTTGTCTCAAAATCTAAGGGGCAGAATTTTCCTCAAACTACCAACACCTACAGGAATACGGATA CCGCTGAGCACCTGATCATGTGGGGTATCCACCACCCTTCGTCCACCCAGGAAAAAAATGACCTATATGGTACCCAGTCACTGAGTATCAG CGTAGGCAGTTCTACATATCGTAACAACTTTGTCCCAGTAGTCGGGGCCCGCCCTCAAGTGAACGGCCAATCTGGACGTATTGATTTCCACT GGACACTGGTGCAACCCGGAGACAACATCACTTTCTCCCACAACGGAGGGCTGATAGCCCCCTCCAGGGTCTCCAAGTTGATCGGACGTG GACTTGGGATACAGTCTGATGCCCCAATTGATAATAACTGTGAGTCTAAATGCTTCTGGAGGGGGGGCAGTATCAATACACGCCTGCCTTTC CAGAACCTGTCGCCACGGACAGTTGGACAGTGTCCAAAATATGTGAACCGCAGGTCGCTGATGTTGGCAACCGGAATGAGGAATGTGCCC GAGCTGATTCAGGGTCGGGGATTATTTGGCGCCATCGCAGGGTTCCTGGAAAATGGATGGGAAGGCATGGTGGATGGATGGTACGGCTTT AGACACCAAAACGCCCAGGGAACAGGGCAAGCAGCAGACTACAAATCCACGCAAGCTGCTATAGACCAGATTACTGGTAAGCTCAACCGTC TCGTTGAGAAAACAAATACGGAATTTGAGAGTATCGAGTCAGAGTTTTCTGAAATCGAACATCAAATAGGGAATGTAATTAATTGGACAAAAG ACTCAATCACTGACATATGGACCTATCAGGCTGAACTGCTAGTTGCAATGGAGAATCAACATACCATCGACATGGCTGACAGCGAGATGCTG AATTTGTACGAACGGGTCAGAAAACAACTGCGGCAGAATGCTGAGGAAGATGGAAAGGGGTGTTTCGAAATATATCACGCTTGTGATGACT CTTGCATGGAATCCATCCGGAATAATACCTACGATCACTCGCAGTATCGAGAGGAGGCCCTGCTGAATCGCTTAAACATTAATCCAGTGACA CTGTCTTCCGGGTATAAAGATATCATTCTGTGGTTTTCCTTTGGCGCATCCTGCTTCGTATTACTAGCCGTTGTCATGGGATTGTTTTTCTTTT GTCTGAAAAACGGGAACATGCGGTGTACCATTTGTATT
2980	ATGTACAAGATCGTGGTCATCATCGCTCTGTTGGGCGCCGTGAAAGGGCTTGATAAGATCTGTCTTGGGCACCACGCCGTGGCCAACGGG ACCATTGTGAAAACCCTCACGAACGAGCAGGAGGAAGTGACAAACGCAACTGAGACCGTTGAGTCAACCGGTATTAATCGCTTGTGCATGA AAGGACGAAAGCACAAGGACTTGGGCAACTGCCATCCAATTGGGATGCTGATCGGTACCCCAGCATGTGATCTGCATTTAACAGGCATGTG GGATACCCTTATTGAGCGAGAAAATGCCATCGCGTATTGCTATCCTGGAGCCACTGTGAATGTGGAGGCACTGCGTCAGAAGATTATGGAA TCTGGCGGGATCAATAAAATAAGCACCGGCTTTACATACGGCAGCTCGATTAATTCCGCTGGCACCACTAGAGCTTGCATGCGCAATGGCG GCAACTCATTCTATGCCGAGCTGAAGTGGTTGGTGAGTAAGAGCAAAGGGCAGAACTTTCCTCAGACCACCAACACCTATCGAAATACGGA CACAGCTGAACACCTTATAATGTGGGGGATCCACCACCCGTCATCTACCCAGGAAAAGAACGACTTATACGGAACACAGTCTCTGTCCATCT CGGTTGGATCCTCTACTTACCGAAATAATTTTGTGCCCGTCGTGGGGGCCAGACCCCAGGTAAACGGACAGTCTGGCAGGATCGATTTCCA TTGGACTCTGGTCCAGCCCGGAGATAATATCACCTTTAGTCACAACGGTGGACTGATCGCTCCCAGTAGAGTGTCCAAGCTGATAGGCAGA GGGCTTGGAATCCAAAGCGACGCCCCAATCGATAATAACTGCGAAAGCAAGTGTTTCTGGCGGGGCGGATCCATCAACACCAGGCTTCCCT TCCAGAACTTGTCCCCAAGGACTGTTGGACAGTGCCCAAAGTATGTTAATCGACGGAGTCTGATGCTCGCTACTGGCATGAGAAACGTACC CGAGCTGATACAAGGGCGCGGGCTGTTCGGCGCAATAGCCGGATTCCTGGAGAACGGATGGGAGGGTATGGTGGACGGCTGGTATGGCT TTCGGCATCAGAATGCCCAGGGTACTGGTCAGGCAGCCGATTATAAGTCAACCCAGGCTGCCATCGATCAGATTACAGGGAAACTGAATCG GCTGGTGGAAAAAACCAATACCGAATTTGAGTCCATTGAATCTGAATTCAGCGAGATCGAGCACCAGATCGGTAATGTCATAAATTGGACTA AGGATTCCATCACAGATATCTGGACATACCAAGCTGAGCTGCTTGTGGCCATGGAAAACCAGCATACAATAGACATGGCCGACAGTGAGAT GCTAAATCTGTATGAGAGAGTGAGAAAACAACTAAGACAGAATGCTGAGGAGGACGGCAAAGGTTGTTTTGAGATCTACCACGCTTGCGAC GACTCGTGTATGGAATCCATACGCAACAATACTTACGACCACTCTCAGTACCGCGAGGAAGCTTTACTTAACCGCCTGAATATTAACCCTGT TACCTTATCCAGTGGGTACAAAGACATAATATTGTGGTTCAGTTTTGGCGCGTCGTGTTTTGTTCTCCTGGCCGTTGTGATGGGCCTGTTTTT TTTCTGCCTGAAGAACGGGAATATGCGCTGTACTATCTGCATA
2981	ATGTACAAGATAATCGTCATTATCGCACTGTTGGGCGCCGTTAAAGGACTGGATAAAATCTGCCTGGGCCATCACGCAGTCGCTAACGGTA CAATCGTGAAAACTCTGACCAATGAGCAGGAGGAAGTCACCAACGCAACCGAAACCGTAGAGTCCACCGGCATCAACCGGCTCTGTATGAA GGGCCGCAAGCATAAAGACCTGGGTAACTGCCACCCCATAGGCATGCTTATTGGCACTCCGGCATGCGACTTGCACCTGACGGGGACCTG GGACACATTGATTGAACGGGAAAATGCTATTGCTTATTGCTATCCTGGAGCCACAGTGAATGTGGAGGCTCTCCGCCAAAAGATCATGGAAT CTGGGGGGATAGATAAGATTAGTACTGGGTTTACGTATGGGAGTTCTATCAACAGCGCCGGGACCACAAGAGCCTGTATGAGAAACGGAG GAAACAGCTTTTACGCAGAGTTGAAATGGCTTGTTAGTAAATCTAAGGGGCAGAATTTCCCTCAGACCACGAACACTTACAGGAATACGGAT ACTGCGGAGCACCTTATTATGTGGGGCATCCATCATCCCTCCTCTACTCAGGAGAAAAACGACCTCTATGGCACTCAATCCTTGTCCATCTC CGTCGGTTCCTCAACCTATAGGAACAACTTTGTGCCAGTTGTAGGCGCAAGGCCACAGGTTAACGGGCAGAGCGGTCGCATCGACTTCCAC TGGACCCTGGTGCAGCCAGGCGATAACATCACCTTTTCACACAACGGCGGGCTGATCGCACCTTCCCGAGTGTCTAAGCTGATAGGGAGG GGCCTCGGCATCCAGTCTGACGCACCAATTGACAATAATTGTGAGTCCAAGTGTTTTTGGCGCGGAGGTTCAATTAACACGCGCCTGCCCT TTCAGAATCTGTCTCCCCGTACTGTGGGCCAGTGTCCAAAGTATGTTAATAGGCGGTCCCTCATGCTGGCTACGGGAATGAGAAACGTGCC TGAGCTGATACAGGGCCGAGGACTATTCGGGGCGATCGCCGGCTTTTTAGAAAACGGGTGGGAGGGCATGGTCGACGGATGGTATGGCTT CAGACACCAGAATGCGCAGGGGACTGGCCAGGCTGCTGACTATAAGAGTACCCAGGCCGCTATAGACCAGATCACGGGTAAGCTGAATCG TCTGGTTGAAAAGACTAATACTGAATTTGAAAGCATCGAAAGCGAGTTTAGCGAAATTGAGCATCAGATCGGGAACGTCATTAATTGGACTA AAGACTCCATTACTGATATCTGGACGTACCAGGCTGAGTTGTTAGTGGCCATGGAAAATCAGCATACCATTGATATGGCAGATAGCGAAATG CTGAATCTGTACCAGAGAGTTCGTAAGCAGCTGCGCCAGAACGCTGAAGAGGACGGAAAGGGATGCTTTGAGATCTACCACGCCTGCGAC GACAGCTGTATGGAATCGATCCGGAACAACACCTATGACCACTCCCAGTATAGGGAAGAAGCCCTGTTAAACCGCCTGAATATTAATCCCGT CACTCTGTCATCCGGCTATAAGGACATAATTCTTTGGTTTAGCTTCGGCGCTAGTTGCTTTGTGCTACTGGCTGTCGTGATGGGCCTCTTTTT TTTTTGCCTTAAAAATGGCAATATGAGGTGTACAATCTGTATC

2982	ATGTATAAGATCGTCGTAATCATCGCCCTGCTGGGCGCTGTTAAGGGGCTGGATAAGATCTGCCTAGGCCACCATGCCGTAGCCAATGGTA CAATCGTTAAGACATTAACCAATGAGCAGGAGGAGGTTACAAACGCTACCGAAACCGTTGAGAGCACCGGGATTAACCGACTCTGTATGAA GGGTAGGAAACATAAGGACCTCGGCAATTGCCATCCTATTGGCATGCTCATAGGGACCCCAGCGTGTGACTTGCACCTCACTGGAATGTGG GACACTCTGATCGAGAGGGAAAATGCCATAGCTTATTGTTACCCAGGAGCCACAGTGAATGTGGAAGCCCTGAGACAGAAGATCATGGAGT CCGGCGGAATCAACAAGATCAGCACTGGTTTCACCTATGGGAGTAGCATAAACAGCGCTGGTACTACCCGAGCATGCATCCGCAATGGAG GCAATAGCTTTTACGCTGAACTGAAATGGCTGGTCTCAAAATCGAAAGGGCAGAATTTTCCTCAGACGACAAATACCTATCGCAATACGGAC ACCGCAGAACACCTGATTATGTGGGGTATCCACCATCCTAGCTCCACACAGGAGAAAAATGATCTGTACGGAACCCAGTCCCTGTCTATCTC CGTGGGATCCTCCACTTACCGGAACAATTTTGTTCCTGTCGTGGGGGCTCGGCCGCAAGTGAACGGTCAGAGCGGACGAATTGATTTCCAC TGGACTCTGGTGCAGCCTGGGGACAATATCACGTTCTCCCATAACGGAGGACTGATCGCCCCCAGCCGCGTGAGCAAGTTGATTGGGCGC GGCCTTGGAATTCAGTCAGATGCGCCAATAGATAATAATTGTGAATCTAAGTGTTTCTGGAGGGGAGGCTCCATCAATACCCGGCTCCCTTT CCAAAATTTGTCCCCGAGAACAGTGGGTCAGTGCCCAAAATATGTGAACCGCCGATCTCTGATGCTGGCCACCGGCATGAGGAACGTGCCT GAGCTGATTCAGGGTAGAGGCCTGTTTGGAGCCATCGCCGGCTTTTTAGAGAACGGCTGGGAAGGAATGGTCGACGGATGGTATGGCTTC AGGCACCAGAACGCCCAAGGGACTGGCCAGGCGGCTGACTACAAGTCCACTCAAGCGGCTATTGACCAAATTACAGGAAAGCTGAATAGA CTCGTTGAAAAAACTAATACAGAGTTTGAAAGTATAGAATCTGAGTTCTCAGAAATAGAGCATCAGATTGGAAACGTGATCAACTGGACTAAA GACTCTATCACCGATATTTGGACGTATCAGGCGGAGTTGCTCGTCGCGATGGAGAACCAACACACAATAGATATGGCTGATTCTGAGATGCT GAACTTGTACGAACGGGTGCGTAAACAACTGCGCCAAAATGCTGAGGAGGACGGGAAGGGGTGTTTTGAGATCTACCATGCTTGCGACGA TAGCTGCATGGAGTCAATTCGAAATAACACGTATGACCACAGCCAGTACAGGGAAGAGGCCCTGCTGAACAGGCTGAATATAAATCCCGTT ACCCTCTCCTCTCGATACAAAGACATCATCCTCTGGTTTTCATTCGGAGCTAGTTGCTTCGTACTACTAGCTGTTGTAATGGGCCTGTTCTTC TTCTGTCTCAAGAATGGTAACATGCGGTGTACGATCTGCATC
2983	ATGTATAAGATCGTGGTGATCATTGCCCTCTTGGGCGCCGTGAAGGGTCTGGATAAAATATGTCTTGGTCACCACGCTGTGGCTAATGGTAC CATCGTGAAGACGCTTACCAACGAACAAGAGGAAGTGACAAATGCTACCGAGACCGTGGAGTCTACCGGGATCAACCGACTGTGCATGAAA GGACGAAAGCACAAAGACCTCGGCAATTGCCACCCTATTGGGATGTTGATTGGCACTCCAGCTTGCGACCTTCACCTGACCGGAATGTGGG ACACGCTCATCGAGCGCGACAACGCCATTGCCTACTGCTACCCCCGGGCCACCGTTAACGTGGAAGCACTTAGGCAAAAAATTATGGAATC CGGTGGAATAAACAAGATTAGCACTGGTTTCACTTATGGGAGTAGTATTAACTCTGCCGGGACAACCAGGGCTTGTATGAGAAATGGCGGC AATTCATTTTACGCAGAGCTCAAGTGGCTGGTTTCAAAAAGTAAAGGGCAGAATTTCCCGCAAACCACTAATACCTACAGGAACACAGACAC GGCCGAGCACCTGATCATGTGGGGTATTCACCACCCTAGCAGCACGCAAGAGAAAAATGACCTTTATGGAACCCAGAGCCTCAGTATTTCA GTGGGAAGCAGCACGTACCGTAATAATTTTGTTCCGGTAGTCGGCGCCCGACCCCAGGTGAACGGGCAGTCGGGACGAATCGACTTTCAT TGGACGCTTGTGCAGCCTGGCGACAACATCACCTTTAGCCACAATGGCGGCCTGATTGCTCCCAGCCGCGTCAGCAAGCTGATCGGGCGC GGACTGGGCATTCAGTCTGATGCCCCTATTGATAACAACTGCGAGAGCAAGTGTTTTTGGCGGGGAGGCTCCATAAATACTCGCTTACCTTT CCAGAACCTCAGTCCAAGGACCGTAGGGCAGTGTCCTAAGTATGTTAACAGAAGGAGTCTCATGTTAGCCACAGGAATGCGCAATGTACCA GAACTCATTCAGGGCAGGGGACTCTTTGGTGCAATTGCTGGGTTTCTGGAGAACGGTTGGGAAGGTATGGTCGACGGCTGGTACGGCTTT CGACATCAAAATGCACAGGGCACAGGCCAGGCGGCTGATTATAAATCCACTCAGGCCGCCATCGACCAAATCACAGGAAAGCTCAACCGTC TGGTCGAGAAGACCAATACTGAGTTCGAAAGCATCGAGTCAGAGTTTAGCGAGATTGAGCATCAGATTGGCAATGTCATCAACTGGACTAAA GATAGCATAACCGATATTTGGACGTACCAAGCAGAGCTACTTGTTGCTATGGAGAATCAGCATACCATTGATATGGCCGATAGCGAGATGCT GAATCTGTATGAGAGAGTGAGAAAGCAGTTGAGACAAAACGCGGAGGAGGATGGGAAGGGCTGTTTCGAGATCTACCACGCCTGCGACGA CTCCTGTATGGAGTCTATACGCAACAACACATACGACCACAGCCAGTATAGGCAGGAAGCCTTCCTGAATAGGCTGAATATCAATCCTGTAA CGCTGTCGAGTGGGTACAAAGACATCATTTTATGGTTCAGTTTTGGAGCCTCTTGTTTCGTTTTGCTGGCCGTCGTCATGGGACTATTTTTTT TTTGTTTAAAGAACGGGAACATGAGATGCACTATTTGCATC
2984	ATGTACAAGATTGTGTTGGTCCTAGCCCTCCTCGGCGCTGTGCATGGGCTGGATAAGATCTGCCTTGGGCACCACGCCGTGCCTAATGGCA CTATCGTGAAAACACTCACAAACGAAAAAGAGGAAGTAACAAATGCCACCGAGACTGTGGAAAGCAAGAGCCTTGATAAACTCTGTATGAAA AACAGAAATTATAAAGATTTAGGCAACTGTCATCCCATTGGTATGGTTGTCGCAACGCCCGCTTGTGATCTGCATTTAACGGGTACATGGGA TACCCTCATCGAGCGAGATAACTCAATCGCCTACTGCTACCCGGGCGCCACTGTGTCAGAGGAAGCCTTGAGACAGAAGATTATGGAATCC GGCGGTATTGACAAGATTTCTACCGGGTTTACGTATGGGTCTTCCATAAACTCCGCAGGCACCACAAAGGCATGCATGAGAAATGGTGGTA ATTCTTTCTACTCAGAGCTTAAATGGCTGGTGTCTAAGAACAAAGGCCAGAACTTTCCCCAGACGACCAACACATACCGTAACACGGACAGC GTCGAGCACCTGATCATCTGGGGCATTCACCATCCTTCCTCAACTCAGGAGAAAAACGATCTCTATGGTACCCAAAGCCTCTCTATTTCAGT CGGATCAAGCACATACCAGAACAACTTTGTGCCCGTCGTTGGAGCCAGGCCACAGGTGAACGGCCAGAGTGGTCGAATTGATTTTCACTGG ACAATGGTGCAACCTGGGGATAATATCACATTCAGCCATAATGGTGGTCTGATCGCCCCAAATAGAGTTTCAAAGCTTAAGGGCAGGGGTCT GGGGATTCAAAGCGGTGCTTCCGTCGATAATGACTGTGAGTCAAAATGCTTTTGGAAGGGGGGCAGCATCAATACAAAGCTACCGTTTCAG AACCTCAGTCCCAGGACAGTTGGGCAGTGTCCTAAGTATGTGAATAAAAAATCCTTACTGTTGGCCACTGGAATGAGAAACGTTCCAGAGGT AGCTCAGGGCCGTGGTCTCTTTGGGGCCATCGCCGGCTTTATTGAAAACGGGTGGGAAGGAATGGTTGATGGGTGGTATGGGTTCAGGCA TCAGAACGCCCAGGGCACTGGACAAGCAGCTGACTATAAGAGCACACAGGCTGCTATTGACCAGATTACTGGAAAACTGAATAGGTTGATA GAGAAGACCAACACTGAGTTCGAGTCTATCGAGTCAGAGTTTTCAGAAATCGAGCATCAGATCGGTAACGTAATCAACTGGACCAAGGACTC AATAACCGACATCTGGACATATCAAGCCGAGTTGCTGGTAGCTATGGAGAATCAACATACTATCGACATGGCTGATTCCGAAATGCTCAACT TGTATGAGCGCGTGCGTAAGCAGTTGAGACAGAATGCTGAGGAAGATGGCAAGGGCTGCTTCGAGATTTACCACAAGTGTGACGATAATTG TATGGAGAGCATTCGAAACAACACTTACGACCACACCCAATACCGCGAGGAAGCCCTGTTGAATCGGCTCAACATCAATCCCGTAAAGTTGT CCAGTGGTTATAAAGACGTCATCTTATGGTTTAGTTTTGGAGCTTCTTGTTTTGTGCTCCTGGCAGTCATTATGGGACTGGTATTCTTCTGCC TGAAAAATGGGAATATGAGATGCACTATCTGTATC

2985	ATGTATAAAATTGTCGTGATTATTGCTCTGCTCGGTGCCGTTAAGGGACTCGATAAGATCTGCTTAGGACATCACGCTGTAGTGAACGGCAC AATCGTGAAAACACTCACGAACGAGCAGGAAGAGGTCACAAACGCCACAGAAACTGTCGAGAGCACCGGTCTAAACCGACTGTGTATGAAA GGAAGGAATCATAAGGACCTAGGCAATTGTCACCCAATCGGAATGCTTATCGGCACCCCTGCATGCGATTTGCACCTGACAGGTACATGGG ATACGCTGATCGAGCGCGAGAATGCCATCGCCTATTGTTACCCCGGTGCAACCGTAAACGAGGAGGCCCTGAGGCAAAAGATAATGGAGT CACGCCGCATAAATAAGATTAGCACCGGTTTTACCTATGGGAGCAGTATAAATAGTGCCGGTACTACGCGCGCATGTATGCGCAATGGGGG CAACTCTTTTTACGCCGAACTCAAGTGGCTAGTGTCAAAATCGAAGGGCCAGAACTTTCCACAGACCACTAACACTTATAGGAACACGGACA CAGCCGAACATTTAATAATGTGGGGTATCCATCACCCAAGTTCCACTCAGGAGAAAAACGATTTATACGGAACTCAGTCGCTGTCAATCAGC GTAGGGAGCTCAACATACCAGAACAACTTTGTTCCAGTGGTGGGCGCACGTCCCCAGGTGAACGGTCAAAGCGGCCGGATAGATTTTCACT GGACCCTGGTGCAGCCTGGAGATAATATTACTTTTAGTCACAACGGTGGCCTAATCGCACCCAGTCGTGTATCGAAGCTTATCGGACGGGG CCTCGGGATACAGTCTGATGCTCCCATAGACAACAACTGTGAGTCTAAGTGCTTTTGGCGGGGCGGGTCTATTAACACCAGGCTCCCATTT CAGAACCTCAGTCCTCGGACCGTGGGTCAGTGTCCGAAGTATGTTAATAAGCGTTCTCTCATGCTCGCCACTGGAATGAGGAACGTGCCAG AGCTAATGCAGGGAAGAGGACTGTTCGGGGCCATAGCCGGCTTCATTGAAAATGGCTGGGAGGGTATGGTTGACGGATGGTACGGCTTCC GACACCAGAATGCTCAAGGCACAGGCCAGGCTGCAGATTATAAATCAACACAGGCTGCTATTGACCAAATCACTGGCAAGCTCAATCGATTA ATTGAAAAAACAAATACCGAGTTTGAAAGCATCGAGAGTGAATTTTCCGAGATCGAACACCAAATCGGAAACGTGATCAACTGGACCAAGGA CTCTATCACCGATATATGGACCTATCAGGCAGAACTTTTAGTTGCCATGGAGAACCAGCATACCATTGATATGGCCGACAGCGAAATGCTGA ACTTGTATGAGCGGGTCCGGAAACAATTACGCCAGAATGCCGAGGAAGACGGAAAAGGGTGCTTTGAGATCTATCATGCTTGCGACGACTC TTGTATGGAGTCAATCAGAAATAATACTTATGACCATAGTCAGTATAGGGAAGAGGCGCTGCTCAACAGACTAAATATCAACCCCGTTACGC TGAGCTCCGGTTATAAGGACATCATTCTTTGGTTTTCCTTTGGCGCAAGCTGCTTTGTTCTGCTAGCTGTAGTGATGGGACTCGTCTTTTTCT GCCTGAAGAACGGAAACATGCGCTGTACAATCTGTATT
2986	ATGTATAAGATCGTTGTGATCATCGCCCTGCTGGGCGCCGTGAAAGGGCTGGACAAGATTTGTCTGGGTCACCACGCCGTCGCCAACGGG ACAATTGTTAAAACACTGACTAACGAGAAGGAGGAGGTGACTAACGCAACCGAGACAGTTGAGTCCACAGGATTGAATCGGCTTTGTATGA AAGGCCGGAAGCACAAAGACCTGGGAAACTGCCACCCTATAGGGATGCTGATCGGAAGTCCTGCTTGTGATCTACACCTGACAGGTACTTG GGACACACTGATCGAGCGGGAGAATGCCATTGCTTACTGCTACCCAGGAGCCACCGTGAATGGTGAAGCTCTGAGACAGAAGATTATGGAA TCCGGAGGCATTGATAAAATCTCTACAGGATTTACATACGAGTCCTCAATCAATTCAGCTGGCACAACTCGAGCATGTATGCGCAATGGCGG GAATAGTTTTTATGCCGAGCTCAAGTGGCTCGTCTCTAAAAGCAAGGGGCAGAACTTTCCCCAGACCACGAACACCTATAGAAATACCGATA CAGCCGAGCATCTTATTATGTGGGGCATACATCATCCCTCCTCTACACAAGAGAAGAATGATCTTTACGGAACGCAGTCCCTCTCGATCAGC GTGGGAAGCTCCACATATAGGAACAATTTTGTTCCCGTGGTTGGCGCTCGTCCTCAGGTGAACGGACAGAGTGGCCGAATCGACTTTCACT GGACCCTTGTTCAGCCCGGCGACAACATAACGTTTTCGCATAATGGCGGCCTCATTGCACCTTCCCGTGTCAGCAAACTGATTGGCCGCGG GCTGGGCATCCAATCTGACGCTCCCATCGATAACAACTGCGAATCAAAATGCTTCTGGAGGGGAGGGTCTATTAATACTCGGCTGCCATTC CAAAACCTTTCCCCCCGGACAGTTGGCCAGTGTCCGAAATATGTGAACAAACGGAGTTTGATGCTCGCTACCGGGATGCGGAACGTGCCTG AATTGATGCAGGGAAGAGGTCTTTTCGGCGCCATTGCCGGCTTCTTGGAAAATGGTTGGGAAGGGATGGTCGACGGATGGTATGGATTTCG GCATCAGAACGCTCAAGGCACTGGGCAGGCAGCTGATTATAAATCAACCCAGGCTGCCATTGACCAGATAACTGGCAAACTCAACAGGCTG GTGGAGAAGACCAACACCGAATTCGAGAGTATTGAGTCTGAATTCAGCGAGATCGAGCACCAGATAGGGAACGTTATTAACTGGACCAAGG ATTCTATTACGGACATTTGGACATACCAAGCTGAGCTCTTAGTGGCCATGGAGAATCAGCATACCATTGACATGGCCGATTCTGAGATGCTT AATCTCTATGAGAGGGTGAGAAAGCAGCTGCGGCAGAATGCCGAAGAAGATGGCAAGGGGTGCTTTGAGATTTACCACGCTTGCGATGATT CATGTATGGAATCTATTCGCAATAACACCTACGATCATTCTCAGTATCGGGACGAAGCCCTGCTGAATCGCCTGAACATTAATCCCGTCACA CTTTCAAGTGGTTACAAGGACATAATTCTATGGTTCAGTTTTGGGGCTAGTTGTTTTGTGCTCCTAGCCGTTGTGATGGGTCTTGTCTTCTTT TGTTTGAAGAACGGAAATATGCGGTGTACAATCTGTATA
2987	ATGTACAAGATCGTGGTGATCATTGCGCTCCTGGGCGCCGTCAAGGGACTGGACAAGATCTGCCTGGGCCATCACGCTGTGGCAAATGGG ACAATCGTGAAAACGCTGACTAACGAGCAAGAGGAGGTGACAAATGCGACGGAGACAGTGGAATCGACTGGGATTAACCGACTGTGTATG AAAGGAAGAAAACATAAAGACCTCGGAAATTGCCATCCCATTGGGATGCTGATCGGAACTCCAGCCTGTGACCTGCACCTGACAGGAATGT GGGATACCCTGATTGAGCGGGAGAATGCTATCGCATATTGTTACCCCGGGGCTACCGTAAACGTCGAAGCCCTTAGGCAGAAGATAATGGA AAGCGGTGGCATCAATAAGATTTCGACAGGCTTCACTTATGGAAGTAGTATTAATAGCGCAGGAACCACTCGAGCCTGTATGAGAAACGGC GGTAACAGCTTTTATGCTGAGCTCAAGTGGCTTGTCTCCAAGTCTAAGGGACAGAATTTCCCACAGACTACAAACACTTACAGGAATACCGA TACAGCTGAGCATCTAATTATGTGGGGAATACATCACCCTTCTTCCACCCAGGAAAAAAACGACCTCTACGGCACTCAATCTCTCAGCATAT CGGTGGGGAGTTCGACTTACCGAAACAACTTTGTCCCGGTTGTCGGTGCCAGGCCCCAGGTGAACGGACAAAGCGGTCGGATCGATTTTC ATTGGACTTTCGTCCAGCCTGGGGACAATATCACATTCTCACATAACGGCGGATTAATTGCCCCCTCTCGCGTGTCAAAGCTCATAGGCCG GGGCCTGGGGATTCAGTCTGACGCTCCAATTGATAATAATTGCGAGTCAAAATGCTTCTGGCGCGGGGGGTCTATTAATACTCGATTGCCAT TCCAGAATCTCAGTCCCCGGACCGTGGGGCAGTGCCCAAAGTATGTGAACCGGCGGAGCCTGATGCTGGCTACTGGAATGCGGAATGTGC CCGAACTTATTCAGGGCCGAGGTCTGTTCGGCGCCATCGCCGGGTTCCTGGAGAATGGGTGGGAGGGCATGGTCGATGGATGGTATGGAT TTCGTCACCAGAACGCACAGGGCACAGGACAGGCTGCCGACTATAAGTCTACACAAGCGGCTATCGACCAGATTACTGGTAAACTCAACCG GCTGGTGGAGAAGACCAACACTGAGTTCGAATCCATTGAATCTGAATTTTCGGAGATCGAACACCAGATTGGAAACGTGATCAACTGGACCA AGGACTCAATCACTGACATCTGGACTTACCAGGCCGAACTGCTTGTGGCTATGGAGAATCAGCATACTATCGATATGGCCGACTCCGAAAT GCTCAACCTTTACGAGCGAGTGCGGAAACAATTGAGACAAAACGCCGAAGAGGATGGGAAGGCGTGTTTTCAAATATACCACGCCTGCGAT GATAGTTGTATGGAGTCAATCAGGAATAACACTTATGATCATAGTCAGTATAGAGAGGAAGCTCTTCTGAATCGACTGAACATCAACCCAGTA ACACTTAGTAGTGGGTACAAAGACATCATCTTGTGGTTCTCATTCGGAGCCTCTTGTTTTGTGCTGCTTGCCGTCGTAATGGGACTGTTTTTT TTTTGTCTCAAGAACGGCAACATGAGGTGCACTATTTGCATT

2988	ATGTACAAAATCGTGGTGATCATCGCACTGCTAGGGGCAGTGAAAGGCCTGGATAAAATCTGCCTGGGACACCACGCAGTAGCCAACGGTA CCATTGTGAAAACACTGACAAACGAACAGGAGGAGGTCACAAATGCTACAGAGACCGTGGAGTCTACAGGTATTAATCGGCTGTGCATGAA GGGAAGAAAGCACAAGGATCTGGGGAACTGCCACCCCATTGGAATGCTCATCGGGACACCTGCATGCGATCTGCACCTCACTGGCATGTG GGACACATTGATCGAAAGAGAAAATGCCATTGCATATTGCTATCCTGGAGCAACGGTCAATGTGGAAGCACTGAGGCAAAAGATCATGGAG TCCGGAGGCATCAACAAAATCTCAACCGGTTTCACATACGGAAGCAGTATTAACTCTGCCGGCACAACCCGTGCTTGCATGCGAAATGGAG GGAACTCTTTCTACGCGGAACTCAAATGGTTGGTGTCTAAGTCGAAAGGACAGAACTTTCCGCAAACTACAAATACTTACAGGAATACTGAC ACTGCAGAACATCTGATCATGTGGGGCATCCACCACCCCTCTTCAACACAGGAAAAGAATGACCTGTACGGGACCCAGAGTCTGTCAATCT CCGTGGGCTCCTCAACTTATCGTAACAACTTTGTCCCTGTTGTGGGAGCACGACCTCAAGTGAACGGACAGAGCGGGAGAATCGATTTTCA CTGGACCCTCGTGCAGCCTGGAGATAACATCACATTTAGCCATAACGGAGGGTTGATCGCTCCGTCCCGTGTGTCTAAACTTATCGGACGG GGCCTGGGTATCCAGAGTGATGCACCAATCGATAACAACTGTGAGAGCAAGTGTTTCTGGCGTGGTGGTTCTATAAACACACGACTGCCCT TCCAGAACCTGTCGCCTAGAACCGTGGGACAGTGTCCTAAATACGTCAACCGAAGAAGCCTAATGCTGGCCACAGGAATGCGCAACGTGC CCGAGCTAATTCAGGGCAGGGGCCTCTTCGGCGCAATCGCAGGCTTTCTGGAGAATGGGTGGGAGGGGATGGTGGATGGTTGGTACGGG TTCCGACATCAGAACGCCCAGGGGACGGGGCAGGCAGCCGACTACAAATCCACTCAGGCCGCAATAGACCAGATTACCGGGAAACTGAAT AGACTGGTGGAGAAAACAAATACTGAGTTCGAGAGTATTGAATCAGAGTTCTCCGAAATCGAGCATCAGATCGGCAACGTGATTAACTGGAC TAAGGACTCTATCACAGATATATGGACATATCAGGCCGAGCTGCTGGTGGCCATGGAGAACCAGCACACCATCGACATGGCCGACTCAGAA ATGTTAAACCTCTATGAGCGGGTTCGTAAGCAGTTACGCCAAAACGCCGAAGAGGATGGCAAGGGGTGCTTCGAAATTTATCACGCTTGTG ATGATTCCTGTATGGAGTCCATTCGGAACAACACCTACGACCACAGCCAGTACCGCGAGGAGGCCCTTCTGAATCGTCTGAACATAAACCC GGTCACCCTCAGTTCAGGATACAAGGACATAATTCTTTGGTTCTCCTTTGGCGCCTCATGCTTCGTGCTCCTGGCCGTGGTGATGGGGCTC TTTTTCTTCTGCTTGAAGAACGGGAACATGCGGTGTACTATCTGCATT
2989	ATGTACAAAGTGGTGGTGATCATCGCCTTACTCGGGGCAGTGCGAGGGCTGGATAAGATTTGCCTGGGTCACCACGCTGTGGCAAACGGG ACAACAGTGAAAACTCTCACAAATGAACAGGAGGAAGTCACCAACGCAACAGAGACCGTCGAAAGTACCTCACTCAATAAGTTGTGTATGAA AGGTCGCCGGTACAAGGACCTCGGGAACTGCCACCCAATTGGAATGCTGATTGGTACGCCCGTGTGTGACCTGCACCTGACCGGGACGTG GGACACCCTGATCGAGCGCGAGAATGCAACTGCTTACTGCTACCCCGGCGTGACCATTAACGAAGAAGCGCTGCGCCAGAAAATCATGGA ATCAGGGGGGATTAGCAAAATGAGGACAGGGTTCACATATGGTCCTAGCATCAATAGCGCCGGAACGACACGGAGCTGCATGCGCAACGG AGGCAACTCATTTTATGCCGAACTTAAGTGGCTGGTGTCTGGGACAAAGGGGCAGAACTTCCCACAGACTACTAATACATATCGAAATACCG ACACTGCCGAGCACCTTATCATCTGGGGAATCCACCATCCTAGCTCAACTCAGGAAAAGAATGACCTCTACGGTACTCAGTCTCTGTCCATT AGCGTGGGATCCAGCACCTACCAAAACAATTTTGTTCCTGTTATCGGGGCAAGACCCCAAGTGAATGGCCAATCTGGGAGAATCGAATTCC ACTGGACACTGGTCAGACCAGGCGACAACATAACTTTTTCACACAATGGAGGACTTATCGCCCCAGACAGGGTGTCAAAGCTGATCGGAAA GGGCATCGGAATTCAGAGTGGTGCGGTGATTGATAAGGATTGTGAAAGCAAGTGCTTTTGGCGCGGGGGCTCCATTATCACCGAGCTCCCT TTTCAAAACCTTAGCCCCCGGACAGTCGGCCAATGCCCAAAATATGTCAAAAAGCGCTCACTGCTCCTTGCCACGGGGATGCGGAACGTGC CGGAAGTGGTGCAAGGTCGAGGCCTCTTTGGCGCGATCGCCGGATTCATCGAGAATGGTTGGGAAGGCATGGTGGACGGGTGGTACGGG TTTAGACACCAGAATGCGCAGGGCATTGGGCAGGCTGCTGACTACAAATCAACTCAGACTGCGATTGATCAGATCACGGGAAAACTGAACC GTCTCATCGAAAAAACAAACACTGAGTTCGAATCCATCGAGTCGGAATTTAGTGAAATCGAACACCAGATCGGCAACGTCATTAATTGGACG AAGGACTCTATTACAGATATCTGGACCTATCAGGCTGAGCTCTTGGTAGCAATGGAAAACCAACACACTATCGATATGGCCGATAGCGAGAT GCTGAACCTGTATGAAAGGGTTAGAAAGCAGCTGAGACAGAACGCTGAAGAAGACGGTAAGGGCTGCTTTGAGATCTATCATACATGCGAC AATTCATGTATGGAGTCCATACGGAATAATACATATGACCATTCGCAGTACCCAGAGGAGGCCTTACTGAACAGACTGAACATCAATCCTGT TAAGCTGTCAAGCGGTTACAAGGACATAATTCTCTGGTTCTCCTTTGGTGCCTCCTGTTTCGTACTTCTCGCCGTCATCATGGGGTTGGGCT TCTTTTGCCTGAAAAATGGCAACATGAGATGCACTATATGCATT
2990	ATGTATAAGATAGTGGTAATAATTGCCCTCTTGGGGGCAGTGAAAGGACTCGATAAGATATGCCTGGGACACCATGCCGTGGCTAACGGCA CAATTGTGAAAACGCTAACAAACGAGCAGGAGGAAGTTACCAACGCTACCGAAACCGTTGAGAGTACCGGCATTAATAGACTTTGTATGAAG GGCCGAAAACACAAAGATCTCGGGAACTGTCATCCGATCGGCATGTTGATTGGAACACCTGCATGTGATCTCCATTTAACAGGCATGTGGG ATACATTGATTGAGCGGGAGAACGCTATAGCCTATTGCTATCCCGGAGCCACAGTGAATGTTGAGGCGCTTCGTCAGAAAATCATGGAGAG CGGAGGCATCAACAAGATCTCCACCGGGTTTACCTATGGCTCGTCTATAAACTCAGCTGGTACAACCAGGGCTTGCATGAGAAATGGGGGA AATTCCTTTTATGCAGAGCTGAAGTGGCTGGTGTCAAAGAGCAAAGGACAGAATTTCCCGCAGACCACAAACACCTATCGTAATACCGACAC CGCCGAGCACCTTATTATGTGGGGCATACACCATCCAAGCTCAACCCAGGAGAAGAACGATCTGTACGGCACGCAGAGTCTGAGCATCTCA GTCGGCTCTTCCACTTATAGGAACAACTTCGTTCCAGTTGTCGGGGCGCGCCCTCAAGTCAACGGACAATCTGGCAGGATCGACTTCCACT GGACTCTTGTACAACCCGGCGATAACATCACCTTCTCCCATAACGGGGGTCTTATCGCCCCATCCCGGGTGAGCAAGCTCATTGGTCGCGG ACTGGGGATTCAGAGCGATGCGCCTATTGACAATAACTGCGAGAGTAAATGCTTTTGGAGGGGCGGTAGCATCAACACCCGGTTGCCTTTC CAAAACCTTTCCCCTAGAACAGTTGGCCAGTGTCCGAAGTATGTAAACAGGCGAAGTCTCATGCTCGCGACAGGCATGCGGAATGTTCCCG AGCTGATTCAGGGCAGGGGACTGTTTGGAGCTATCGCGGGATTTCTGGAAAACGGTTGGGAAGGGATGGTGGACGGATGGTATGGATTCC GCCACCAGAATGCACAAGGAACGGGCCAGGCCGCCGACTACAAGTCCACTCAGGCCGCTATTGATCAGATTACCGGCAAGCTCAATCGCC TGGTGGAGAAGACTAATACCGAGTTCGAGAGCATCGAGTCCGAATTCTCGGAAATTGAGCATCAGATCGGCAATGTCATAAATTGGACCAAA GACTCAATCACAGATATATGGACTTATCAGGCTGAGTTGCTTGTGGCTATGGAAAACCAGCACACAATCGATATGGCTGATTCCGAGATGCT TAATCTGTATGAACGCGTAAGGAAGCAGTTACGGCAAAATGCAGAAGAAGCATGGCAAGGGCTGTTTTGAGATCTATCATGCCTGCGATGAC AGTTGCATGGAGTCAATTAGGAATAATACGTATGATCACTCTCAGTATCGTGAGGAAGCTCTGCTGAACCGTCTGAACATCAATCCTGTGAC CTTATCGTCCGGATATAAGGATATCATTCTATGGTTCAGCTTTGGCGCTAGCTGCTTTGTCTTACTCGCTGTGGTCATGGGCCTGTTTTTTTT CTGCTTAAAGAACGGGAACATGCGTTGTACAATCTGCATT

2991	ATGTATAAAATCGTCGTTATAATCGCGTTATTGGGAGCAGTGAAGGGCTTAGACAAGATTTGTCTGGGACATCATGCAGTGGTGAACGGAAC CATCGTGAAGACACTAACAAATGAACAGGAGGAAGTCACCAACGCAACCGAAACTGTGGAATCTACAGGTCTAAACAGGTTGTGTATGAAA GGCAGGAATCATAAAGACCTCGGCAACTGCCATCCAATTGGTATGCTTATTGGAACGCCCGCATGTGACCTGCATCTCACAGGTACATGGG ACACACTCATTGAGCGCGAAAATGCTATCGCCTACTGCTATCCTGGCGCTACCGTCAATGAGGAGGCACTGAGACAGAAGATCATGGAGAG CGGCGGAATAAACAAAATTTCTACAGGCTTCACCTATGGGAGCAGCATCAATTCTGCCGGCACAACAAGGGCCTGTATGCGTAACGGGGGA AACTCCTTTTACGCCGAGCTGAAATGGTTGGTGTCCAAGAGCAAGGGGCAGAACTTTCCCCAGACCACAAATACATATAGGAATACGGATAC TGCCGAGCACCTCATCATGTGGGGCATCCATCACCCAAGCTCCACCCAGGAGAAGAATGACCTCTACGGGACCCAGTCCCTTAGCATTTCT GTGGGCAGTAGCACTTATCAAAATAACTTCGTCCCCGTCGTAGGGGCGAGACCACAGGTTAACGGACAAAGCGGCAGAATTGACTTCCACT GGACACTCGTTCAGCCAGGTGATAATATTACTTTTAGCCATAATGGGGGACTGATAGCACCCAGCAGAGTGAGTAAGTTGATAGGCAGGGG GCTTGGTATTCAGTCAGATGCGCCCATTGATAACAACTGTGAGAGTAAGTGCTTCTGGAGAGGGGGCTCCATTAATACCCGCTTGCCATTTC AAAACCTGTCCCCCAGGACTGTAGGGCAGTGCCCTAAGTACGTGAACAAACGGTCCTTAATGCTGGCCACTGGTATGCGCAATGTGCCAGA GCTCATGCAGGGCAGAGGGCTCTTTGGAGCCATAGCAGGGTTTATTGAAAATGGGTGGGAGGGCATGGTGGACGGATGGTACGGATTTAG ACACCAAAACGCGCAGGGCACTGGACAGGCGGCCGACTATAAGTCCACCCAGGCCGCGATCGATCAGATCACCGGAAAGTTGAACCGCTT AATAGAGAAGACAAACACCGAGTTCGAAAGCATAGAGTCCGAATTTTCCGAAATCGAACACCAGATCGGCAACGTGATCAACTGGACGAAG GATTCGATTACCGACATTTGGACTTATCAGGCTGAACTACTCGTGGCTATGGAGAACCAACATACTATCGATATGGCAGATTCTGAAATGCTT AACCTCTATGAACGGGTTAGGAAACAGCTGAGGCAGAACGCAGAAGAAGACGGCAAGGGGTGCTTCGAAATTTATCATGCATGTGACGATA GTTGCATGGAAAGCATTAGAAACAACACCTATGACCATTCTCAGTATCGGGAGGAGGCACTCCTGAATCGTCTGAATATCAACCCAGTCACA CTCAGTTCAGGGTATAAAGACATAATTCTCTGGTTCTCATTCGGAGCCTCCTGTTTTGTGCTACTCGCAGTTGTCATGGGACTGGTCTTTTTT TGCCTGAAAAATGGCAATATGAGATGTACAATTTGCATC
2992	ATGTATAAAATTGTGGTCATAATTGCACTGCTCGGCGCAGTAAAGGGGCTCGATAAGATTTGCCTGGGCCACCACGCCGTCGCTAACGGGA CGATTGTCAAGACCCTGACTAACGAACAAGAGGAGGTGACTAATGCTACCGAAACCGTCGAATCCACCGGCATTAACAGACTCTGTATGAA AGGCAGAAAACATAAGGATCTGGGAAATTGCCACCCAATCGGAATGCTGATCGGGACACCCGCCTGCGACCTGCACCTGACTGGGACTTG GGATACACTTATCGAAAGGGAGAATGCCATTGCTTATTGTTACCCCGGCGCAACAGCGAATGTTGAAGCCCTGCGGCAAAAAATTATGGAAT CCGGTGGCATTGACAAGATCAGCACCGGGTTCACCTACGGCTCATCTATTAACAGTGCTGGGACCACCAGAGCTTGCATGCGGAACGGTG GCAACAGTTTCTACGCTGAACTCAAGTGGTTAGTCTCAAAGTCTAAAGGACAGAACTTCCCCCAGACTACTAATACGTATAGGAACACTGAT ACTGCTGAACATCTGATTATGTGGGGCATTCACCATCCAAGCTCAATTCAGGAGAAAAACGACCTGTATGGCACCCAGTCGCTGTCTATCAG CGTGGGTTCCAGCACCTACCGGAACAATTTTGTACCGGTGGTTGGCGCGCGTCCTCAGGTCAATGGTCAATCTGGGCGGATTGACTTCCAC TGGACTCTGGTTCAACCCGGCGACAACATAACTTTCTCTCACAATGGTGGATTGATCGCTCCTTCCCGCGTTTCAAAACTGATAGGCCGCGG CTTAGGAATTCAGAGCGACGCCCCGATAGACAATAATTGCGAGTCCAAGTGCTTCTGGAGGGGAGGAAGCATTAACACGCGGTTGCCTTTC CAGAACCTGTCCCCAAGGACAGTGGGACAATGCCCAAAATATGTCAATAGAAGAAGTCTCATGCTGGCGACTGGTATGCGCAACGTGCCTG AGCTTATCCAGGGCCGCGGATTGTTCGGGGCAATCGCCGGCTTCCTGGAAAATGGGTGGGAAGGAATGGTGGATGGTTGGTACGGTTTTC GACATCAGAATGCCCAGGGTACTGGCCAGGCTGCAGACTATAAAAGCACTCAGGCTGCCATTGACCAGATCACGGGTAAACTGAATAGGCT GGTGGAGAAGACGAACACAGAGTTCGAGTCTATTGAATCTGAGTTTTCTGAGATAGAGCACCAGATCGGAAACGTCATCAACTGGACCATG GATTCCATAACTGACATCTGGACTTACCAGGCGGAACTGCTGGTTGCAATGGAGAATCAGCACACTATTGACATGGCCGATTCCGAGATGC TCAATCTTTACGAGCGGGTCCGTAAACAGCTAAGGCAGAATGCCGAGGAAGATGGGAAGGGTTGCTTCGAAATATATCATGCATGTGATGA TTCTTGTATGCAGTCGATTCGTAATAACACCTACGATCACAGTCAATACAGACAAGAAGCTCTTCTTAATCGCCTAAACATTAACCCAGTCAC ATTGAGCTCAGGATACAAGGACATCATCCTGTGGTTCTCATTCGGGGCTTCCTGTTTTGTCCTGCTGGCCGTGGTCATGGGGCTGGTTTTTT TTTGCCTGAAGAACGGCAATATGAGATGCACAATTTGCATC
2993	ATGTATAAGGTGGTTGTGATTATTGCGCTGCTCGGGGCCGTTCGAGGCCTGGATAAAATCTGTCTGGGCCATCACGCTGTCGCCAATGGCA CGACTGTGAAAACTTTAACAAATGAGCAGGAAGAGGTGACAAATGCCACTGAAACCGTCGAGAGTACATCACTGAATAAACTGTGCATGAAG GGACGCCGCTATAAGGACCTGGGTAATTGCCATCCAATTGGTATGTTGATCGGGACCCCTGTGTGCGACTTACACCTAACTGGCACTTGGG ACACACTGATCGAGAGGGAGAATGCCACCGCCTATTGTTACCCTGGTGTGACCATTAACGAAGAGGCTCTCAGACAAAAGATTATGGAGTC TGGAGGAATCTCTAAGATGCGTACTGGGTTTACCTATGGGCCAAGCATCAACTCGGCTGGCACTACTCGCTCCTGTATGAGAAATGGAGGA AACTCCTTTTATGCTGAGTTGAAGTGGCTCGTTTCCGGCACCAAGGGGCAAAACTTTCCACAGACGACCAATACTTATAGGAATACGGACAC CGCTGAACACCTGATTATTTGGGGAATCCATCACCCCAGCTCTACGCAGGAGAAGAACGACCTTTACGGTACCCAGAGTCTTTCGATTAGC GTGGGCAGCAGCACATACCAGAATAATTTCGTGCCCGTAATAGGTGCACGGCCTCAGGTGAATGGACAGTCGGGCCGGATAGAATTCCAC TGGACGCTCGTTAGGCCCGGAGACAACATAACTTTTTCTCATAATGGCGGACTTATCGCTCCAGATCGGGTCAGCAAGCTTATAGGGAAGG GGATCGGCATCCAGTCCGGTGCTGTCATCGATAAGGACTGTGAATCCAAGTGTTTCTGGCGTGGCGGTTCAATTATCACCGAACTACCTTTT CAAAACTTGTCACCAAGAACCGTGGGACAGTGTCCAAAGTATGTCAAAAAAAGATCCTTGTTACTGGCTACCGGGATGAGGAATGTGCCTGA AGTCGTTCAGGGCCGCGGGCTGTTCGGAGCCATCGCGGGATTTATCGAAAACGGGTGGGAGGGGATGGTGGACGGTTGGTACGGATTCA GGCATCAGAATGCTCAGGGGATTGGGCAGGCTGCCGACTACAAGTCTACCCAGACGGCTATTGACCAGATCACCGGCAAGTTGAATCGCC TGATAGAAAAAACTAATACGGAATTCGAAAGCATCGAATCCGAGTTCTCAGAAATCGAGCACCAGATAGGTAATGTAATCAATTGGACAAAG GACTCTATCACCGATATCTGGACCTATCAGGCAGAATTGCTGGTGGCAATGGAGAACCAGCACACAATCGATATGGCTGATTCAGAAATGCT GAATTTGTACGACAGGGTCCGGAAGCAGCTCCGACAGAACGCAGAAGAAGATGGGAAGGGTTGTTTCGAGATATATCATACCTGCGACAAC TCGTGTATGGAATCAATACGCAATAATACATACGATCACAGTCAGTATCGTGAGGAGGCTCTCCTGAACAGGCTGAATATTAATCCTGTGAA ACTGAGCTCCGGGTACAAGGATATTATTCTGTGGTTCAGTTTTGGCGCTTCATGTTTCGTGTTGCTGGCAGTTATTATGGGGCTTGGTTTCTT CTGCCTGAAGAATGGGAACATGCGCTGCACGATTTGCATA

2994	ATGTATAAGGTGGTCGTAATCATCGCCCTTCTGGGTGCTGTCCGGGGGCTTGGACAAAATTTGTCTCGGGCATCACGCAGTGGCTAACGGCA CAATAGTGAAGACCCTCACTAACGAACAGGAGGAAGTGACAAACGCGACAGAAACAGTTGAGAGTAAATCCCTCGGCAAGCTGTGCATGAA AGGGAGGAGTTATAATGACCTCGGCAATTGTCATCCGATAGGTATTCTGATCGGCACTCCTGCGTGCGACCTGCACCTGACCGGGACCTGG GATACCCTCATCGAGAGGGAGAACGCCGTGGCCTACTGCTACCCAGGCGCCACAGTAAATGAGGAAGCGCTGCGGCAGAAAATTATGGAA TCTGGAGGAATTACCAAGATTTCTACTGGCTTCACCTATGGGACAACCATCAACAGCGCCGGTACCACGAAGGCCTGTATGAGAAACGGTC GGAATTCTTTTTACGCAGAGCTCAAATGGCTGGTGAGCAAAAACAAGGGGCAAAATTTCCCCCAGACCACAAACACATATAGGAATACAGAC ACAGCTGAGCATTTAATTATTTGGGGCATCCATCACCCCTCCAGCACACAAGAAAAGAATGACCTGTACGGTACCCAGTCTCTCTCTATTAG CGTTGGCTCTTCTACATACCAGAATAACTTTGTTCCAGTCGTCGGAGCACGGCCACAGGTTAACGGGCAGAGTGGACGTATCGATTTCCAC TGGACTCTTCTACAGCCTGGCGACAATATTACTTTTAGTCACAACGGCGGCCTTATAGCCCCGTCCAGAGTCTCCAAGCTGATCGGCAGAG GGCTGGGCATTCAGTCCGAGGCTCCAATTGATAATGGCTGCGAAAGCAAATGCTTTTGGAAAGGCGGGAGTATCAACACAAAACTGCCCTT TCAGAATCTGAGCCCTCGAACTGTGGGTCAGTGCCCAAAATACGTTAATAAGCGGAGTCTGATGTTAGCTACTGGCATGCGGAACGTTCCA GAAATCATGCACGGCCGCGGACTCTTTGGAGCTATCGCCGGGTTTATAGAAAATGGCTGGGAAGGCATGGTTGACGGCTGGTACGGATTC AGGCACCAAAACGCCCAGGGAACCGGACAGGCCGCCGATTACAAAAGCACGCAAGCAGCTATTGATCAGATCACTGGAAAACTCAATCGA CTGATCGAAAAGACTAACACAGAATTTGAATCCATTGAGTCCGAATTTAGTGAGATCGAGCACCAAATCGGCAACATCATTAACTGGACGAA GGACAGCATAACAGACATATGGACATATCAGGCAGAGCTCCTGGTCGCAATGGAAAACCAGCATACCATAGATATGGCTGATTCTGAGATG CTCAATCTGTATGAACGAGTCAGAAAGCAATTAAGACAAAACGCTGAGGAGGATGGTAAGGGATGCTTCGAGATTTACCATGCTTGCGACG ATTCCTGCATGGAAAGCATCCGTAATAACACCTACGACCACAGCCAATACCGTGAAGAAGCCCTCCTAAATCGATTGAACATTAATCCTGTG AAATTGAGTTCCGGCTACAAGGACATCATCCTGTGGTTCTCATTTGGCGCATCATGCTTGATTCTTTTGGCCGTGGTGATGGGCCTGGTTTT CTTCTGTCTGAAAAATGGCAACATGAGATGTACTATCTGCATT
2995	ATGTACAAGATCGTGGTTATAATTGCCCTCCTGGGCGCCGTTAAGGGCCTTGATAAGATATGTCTGGGGCACCATGCCGTGGCTAATGGGA CCATAGTGAAAACCCTGACTAATGAGCAGGAGGAGGTCACTAACGCCACCGAGACTGTGGAATCCACCGGCATCAACAGGCTGTGTATGAA GGGGCGCAAACACAAGGACCTAGGGAATTGCCACCCCATCGGTATGCTTATCGGCACGCCAGCCTGTGACCTTCACCTGACTGGGATGTG GGACACTCTGATTGAAAGAGAAAACGCTATAGCGTACTGCTACCCCCGGCGCAACTGTGAACGTAGAAGCGTTGCGGCAGAAGATCATGGA GTCTGGGGGAATCAATAAGATCTCAACAGGGTTTACATACGGCTCATCTATCAATTCCGCGGGCACGACCAGGGCGTGCATGAGGAACGG CGGGAACTCTTTTTACGCCGAGCTTAAATGGTTAGTATCTAAGAGCAAGGGGCAGAACTTTCCACAAACGACCAACACATACCGCAACACCG ATACCGCAGAACACCTCATCATGTGGGGAATCCACCATCCCAGTTCCACTCAAGAAAAAAACGATCTATACGGCACTCAGTCCCTCTCAATC TCCGTGGGAAGTAGCACATATCGGAACAACTTCGTGCCCGTGGTTGGGGCCCGCCCCCAGGTCAATGGCCAGTCAGGAAGAATTGACTTC CATTGGACACTTGTGCAGCCAGGCGACAATATCACATTTTCCCACAACGGCGGATTGATCGCCCCCTCCAGAGTTAGCAAGCTGATCGGGA GAGGCCTGGGTATCCAGAGTGACGCGCCAATCGATAACAACTGCGAGAGCAAGTGCTTTTGGCGCGGAGGCTCAATTAACACACGGCTAC CTTTCCAAAACCTCAGTCCGCGTACAGTTGGCCAGTGCCCCAAGTATGTGAACCGGAGATCTTTGATGCTCGCTACCGGCATGCGAAATGT CCCGGAGCTGATTCAAGGCCGCGGGCTCTTTGGAGCGATTGCGGGGTTTCTGGAGAACGGATGGGAAGGAATGGTGGATGGTTGGTACG GATTTCGGCATCAAAATGCACAGGGGACTGGCCAAGCCGCAGACTATAAGAGCACCCAGGCTGCGATCGACCAGATCACCGGAAAGCTCA ACCGCCTGGTGGAGAAGACCAATACGGAATTTGAATCAATCGAAAGCGAGTTTAGTGAAATAGAACACCAGATTGGTAATGTGATTAATTGG ACAAAAGACAGCATTACTGATATTTGGACATATCAGGCTGAGTTACTGGTAGCTATGGAGAATCAGCACACCATTGATATGGCCGACTCCGA AATGCTGAATCTGTACGAACGGGTTCGGAAACAGCTCAGGCAGAACGCCGAAGAAGACGGTAAGGGATGCTTCGAAATCTACCACGCCTG CGATGATAGTTGCATGGAGTCTATACGGAACAACACCTATGACCATTCTCAGTATAGAGAAGAGGCCCTACTGAACCGTTTGAACATTAATC CAGTGACCCTGTCCTCCGGTTACAAGGACATAATCCTGTGGTTCTCGTTTGGTGCTAGCTGCTTTGTGCTGCTGGCGGTGGTAATGGGACT GTTTTTCTTTTGCCTGAAAAATGGCAATATGAGATGCACCATTTGCATC
2996	ATGTACAAGATTGTGGTGATTATAGCCCTTCTGGGGGCTGTCAAGGGACTGGACAAAATCTGCCTGGGACATCATGCTGTGGCAAACGGAA CTATTGTGAAGACTTTGACTAACGAACAGGAGGAGGTGACCAACGCCACCGAAACCGTGGAATCCACAGGCATCAACCGGCTCTGCATGAA AGGAAGAAAGCACAAGGACCTGGGCAATTGTCATCCAATCGGAATGCTTATCGGGACTCCGGCCTGTGATTTGCATCTGACCGGCATGTGG GATACGTTAATCGAACGAGAGAACGCCATTGCCTATTGCTACCCTGGCGCGACCGTGAATGTCGAGGCCTTGAGGCAGAAGATCATGGAGA GCGGCGGAATCAACAAAATAAGCACTGGTTTCACCTACGGTAGTAGCATAAATTCCGCTGGGACCACGCGGGCCTGCATGCGGAACGGCG GCAATTCCTTTTACGCAGAGCTTAAATGGTTAGTGAGCAAAAGCAAGGGACAGAATTTCCCCCAGACCACAAACACATATCGTAACACCGAC ACCGCGGAACATCTGATTATGTGGGGCATACATCACCCCTCTTCCACGCAAGAAAAGAATGACCTGTACGGAACACAATCCTTATCCATTAG CGTAGGAAGCAGCACTTATCGGAACAACTTTGTGCCCGTGGTGGGAGCTCGGCCTCAGGTAAACGGCCAGTCCGGTCGGATAGATTTCCA TTGGACTTTGGTTCAGCCGGGGGATAACATTACATTTTCCCATAACGGCGGCCTCATTGCTCCCTCGCGGGTCTCTAAACTTATTGGGCGG GGCCTGGGGATCCAGTCAGATGCCCCTATCGATAATAATTGCGAATCCAAATGTTTTTGGAGAGGAGGCAGTATTAACACCCGGCTGCCTTT CCAGAATCTCAGCCCCCGTACAGTCGGACAGTGTCCCAAATATGTGAATAGACGCTCGCTAATGCTGGCCACCGGCATGCGGAACGTTCCT GAGCTGATCCAGGGACGGGGGCTCTTTGGTGCAATTGCTGGATTTCTGGAAAATGGCTGGGAAGGCATGGTGGATGGCTGGTACGGCTTT AGGCACCAGAATGCCCAAGGCACGGGGCAGGCAGCCGACTACAAATCCACACAAGCCGCCATTGACCAAATCACCGGGAAATTAAATAGG CTTGTGGAGAAAACCAATACCGAATTCGAAAGCATCGAGAGCGAATTTTCAGAGATCGAGCACCAGATCGGGAATGTGATAAATTGGACCA AAGATTCCATTACCGATATTTGGACTTACCAAGCTGAGCTGCTCGTGGCCATGGAAAACCAGCACACAATCGATATGGCCGACTCTGAAATG TTGAACTTGTATGAAAGAGTGCGAAAACAGTTGCGTCAAAACGCCGAGGAACACGGGAAAGGTTGTTTCGAAATCTATCACGCGTGCGATG ACAGCTGTATGGAAAGTATTAGGAACAATACATATGACCACTCACAGTATCGCGAGGAGGCACTGTTGAATCGCCTAAACATCAACCCCGTT ACATTGTCTTCTGGTTACAAAGACATTATCTTATGGTTTTCGTTTGGAGCCTCGTGTTTCGTGCTCCTAGCGGTAGTGATGGGGCTCTTCTTC TTCTGTCTGAAGAACGGGAACATGCGCTGCACTATCTGTATC

2997	ATGTACAAGATCGTAGTCATCATCGCCCTGCTGGGGGCCGTCAAAGGACTGGACAAGATTTGCCTGGGCCATCATGCAGTCGCAAATGGCA CTATCGTGAAGACTCTGACAAACGAACAGGAAGAGGTGACAAATGCCACTGAAACAGTCGAGTCTACTGGGATCAACAGACTTTGCATGAA GGGCCGCAAGCACAAGGACCTGGGTAATTGCCACCCCATTGGAATGCTTATTGGCACACCCGCTTGCGACTTACACCTAACCGGTATGTGG GATACTCTGATCGAAAGGGAAAACGCCATTGCATACTGCTACCCCGGCGCTACAGTGAACGTCGAGGCTCTGCGCCAGAAAATTATGGAGT CTCGTGGAATCAATAAGATCTCAACTGGCTTCACCTATGGGAGTTCCATCAACAGCGCTGCGACTACGCGTGCGTGCATGCGTAACGGCGG AAACAGCTTTTACGCAGAGCTGAAATGGCTCGTCAGCAAGTCCAAGGGTCAGAATTTCCCACAGACTACGAATACCTACCGAAATACTGACA CTGCGGAGCATCTCATCATGTGGGGAATCCACCATCCAAGCAGCACCCAGGAAAAGAACGATCTGTACGGCACTCAGTCCCTAAGCATCTC CGTCGGCAGCAGTACTTATCGCAATAACTTTGTCCCCGTGGTAGGAGCCCGTCCTCAGGTGAACGGACAGAGCGGACGTATCGACTTTCAT TGGACACTCGTGCAACCCGGGGACAATATAACGTTCAGCCATAACGGAGGTTTAATAGCGCCTAGCCGTGTGTCCAAACTCATCGGCCGTG GCCTGGGGATCCAATCCGATGCACCAATCGACAATAACTGCGAGTCTAAGTGCTTCTGGCGGGGAGGCTCAATTAATACCCGCCTACCCTT CCAGAATCTGTCACCCAGGACCGTGGGCCAGTGTCCTAAGTACGTCAACAGGCGATCTCTCATGTTGGCCACTGGCATGCGAAACGTGCCT GAGCTGATCCAAGGTAGAGGGCTGTTCGGAGCAATCGCCGGATTCCTGGAGAACGGCTGGGAGGGAATGGTGGACGGATGGTATGGTTT CAGGCACCAAAACGCTCAAGGAACTGGCCAGGCAGCTGACTATAAGTCCACTCAGGCTGCAATCGATCAAATCACCGGCAAGCTGAACAG GCTCGTTGAGAAAACGAACACGGAATTTGAGTCCATCGAATCTGAATTTTCCGAGATCGAACATCAGATAGGCAACGTTATCAACTGGACGA AGGACTCTATCACTGACATTTGGACGTATCAGGCGGAGCTCTTAGTGGCCATGGAAAACCAGCACACTATCGACATGGCCGACAGTGAAAT GCTGAACCTCTACGAAAGAGTTCGCAAGCAGCTCCGGCAGAATGCAGAAGAGGATGGAAAAGGCTGTTTTGAAATCTACCACGCATGTGAC GACTCATGCATGGAGTCTATACGAAATAATACTTACGACCATTCGCAATATCGCGAGGAGGCTCTGCTAAATAGGTTGAACATCAATCCCGT CACACTGTCCAGTGGCTACAAGGATATCATTCTGTGGTTCTCATTTGGAGCCTCCTGCTTCGTTCTGCTGGCCGTGGTTATGGGATTATTCT TCTTTTGCCTCAAGAACGGAAATATGCGCTGCACAATCTGCATC
2998	ATGTACAAGATTGTGGTGATTATCGCATTGCTGGGCGCCGTGAAGGGGCTGGATAAAATATGTCTGGGGCACCATGCCGTAGCAAACGGTA CGATAGTTAAGACTCTTACCAACGAACAGGAAGAGGTGACGAACGCGACAGAGACCGTTGAGAGTACAGGTATTAATCGACTCTGTATGAA GGGCCGCAAGCACAAGGACCTGGGCAATTGCCACCCCATCGGGATGCTGATCGGCACTCCTGCGTGTGATCTGCACTTGACCGGCACTTG GGACACTTTGATTGAGAGAGAAAACGCGATTGCTTATTGCTACCCTGGAGCAACCGCCAATGTTGAAGCCCTGAGGCAGAAGATTATGGAG TCGGGAGGTATTGATAAGATCTCCACTGGATTCACTTACGGCTCATCTATAAATAGCGCCGGGACTACCCGCGCTTGCATGAGGAACGGGG GGAACAGCTTCTATGCCGAGCTCAAATGGCTTGTGTCTAAGTCGAAAGGTCAAAACTTCCCACAGACAACTAACACATACCGCAACACCGAC ACTGCAGAGCACTTGATTATGTGGGGCATCCATCATCCGTCTAGCATCCAGGAGAAGAATGATCTCTATGGGACCCAGAGCCTCAGCATTT CAGTCGGAAGTTCCACCTACAGGAATAATTTCGTGCCTGTGGTGGGCGCACGGCCTCAAGTGAACGGCCAGTCCGGAAGAATAGACTTTCA CTGGACTCTGGTGCAGCCGGGGGATAACATTACATTCTCGCACAACGGTGGGCTTATAGCCCCCAGCAGAGTAAGTAAACTTATCGGTCGG GGTTTGGGCATTCAGTCGGATGCCCCTATCGATAATAATTGTGAATCTAAATGTTTTTGGAGGGGCGGGAGCATCAACACCAGGCTGCCTTT CCAGAATCTGTCCCCCCGAACTGTGGGACAGTGCCCCAAGTATGTTAACAGACGGTCCCTTATGCTGGCCACTGGCATGCGCAACGTCCC CGAACTCATTCAGGGGCGCGGTCTGTTTGGTGCAATTGCCGGGTTTCTGGAGAACGGCTGGGAAGGGATGGTAGACGGTTGGTATGGCTT CCGGCACCAAAACGCCCAGGGAACTGGCCAAGCTGCCGACTACAAATCTACCCAGGCTGCCATAGATCAGATCACTGGCAAACTCAATCG CCTGGTGGAAAAAACCAACACAGAGTTTGAGAGCATCGAGTCCGAATTCTCCGAAATCGAACACCAGATCGGTAATGTGATTAACTGGACCA TGGATAGTATCACCGACATATGGACATATCAGGCTGAACTCCTGGTGGCCATGGAGAACCAACATACTATTGATATGGCCGACTCAGAAATG TTAAATCTGTATGAGCGTGTGAGAAAACAGCTCAGGCAGAACGCTGAGGAAGACGGAAAGGGCTGCTTCGAAATCTACCACGCATGCGATG ACTCTTGCATGGAATCAATCCGGAACAACACATATGACCACAGCCAGTACCGGGAGGAGGCCCTTTTGAATAGGCTCAATATCAATCCCGT CACGCTTTCTTCCGGTTACAAAGATATAATTCTGTGGTTCTCATTCGGGGCAAGCTGCTTTGTGCTTCTCGCTGTCGTGATGGGCCTCGTTTT CTTTTGCTTGAAGAATGGAAATATGCGGTGCACGATCTGTATC
2999	ATGTACAAGATCATCGTTATAATCGCACTCCTGGGCGCCGTCAAAGGCCTGGACAAGATTTGCCTAGGGCATCACGCCGTAGCCAATGGAA CCATTGTCAAGACCCTGACCAACGAGCAGGAAGAGGTGACAAACGCCACAGAGACAGTAGAGTCCACCGGAATCAATCGCTTGTGCATGAA AGGACCGAAGCATAAAGACTTAGGAAATTGTCATCCAATCGGCATGCTGATTGGGACACCCGCCTGTGACTTGCATTTGACAGGGACCTGG GACACCCTTATCGAGCGTGAGAATGCCATCGCGTACTGCTACCCCGGGGCTACGGTGAACGTCGAGGCTTTAAGACAGAAGATCATGGAG TCCGGCGGAATCGATAAGATTTCTACCGGTTTCACTTACGGGAGTTCTATCAACTCTGCAGGAACCACTCGGGCCTGCATGAGGAATGGAG GCAACTCATTCTACGCAGAATTGAAATGGCTGGTAAGCAAGAGCAAGGGCCAAAATTTCCCCCAGACCACGAATACCTATAGGAATACTGAC ACTGCGGAGCATCTGATTATGTGGGGCATCCACCACCCGAGCTCAACTCAAGAGAAAAATGATCTTTACGGAACCCAGAGCCTGTCAATTTC CGTGGGGAGCAGTACTTATCGCAATAATTTCGTACCTGTCGTAGGTGCCAGACCGCAGGTCAATGGACAGAGCGGCCGTATCGACTTCCAC TGGACACTGGTGCAGCCAGGGGATAACATCACTTTTTCCCACAACGGCGGCCTGATCGCTCCCAGCAGGGTGTCGAAGCTGATCGGTCGG GGGCTGGGGATACAGTCAGACGCACCCATTGATAACAATTGCGAGAGCAAATGTTTCTGGCGTGGTGGAAGCATAAACACAAGACTTCCGT TTCAGAACCTTAGTCCACGCACTGTCGGCCAGTGCCCAAAATACGTAAACCGACGTTCCCTGATGCTCGCTACGGGGATGCGAAACGTACC AGAGTTAATCCAGGGCCGCGGGCTGTTCGGAGCTATTGCCGGCTTCCTCGAAAACGGGTGGGAAGGCATGGTAGACGGGTGGTACGGGT TTAGGCATCAAAACGCCCAGGGCACAGGCCAGGCGGCCGACTATAAAAGCACCCAGGCTGCAATTGACCAGATTACCGGGAAACTGAACA GACTTGTAGAAAAAACCAATACAGAATTTGAGTCTATCGAATCAGAGTTCTCTGAGATAGAGCATCAGATCGGTAACGTGATCAATTGGACTA AAGATTCAATAACTGACATATGGACTTACCAGGCCGAACTTCTGGTGGCTATGGAAAACCAGCATACAATTGACATGGCCGACTCCGAGATG TTGAACCTTTACGAGCGCGTGCGTAAACAGCTGAGGCAAAATGCTGAAGAAGACGGAAAAGGCTGCTTCGAGATATATCACGCTTGTGACG ATTCATGTATGGAATCCATACGGAATAATACTTACGACCATTCGCAATATAGGGAGGAGGCCTTGTTGAACCGGCTGAATATCAATCCAGTG ACCCTTAGCTCCGGCTATAAAGATATTATTTTGTGGTTCTCTTTTGGAGCTTCCTGCTTTGTCCTATTAGCTGTGGTCATGGGCCTCTTTTTTT TCTGCCTTAAAAATGGGAATATGCGGTGCACTATCTGTATC

3000	ATGTATAAGATCGTGGTGATAATCGCCCTGCTGGGGGCTGTTAAAGGTCTCGACAAAATCTGCTTAGGGCATCACGCCGTGGCCAATGGCA CCATAGTTAAGACCCTGACCAATGAACAGGAAGAGGTTACCAACGCAACGGAGACCGTCGAATCTACCGGCATCAACCGATTATGCATGAA GGGTCGCAAGCACAAAGATCTCGGAAACTGCCACCCCATTGGCATGCTCATCGGGACCCCTGCTTGCGACCTGCATTTAACAGGTATGTGG GACACGCTGATTGAAAGAGAGAACGCGATCGCCTACTGTTACCCAGGCGCTACAGTGAACGTGGAGGCTCTGCGACAGAAAATTATGGAGT CGGGCGGCATCAATAAAATCAGCACGGGTTTCACTTATGGCAGTTCAATCAACTCCGCCGGTACTACCAGGGCCTGTATGCGGAATGGGG GCAACTCTTTTTATGCCGAGCTGAAGTGGCTGGTCTCCAAGAGTAAGGGCCAGAATTTCCCACAGACCACAAACACTTACAGGAACACGGA CACCGCCGAGCATCTGATTATGTGGGGCATTCACCACCCATCTAGTACCCAGGAGAAGAACGACTTGTACGGCACACAGTCTCTGTCAATT AGTGTTGGGTCTTCCACATATCGTAATAATTTTGTACCGGTCGTGGGCGCGGGGCCCCAAGTGAACGGTCAAAGCGGTAGGATTGATTTTC ATTGGACCCTCGTGCAACCAGGAGATAATATCACTTTTTCGCACAATGGCGGTCTCATAGCCCCGAGCCGGGTCAGCAAGCTAATCGGACG GGGGCTCGGGATCCAGTCCGATGCTCCCATCGATAATAACTGTGAATCTAAGTGTTTTTGGCGGGGCGGATCCATAAACACCAGGCTTCCC TTCCAGAACTTGTCCCCAAGGACGGTAGGCCAATGCCCAAAGTACGTTAACCGCCGCTCACTCATGTTGGCCACTGGGATGAGAAACGTCC CGGAACTGATTCAGGGCCGGGGCCTGTTTGGCGCAATCGCTGGGTTCTTAGAGAACGGCTGGGAGGGCATGGTGGACGGATGGTACGGG TTCCGCCATCAGAACGCTCAAGGCACTGGGCAGGCTGCAGATTACAAGAGTACCCAGGCTGCCATTGATCAGATTACCGGAAAACTGAATA GGCTGGTCGAAAAGACAAACACTGAGTTCGAAAGCATCGAGAGCGAATTTTCTGAGATCGAGCACCAGATCGGGAATGTGATTAACTGGAC TAAAGATAGCATAACGGATATTTGGACATATCAAGCAGAGTTGTTGGTGGCCATGGAGAATCAGCATACAATCGACATGGCCGATAGTGAAA TGCTGAATCTTTATGAAAGGGTACGGAAGCAGCTGCGCCAAAACGCCGAGGAAGATGGGAAGGGGTGTTTTGAAATCTATCATGCCTGTGA CGATTCCTGCATGGAATCTATTCGTAATAATACTTACGACCATAGCCAATATCGAGAGGAGGCATTGTTGAACAGGCTAAACATCAATCCGG TGACACTCAGTTCCGGATATAAAGATATAATCCTCTGGTTCTCTTTTGCCGCTTCTTGTTTCGTGCTGCTTGCTGTCGTAATGGGACTGTTCT TTTTTTGTCTCAAAAACGGCAACATGAGATGCACCATTTGCATA
3001	ATGTATAAAATCGTTGTGATCATTGCCCTGCTGGGCGCCGTAAAGGGCCTGGACAAAATATGCCTGGGGCACCATGCAGTAGTGAATGGCA CAATTGTCAAGACACTCACTAACGAACAGGAAGAAGTCACCAATGCCACGGAGACGGTGGAGAGTACCGGTCTGAACCGTCTGTGCATGAA GGGCCGCAACCATAAAGACTTGGGTAACTGTCATCCTATAGGTATGCTAATCGGTACTCCGGCCTGCGATCTGCACCTCACCGGTACCTGG GACACCCTGATCGAGCGTCAGAACGCGATCGCCTATTGCTATCCAGGCGCTACTGTAAATGAGGAAGCCCTGAGACAGAAGATCATGGAGT CCGGAGGGATCAATAAGATCTCTACTGGATTCACCTATGGCAGCAGTATTAACTCCGCCGGAACTACGCGGGCCTGCATGAGAAATGGCGG GAATAGCTTCTACGCTGAACTGAAATGGCTCGTTTCGAAGTCAAAAGGACAGAACTTCCCTCAGACAACGAACACATACCGCAACACCGACA CCGCGGAGCATCTAATCATGTGGGGTATCCATCACCCCAGCTCTACACAGGAAAAAAATGATCTTTATGGAACCCAGTCTTTGTCCATTTCT GTGGGCAGCTCCACCTACCAGAACAATTTCGTCCCAGTCGTGGGCGCGCGGCCACAGGTGAACGGACAGTCGGGGCGCATTGACTTTCAC TGGACCCTGGTGCAGCCCGGTGACAATATCACTTTTTCTCACAACGGCGGACTTATTGCTCCCTCCCGGGTGAGCAAACTGATCGGGCGG GGGCTGGGGATCCAGTCCGATGCACCTATCGATAACAATTGCGAGAGTAAATGCTTCTGGCGTGGGGGATCAATAAATACCAGGCTCCCAT TCCAAAATCTTTCACCCCGTACAGTTGGACAGTGCCCCAAATACGTGAACAAGCGCTCTCTTATGCTGGCAACAGGAATGAGAAACGTTCCA GAGCTAATGCAGGGTCGAGGGCTTTTCGGAGCTATTGCTGGATTTATCGAAAATGGCTGGGAGGGAATGGTCGATGGATGGTATGGATTCC GACACCAGAACGCTCAGGGGACCGGCCAGGCCGCGGACTACAAGAGCACACAGGCTGCAATTGATCAGATCACAGGGAAGTTAAATAGGC TCATCGAAAAGACGAACACCGAGTTCGAGTCAATCGAGTCCGAATTCTCTGAGATCGAACACCAAATTGGCAATGTCATCAATTGGACAAAG GATAGTATCACCGATATTTGGACGTACCAAGCAGAGCTTCTGGTTGCCATGGAGAACCAACACACCATCGATATGGCCGACAGCGAGATGC TTAATCTGTACGAGCGCGTGCGAAAGCAACTTAGGCAAAACGCCGAAGAAGATGGTAAGGGCTGCTTTGAGATTTATCACGCTTGCGATGA CTCCTGCATGGAGTCTATTAGAAATAATACCTACGATCATTCTCAGTATAGACAGGAGGCCCTGCTGAACCGGTTGAACATTAACCCAGTTA CACTGAGCTCCGGATACAAAGATATTATTCTGTGGTTTTCCTTTGGCGCGTCCTGCTTTGTGCTGTTAGCCGTGGTGATGGGGCTGGTATTC TTCTGTCTCAAGAACGGTAATATGCGTTGTACCATATGCATC
3002	ATGTATAAGATCGTGGTCATAATTGCACTCTTGGGGGCAGTGAAAGGGCTGGATAAAATTTGTCTGGGACACCACGCCGTTGCTAATGGCA CCATCGTGAAAACCCTTACCAACGAGCAGGAAGAGGTTACTAACGCAACCGAGACGGTCGAGTCCACTGGAATCAATCGTCTTTGCATGAA GGGCCGTAAACATAAGGATTTAGGAAATTGTCATCCCATTGGCATGCTGATCGGAACACCTGCCTGCGACCTGCACCTGACAGGGATGTGG GACACGCTGATCGAGAGGGAGAATGCAATTGCGTACTGTTATCCTGGAGCTACCGTGAATGTTGAAGCACTGCGTCAAAAAATCATGGAAT CCGGCGGGATCAACAAAATTAGCACTGGGTTCACATATGGAAGCAGCATAAACTCCGCCGGCACTACAAGGGCCTGCATGAGAAACGGCG GCAATAGTTTCTACGCTGAACTTAAGTGGCTTGTGTCAAAATCAAAGGGGCAAAACTTTCCTCAGACTACAAATACATATAGAAACACTGACA CCGCCGAGCACCTGATCATGTGGGGCATTCACCATCCTAGCTCTACCCAGGAGAAAAATGATTTATACGGCACTCAGAGTCTGTCTATTTCT GTGGGGTCGTCCACATATAGAAATAATTTCGTGCCAGTCGTGGGGGCTCGGCCGCAAGTGAATGGGCAGAGTGGTAGAATCGATTTCCACT GGACACTTGTGCAGCCCGGGGATAACATTACTTTCTCTCATAACGGCGGCCTGATCGCACCCTCCAGGGTGAGTAAACTAATAGGCAGGGG TCTGGGCATTCAGTCCGACGCTCCAATCGATAACAACTGCGAGTCCAAGTGCTTCTGGCGCGGCGGTTCCATCAACACCCGGTTACCTTTT CAGAACCTGTCCCCCCGCACCGTTGGCCAGTGCCCTAAATACGTAAATCGAAGATCATTAATGCTGGCGACCGGTATGAGGAACGTACCTG AGCTGATTCAGGGGAGGGGGTTATTTGGTGCCATTGCCGGGTTTCTCGAAAACGGGTGGGAGGGAATGGTTGATGGCTGGTACGGGTTCA GGCACCAGAATGCCCAGGGCACAGGACAGGCTGCCGATTACAAGAGCACTCAGGCCGCTATTGACCAGATCACCGGCAAGCTGAATCGGC TGGTAGAGAAAACCAATACGGAGTTTGAGTCCATTGAGTCCGAGTTCTCAGAGATCGAGCACCAGATTGGCAATGTCATCAATTGGACCAAG GATTCAATTACGGATATATGGACATACCAGGCTGAGCTGCTTGTAGCAATGGAGAACCAACATACCATAGACATGGCTGATTCTGAGATGCT GAACCTCTACGAAAGCGTCAGAAAACAACTCAGACAAAACGCAGAAGAAGACGGGAAAGGGTGTTTCGAAATTTACCATGCTTGTGATGAC TCCTGCATGGAGAGCATCAGAAACAACACTTATGACCACAGCCAGTACCGTGAAGAGGCATTGTTGAACCGACTCAACATCAACCCTGTTAC GCTGTCATCTGGGTATAAAGACATTATTCTATGGTTCTCTTTTGGCGCTTCATGTTTTGTTCTGTTGGCGGTAGTGATGGGTCTGTTCTTTTTC TGTTTAAAAAATGGCAATATGAGGTGCACCATCTGCATA

3003	ATGTACAAGATCGTGGTTATTATCGCTCTTCTGGGTGCCGTTAAAGGGCTGGACAAAATTTGTCTGGGGCATCACGCCGTGGCGAATGGTA CTATCGTCAAGACATTGACCAACGAGCAGGAGGAGGTTACAAATGCAACCGAGACCGTTGAGAGTACTGGAATCAACCGGCTGTGCATGAA GGGAAGAAAGCACAAGGATCTGGGGAATTGTCACCCTATCGGAATGTTAATTGGTACACCTGCTTGTGACCTGCACCTGACAGGCATGTGG GACACCCTGATCGAACGGGAGAACGCCATTGCCTATTGTTATCCAGGGGCCACTGTTAATGTCGAGGCACTCAGGCAGAAGATTATGGAAT CCGGTGCAATTAATAAAATTAGCACAGGGTTTACTTACGGTTCCTCAATAAATTCAGCCGGCACTACACGGGCGTGTATGAGGAATGGAGGA AACTCCTTCTATGCCGAACTGAAATGGCTCGTCAGCAAATCCAAGGGTCAAAATTTTCCTCAAACAACTAACACCTATAGAAATACCGATACC GCAGAGCACCTCATAATGTGGGGGATACACCATCCGAGCTCCACACAAGAAAAAAACGACCTGTATGGCACTCAAAGTCTTAGCATTTCCGT CGGAAGTTCTACATACCGGAATAACTTTGTGCCGGTGGTAGGTGCCCGACCCCAAGTCAACGGCCAGAGTGGCCGCATCGATTTCCATTGG ACACTGGTGCAGCCAGGAGATAACATTACCTTTTCTCACAATGGCGGCTTGATCGCTCCTTCGAGGGTGAGCAAGCTGATTGGCCGGGGTC TGGGCATTCAATCCGACGCGCCTATCGACAATAATTGTGAGTCTAAATGCTTTTGGAGAGGAGGGTCAATCAACACAAGACTACCCTTCCAG AATCTTTCCCCCCGAACAGTGGGACAGTGCCCCAAATATGTGAATCGGCGCTCTCTTATGCTGGCCACTGGGATGAGAAACGTGCCTGAGT TAATACAGGGCCGGGGACTTTTCGGAGCAATCGCCGGATTCCTAGAGAACGGTTGGGAGGGGATGGTCGACGGATGGTACGGGTTCAGAC ACCAAAATGCTCAGGGGACAGGGCAGGCAGCAGACTATAAATCAACGCAGGCCGCTATCGACCAGATCACCGGTAAACTGAACCGGCTTG TCGAAAAGACCAACACTGAGTTCGAGAGTATTGAATCCGAGTTTAGCGAGATCGAACACCAGATAGGCAATGTTATCAATTGGACAAAGGAT TCAATCACGGACATATGGACATACCAAGCAGAACTGCTGGTAGCTATGGAAAATCAACATACAATCGATATGGCCGACTCAGAGATGCTCAA CCTCTACGAAAGAGTAAGGAAACAACTCAGACAAAATGCCGAAGAGGACGGCAAGGGTTGCTTTGAAATATATCATGCGTGCGACGACAGC TGCATGGAGAGCATACGGAACAATACGTACGACCATTCCCAGTATCGCGAAGAAGCCTTACTGAACCGCTTGAACATTAACCCCGTGACCC TTTCCTCCGGATATAAAGATATCATTCTCTGGTTCAGTTTTGGAGCCTCATGCTTTGTCCTTTTGGCTGTGGTAATGGGATTATTTTTCTTCTG TCTCAAGAATGGTAACATGCGATGTACAATTTGCATC
3004	ATGTACAAGATCGTGGTGATCATAGCACTGCTGGGCGCCGTGAAAGGCTTAGACAAAATCTGCTTAGGGCATCATGCCGTCGCTAATGGGA CTATAGTCAAGACATTGACCAACGAGCAAGAAGAGGTGACAAACGCAACTGAAACTGTTGAAAGTACCGGGATAAACCGGTTGTGCATGAA AGGGCGAAAGCACAAGGACCTTGGCAACTGTCATCCCATCGGCATGCTTATAGGAACACCTGCTTGCGATCTCCATCTGACTGGCATGTGG GACACCCTGATAGAACGAGAAAATGCCATTGCCTACTGTTACCCAGGCGCCACCGTTAATGTTGAGGCCCTCAGACAGAAGATAATGGAAT CTGGAGGCATCAACAAAATCTCTACGGGCTTTACCTACGGTTCTAGCATTAACTCTGCCGGGACTACCCGAGCTTGTATGCGCAACGGGGG CAACTCTTTCTACGCCGAACTTAAGTGGCTGGTGAGCAAGAGCAAGGGACAGAATTTCCCTCAAACTACTAACACTTATAGGAACACAGACA CCGCCGAACATCTAATAATGTGGGGAATACACCATCCCTCATCCACACAGGAAAAAAACGACCTTTATGGGACCCAGTCTCTGAGCATTTCA GTCGGCTCATCCACTTATCGTAACAACTTTGTGCCCGTGGTTGGGGCTCGGCCCCAGGTGAATGGCCAGTCTGGGCGTATTGATTTCCACT GGACCCTAGTGCAACCCGGAGACAACATCACCTTTAGTCATAATGGTGGCCTTATAGCCCCTAGCAGGGTGTCAAAACTTATTGGGAGAGG CCTCGGGATTCAGAGTGACGCACCCATCGACAACAACTGCGAGAGTAAGTGTTTTTGGAGAGGGGGTTCGATCAACACCCGCCTCCCCTTC CAGAACCTATCGCCACGCACTGTTGGTCAGTGCCCAAAATACGTCAACAGGAGGTCACTTATGCTGGCTACAGGGATGCGGAACGTGCCG GAGCTGATACAGGGACGCGGTCTGTTCGGAGCCATTGCCGGATTCTTGGAGAATGGCTGGGAGGGGATGGTGGACGGCTGGTACGGCTT CCGACACCAGAATGCCCAGGGTACAGGTCAGGCAGCGGACTATAAGAGTACTCAAGCCGCTATCGATCAAATCACTGGGAAGCTCAACCG ACTGGTCGAGAAGACAAACACAGAATTCGAGTCTATCGAGTCAGAATTCTCTGAAATCGAGCACCAGATTGGTAACGTAATCAATTGGACCA AGGATAGTATTACCGACATCTGGACATACCAAGCAGAACTGCTCGTGGCAATGGAAAATCAGCACACAATAGACATGGCCGATAGTGAGAT GCTCAACCTGTACGAACGCGTTCGAAAACAGCTGAGACAGAACGCTGAAGAAGACGGGAAAGGGTGCTTCGAGATCTATCACGCCTGCGA CGATTCATGCATCGAGAGCATTAGAAACAATACCTACGACCACAGCCAGTATAGGGAGGAGGCTCTGCTAAACAGGCTCAACATTAATCCC GTAACACTCTCAAGCGGATATAAAGACATAATTCTGTGGTTTTCCTTCGGAGCTTCCTGCTTCGTCCTGCTGGCGGTAGTCATGGGGTTATT CTTTTTTTGTCTGAAGAACGGGAATATGCGATGTACAATATGCATT
3005	ATGTACAAGGTTGTGGTTATTATAGCACTCCTCGGGGCTGTGCGCGGACTGGATAAGATCTGCCTGGGACATCATGCCGTGGCAAATGGTA CCACTGTAAAAACTCTCACCAATGAGCAGGAAGAGGTGACCAACGCCACTGAGACAGTGGAATCTACAAGCCTGAACAAGCTGTGTATGAA GGGCAGACGATATAAGGATCTGGGTAATTGCCATCCCATCGCATGTTGATCGGAACACCAGTTTGCGATCTGCATCTAACTGGAACCTGG GACACGCTTATCGAGCGGGAAAACGCCACCGCGTACTGCTACCCAGGAGTCACCATCAACGAAGAAGCACTGAGACAGAAGATCATGGAG AGTGGGGGGATCTCCAAGATGCGCACCGGGTTCACTTATGGGCCCAGCATTAACAGCGCTGGGACAACCAGATCCTGCATGAGGAATGGA GGGAACTCTTTTTATGCCGAATTAAAATGGCTGGTTTCAGGAACAAAAGGACAAAATTTCCCGCAGACCACCAATACATATAGAAACACCGAT ACAGCAGAACACTTAATCATCTGGGGGATCCATCACCCTTCATCTACGCAGGAAAAGAATGATCTGTATGGTACCCAGTCTTTGAGTATTAG TGTAGGCTCTTCCACGTACCAGAATAACTTTGTCCCTGTTATAGGTGCTCGACCTCAGGTCAATGGGCAGTCTGGGCGTATCGAGTTCCATT GGACCCTCGTACGACCCGGCGACAACATCACATTTAGCCACAACGGCGGATTGATAGCCCCCGACAGAGTGTCCAAGTTAATCGGCAAAG GAATTGGTATCCAGTCCGGCGCCGTGATCGATAAGGATTGTGAGTCAAAATGCTTCTGGAGAGGCGGCAGCATCATAACAGAGCTTCCGTT TCAAAACCTGAGTCCCCGCACCGTTGGGCAATGTCCTAAGTACGTGAAAAAACGAAGCCTCCTGCTGGCAACCGGCATGCGGAATGTTCCC GAGGTGGTCCAGGGAAGAGGCCTGTTTGGCGCGATTGCTGGGTTCATCGAGAATGGCTGGGAAGGCATGGTTGACGGATGGTATGGGTT CAGGCATCAGAACGCGCAGGGCATCGGGCAGGCTGCTGACTACAAGAGCACTCAGACCGCCATCGATCAGATCACAGGGAAGCTAAATCG GCTGATTGAAAAGACTAACACGGAGTTTGAGAGCATTGAAAGCGAATTTAGTGAAATCGAACACCAGATAGGAAACGTTATTAACTGGACAA AGGATAGTATTACCGACATATGGACGTATCAAGCAGAGCTGCTGGTCGCCATGGAGAACCAGCATACTATAGATATGGCAGACAGCGAAAT GCTCAATTTGTACGAGAGGGTAAGAAAACAACTGAGACAAAATGCTGAAGAGGATGGCAAAGGCTGCTTTGAAATATACCACACCTGTGACA ATTCCTGTATGGAGTCCATTCGGAACAATACGTATGATCATTCCCAGTATCGCGAGGAAGCCCTACTCAACCGCCTCAATATCAACCCCGTT AAGCTATCATCAGGCTACAAGGACATTATCCTGTGGTTCAGCTTCGGAGCCTCCTGTTTCGTCGCTACTTGCTGTGATCATGGGTTTGGGGTT CTTTTGCCTGAAAAATGGAAATATGAGATGCACAATTTGTATT

3006	ATGTATAAGATAGTCGTGATAATCGCCCTTTTGGGAGCGGTGAAGGGACTGGACAAGATTTGTCTCGGCCATCATGCAGTCGCCAACGGTA CAATTGTTAAAACCCTCACCAACGAACAAGAAAAAGTGACAAATGCCACAGAAACTGTGGAGAGCACCGGCCTTAACCGGTTGTGCATGAA GGGGCGAAAGCATAAGGACTTGGGTAATTGTCATCCAATTGGGATGCTGATTGGGACTCCCGCCTGCGATCTGCACCTGACCGGCACATG GGATACCATTATCGAGAGGGAGAACGCGATCGCTTACTGCTATCCTGGGGCTACAGTCAATGAGGAGGCCCTAAGGCAGAAAATAATGGAG TCAGGGGCAATCGATAAAATTTCTACAGGCTTCACTTACGGATCTTCTATTAACAGCGCAGGCACCACGCGCGCCTGCATGAGAAATGGCG GGAATAGTTTTTATGCCGAGCTGAAATGGCTGGTGTCCAAGAGCAAGGGACAGAACTTTCCGCAAACCACTAATACATACCGTAATACTGAC ACAGCAGAACATCTGATTATGTGGGGAATCCACCACCCCTCCAGCACTCAAGAGAAAAATGATCTTTACGGAACACAGTCCCTCAGCATAAG CGTGGGTTCGAGTACATATAGGAACAATTTCGTGCCAGTGGTAGGCGCGCGACCGCAGGTCAATGGACAGTCAGGAAGGATCGACTTCCA TTGGACTCTAGTGCAACCTGGGGATAATATTACCTTTAGCCATAATGGAGGTCTGATAGCCCCATCTCGCGTCTCCAAGCTGATTGGGCGC GGTCTGGGGATTCAGAGTGACGCTCCTATTGATAACAACTGCGAGAGTAAATGTTTCTGGCGCGGCGGCTCGATTAACACCCGTCTTCCCT TTCAGAACCTGAGTCCCCGGACAGTAGGACAGTGCCCTAAATACGTGAACAAAAGAAGTCTTATGCTCGCCACTGGGATGCGAAACGTTCC GGAACTGATCCAGGGTCGCGGCCTGTTCGGCGCAATTGCTGGTTTCCTCGAAAATGGATGGGAAGGTATGGTGGATGGGTGGTACGGGTT CAGACATCAAAACGCTCAGGGCACGGGCCAGGCTGCCGACTACAAGAGCACACAGGCTGCCATCGACCAGATTACTGGGAAACTCAATAG ACTCGTGGAAAAAACTAATACGGAATTCGAAAGCATCGAATCTGAGTTCAGTGAGATCGAACATCAGATCGGAAACGTGATCAACTGGACAA AGGATTCGATTACTGATATTTGGACGTACCAGGCAGAGTTGTTGGTCGCCATGGAGAATCAACACACAATCGATATGGCCGACTCAGAGAT GTTGAATCTTTACGAGAGGGTGCGAAAGCAGCTGAGGCAGAATGCCGAGGAAGATGGAAAGGGGTGTTTTGAGATCTACCACGCCTGCGA CGATTCTTGCATGGAATCCATCCGCAATAATACATACGATCACTCACAGTATAGGGAGGAGGCCCTACTCAACAGACTGAACATCAATCCCG TCACACTTTCAAGTGGTTACAAGGATATCATACTTTGGTTCAGCTTCGGCGCCCAGTTGCTTTGTTCTGCTTGCCGTTGTGATGGGCCTGGTG TTCTTTTGCCTGAAGAATGGTAACATGAGATGTACCATCTGCATT
3007	ATGTATAAAATTGTCGTTATCATTGCCTTGCTCGGCGCTGTGAAGGGGCTGGATAAAATTTGTCTGGGACATCACGCCGTGGCGAACGGGA CTATCGTCAAGACGCTTACTAACGAACAGGAAGAGGTGACTAACGCGACGGAAACCGTGGAGTCAACTGGGATTAATCGCTTGTGCATGAA GGGCAGAAAGCACAAGGACTTAGGCAACTGTCATCCTATCGGCATGCTGATCGGTACCCCAGCCTGCGACCTTCACTTGACTGGCACCTG GGATACCCTGATCGAAAGAGAGAACGCAATAGCATACTGTTATCCAGGCGCCACAGCCAACGTAGAGGCCCTGAGACAGAAGATCATGGAA TCTGGAGGTATCGATAAAATTAGCACCGGATTTACATATGGATCAAGTATAAACTCGGCAGGAACGACGCGGGCGTGCATGCGAAATGGCG GAAATTCCTTCTACGCCGAATTGAAATGGCTAGTGTCCAAAAGCAAAGGCCAAAACTTTCCACAGACTACTAACACCTACAGAAATACTGACA CCGCTGAACATTTGATTATGTGGGGGATCCACCACCCAAGTAGCATTCAGGAGAAGAACGATCTGTATGGCACTCAGTCACTAAGTATTTCC GTGGGCAGCTCTACCTACCGCAATAACTTCGTGCCGGTTGTGGGCGCCAGACCTCAGGTGAACGGACAAAGTGGTCGCATTGATTTCCACT GGACACTTGTGCAACCTGGGGACAATATCACTTTCAGCCACAACGGGGGGCTGATCGCCCCATCAAGAGTGAGCAAATTGATAGGGAGAG GCTTGGGAATCCAGTCAGATGCACCTATCGATAATAATTGTGAAAGCAAATGCTTTTGGCGTGGGGGATCCATAAACACTCGACTGCCATTC CAGAATTTGTCCCCCCGCACCGTTGGCCAGTGTCCCAAGTACGTTAATCGCAGAAGTCTGATGCTGGCCACCGGCATGAGAAACGTGCCC GAACTGATACAGGGACGCGGGCTCTTTGGAGCAATTGCCGGCTTTCTGGAGAACGGATGGGAGGGAATGGTGGACGGATGGTACGGCTTT CGGCACCAGAATGCCCAGGGGACAGGACAAGCCGCCGATTATAAGTCAACACAGGCCGCAATCGACCAGATTACCGGCAAGCTCAACAGG CTCGTGGAAAAGACAAACACTGAGTTTGAGTCAATCGAGTCAGAATTTTCCGAGATCGAGCATCAAATCGGCAACGTGATCAATTGGACTAT GGATTCTATCACCGACATTTGGACTTACCAGGCAGAGCTCCTGGTGGCAATGGAGAACCAACACACTATCGACATGGCAGATTCAGAGATG CTCAACCTCTATGAAAGGGTACGAAAACAGCTTCGGCAGAACGCCGAGGAGGACGGTAAGGGGTGTTTTGAAATCTATCATGCGTGCGACG ATAGCTGTATGGAAAGTATCCGGAATAATACATACGATCACTCCCAGTACCGCGAAGACGCTCTCTTGAACCGGCTTAACATCAACCCGGTA ACTCTCAGTAGCGGGTACAAGGACATCATTCTGTGGTTTTCTTTTGGTGCCTCCTGTTTTGTGCTGCTGGCCGTGGTGATGGGACTCGTGTT TTTTTGCCTCAAGAACGGGAATATGCGGTGCACAATATGCATT
3008	ATGTATAAGATTGTGGTGATTATTGCTCTCTTGGGGGCAGTCAAAGGACTTGATAAGATTTGTCTGGGCCACCATGCTGTCGCAAATGGGAC AATCGTCAAGACCCTTACCAACGAGCAAGAAAAGGTAACCAACGCAACCGAGACTGTTGAGAGTACAGGGCTGAATAGACTCTGTATGAAG GGCCGCAAACATAAGGACCTAGGTAACTGTCATCCAATAGGGATGCTGATTGGAACACCGGCATGTGACTTACACCTCACAGGTACATGGG ATACTATTATAGAGAGAGAAAATGCCATTGCCTACTGTTATCCTGGGGCTACCGTTAACGAGGAAGCCTTACGTCAGAAAATTATGGAATCC GGGGGCATTGATAAAATATCCACTGGATTTACTTATGGAAGCTCAATCAACTCCGCGGGCACTACTCGAGCATGTATGCGCAACGGAGGTA ACTCCTTCTACGCCGAACTGAAGTGGCTCGTTTCAAAATCCAAAGGACAGAACTTTCCGCAGACCACTAATACCTACCGCAATACAGACACA GCAGAACATCTGATCATGTGGGGCATTCATCACCCCAGCTCTACTCAAGAAAAGAATGACCTGTACGGGACTCAAAGTCTGTCCATTTCTGT TGGGTCTAGTACCTACCGGAATAACTTTGTCCCCGTTGTTGGAGCACGCCCTCAGGTTAACGGTCAGAGTGGAAGGATCGACTTTCACTGG ACCCTCGTCCAGCCTGGCGACAATATCACATTTAGTCACAACGGGGGCCTAATCGCTCCCTCTCGGGTGAGCAAGCTGATAGGTCGGGGC TTGGGTATACAGTCTGACGCCCCAATCGATAACAACTGTGAAAGCAAGTGCTTCTGGCGGGGTGGCTCTATCAACACCCGACTGCCATTTC AAAATCTCAGTCCAAGGACAGTGGGGCAATGCCCAAAATACGTCAACAAAAGGTCACTCATGCTCGCCACAGGAATGCGCAATGTGCCTGA ACTGATTCAAGGACGAGGACTGTTTGGCGCTATTGCCGGGTTTTTAGAGAATGGCTGGGAAGGCATGGTAGACGGGTGGTACGGGTTCCG GCATCAGAATGCCCAAGGGACGGGTCAGGCCGCTGACTATAAGTCCACTCAGGCCGCAATAGATCAGATCACCGGAAAGTTGAACAGACT AGTGGAGAAGACAAACACTGAGTTCGAGTCCATTGAGAGCGAATTCAGCGAGATAGAACACCAAATCGGCAATGTGATTAACTGGACCAAA GACAGTATTACTGATATCTGGACTTATCAAGCCGAACTGTTGGTGGCAATGGAGAACCAACACACGATCGACATGGCAGACAGTGAGATGC TCAATCTGTACGAGAGGGTGCGTAAACAGCTGAGGCAGAACGCCGAACAAGACGGAAAGGGGTGTTTCGAGATATACCACGCCTGTGATG ACTCCTGTATGGAAAGCATCAGGAACAACACTTATGACCATAGTCAATATAGGGAAGAAGCACTGCTCAACCGCCTGAACATCAATCCTGTA ACACTTTCATCAGGATATAAGGACATCATCCTCTGGTTTAGTTTCGGAGCTTCATGCTTTGTGCTGCTCGCGGTAGTCATGGGGCTGGTCTT CTTTTGTTTGAAGAACGGTAACATGAGATGCACCATTTGTATC

3009	ATGTACAAAATCGTCGTCATCATCGCCTTACTGGGAGCCGTCAAAGGACTTGATAAAATTTGTCTTGGTCACCACGCTGTTGCCAATGGAAC AATCGTGAAGACACTTACCAATGAGCAGGAGGAGGTGACTAATGCAACAGAGACAGTGGAGTCCACTGGAATAAATCGGCTGTGCATGAAG GGCAGAAAACACAAGGATTTGGGCAACTGCCACCCTATTGGAATGCTCATAGGCACCCCTGCGTGTGACCTTCATCTGACTGGCACTTGGG ACACTCTCATCGAACGCGAGAACGCCATCGCCTACTGCTATCCTGGAGCCACCGCCAACGTGGAGGCTCTACGGCAGAAGATTATGGAGA GCGGAGGGATTGACAAGATCTCCACGGGTTTCACTTACGGCAGCTCCATCAACAGTGCCGGTACAACTCGCGCGTGCATGAGAAACGGGG GCAACAGCTTTTATGCCGAGCTGAAGTGGCTGGTGAGTAAGTCAAAAGGGCAGAACTTTCCACAGACAACAAATACGTACAGAAATACAGAT ACAGCCGAGCACCTGATCATGTGGGGGATTCATCATCCAAGTTCTATCCAGGAGAAAAACGACTTGTATGGGACTCAGAGTCTCAGTATCA GCGTAGGTTCCTCTACCTACAGGAACAACTTCGTGCCCGTCGTCGGGGCCCGTCCCCAAGTCAACGGCCAGAGTGGACGGATTGATTTTC ACTGGACTCTTGTGCAGCCCGGCGACAACATTACCTTCAGTCACAACGGAGGGCTGATTGCACCTTCCAGAGTCTCGAAGCTGATCGGACG TGGATTGGGGATTCAGTCGGACGCTCCTATCGACAATAATTGCGAGAGCAAATGTTTCTGGCGCGGGGGTTCTATTAACACCAGGCTCCCC TTTCAAAATTTGAGCCCGCGAACCGTCGGCCAATGTCCCAAGTATGTCAACCGGAGAAGCCTGATGTTGGCTACAGGAATGCGCAATGTGC CGGAGCTCATCCAGGGCCGTGGCTTGTTCGGTGCAATTGCCGGATTCCTAGAAAATGGATGGGAGGGAATGGTTGATGGGTGGTACGGCT TCCGCCACCAAAACGCTCAGGGCACTGGACAGGCCGCGGACTATAAGTCTACTCAGGCGGCTATCGATCAGATCACTGGGAAACTGAATA GGCTTGTGGAAAAAACCAATACGGAGTTCGAAAGTATCGAGTCAGAGTTTAGCGAAATCGAGCATCAGATTGGTAACGTGATCAACTGGAC CATGGATAGCATTACCGACATCTGGACCTACCAGGCCGAATTGCTGGTGGCCATGGAGAATCAGCACACAATTGACATGGCAGACTCTGAA ATGCTTAACCTATATGAGCGCGTACGGAAACAGCTCAGACAGAACGCGGAGGAAGATGGAAAGGGCTGCTTCGAAATTTATCACGCTTGCG ACGACTCCTGTATGGAGTCTATCAGAAACAATACGTATGATCATTCGCAATACCGGGAGGAAGCACTCCTGAATAGACTCAACATCAACCCC GTTACACTGAGCTCAGGGTACAAAGACATTATTCTCTGGTTTTCCTTCGGAGCTTCTTGCTTCGTACTGTTAGCTGTCGTCATGCGCCTAGTT TTTTTTTGTCTTAAAAACGGCAACATGAGATGCACCATCTGTATA
3010	ATGTATAAAATCGTGGTAATCATTGCACTGCTGGGAGCAGTCAAAGGGCTCGATAAAATATGCCTTGGCCACCACGCCGTAGCAAATGGTAC AATCGTGAAGACCCTCACGAACGAACAGGAGGAAGTCACCAACGCTACAGAAACTGTCGAAAGTACCGGGATCAACCGGCTCTGTATGAAG GGCAGGAAGCATAAAGACTTAGGTAACTGCCATCCTATCGGCATGCTGATTGGCACACCTGCCTGCGACTTGCACTTAACGGGAATGTGGG ACACATTGATTGAGAGAGAGAATGCTATTGCTTACTGTTACCCAGGAGCGACTGTGAACGTGGAGGCACTACGGCAAAAAATAATGGAGTC CGGGGGAATCAATAAGATCTCGACAGGATTTACCTATGGGAGCTCAATCAATTCTGCGGGAACAACCCGGGCCTGCATGCGCAATGGCGG GAATAGCTTCTATGCCGAACTCAAGTGGCTCGTTAGCAAGAGCAAAGGGCAGAATTTCCCGCAAACTACTAACACATACGGAACACAGACA CAGCAGAGCATTTGATTATGTGGGGGATTCACCACCCCTCTAGTACGCAGGAAAAGAACGATCTTTACGGGACCCAGTCCCTTTCTATCAGC GTAGGAAGCAGCACTTACAGGAATAATTTTGTCCCTGTGGTTGGCGCGGGCCCTCAGGTTAATGGGCAATCTGGCCGCATTGATTTTCATT GGACCCTCGTACAGCCTGGCGACAATATAACATTCTCACATAATGGCGGCCTGATCGCACCTTCCCGGGTATCTAAATTGATAGGTAGAGG CCTGGGGATTCAAAGCGATGCTCCAATAGATAATAACTGTGAATCAAAGTGCTTTTGGCGGGGAGGGTCAATTAATACGCGGCTACCCTTTC AGAATCTTAGCCCACGCACCGTTGGCCAATGTCCAAAGTACGTGAATCGTAGATCACTGATGCTGGCTACAGGGATGCGGAATGTGCCAGA ACTCATTCAGGGCCGTGGGCTATTCGGCGCTATCGCCGGGTTTCTGGAAAACGGCTGGGAGGGGATGGTCGATGGCTGGTACGGCTTCC GGCACCAGAACGCTCAGGGTACAGGGCAGGCGGCTGATTATAAAAGCACACAGGCTGCAATCGACCAAATTACTGGAAAACTAAATAGGCT AGTGGAAAAAACCAACACCGAATTCGAAAGTATTGAAAGTGAGTTCAGTGAAATTGAGCACCAGATTGGAAACGTGATCAACTGGACTAAGG ACAGTATCACCGATATTTGGACATATCAGGCAGAACTTCTCGTGGCCATGGAAAACCAGCATACTATTGACATGGCTGACAGTGAAATGTTA AATCTCTATGAGCGCGTACGCAAACAATTGCGGCAGAATGCCGAGGAAGATGGGAAGGGCTGCTTCGAAATCTATCACGCTTGCGACGATT CCTGCATGGAGTCCATCAGGAACAATACATACGATCACAGCCAGTATCGCCGAAGAAGCCCTGCTTAATACACTAAATATTAATCCCGTCACC CTGTCCTCAGGCTATAAGGACATCATCCTTTGGTTTTCATTTGGCGCAAGTTGCTTCGTCTTGCTCGCAGTCGTCATGGGGCTGTTTTTTTTC TGTCTCAAAAACGGAAACATGAGGTGTACCATTTGCATT
3011	ATGTACAAAATTGTGGTTATCATCGCTTTGCTTGGCGCTGTGAAGGGGCTGGATAAGATCTGTCTCGGGCATCACGCTGTCGCCAATGGCA CAATTGTCAAGACATTGACGAACGAGCAGGAGGAAGTGACAAATGCCACTGAAACAGTGGAATCCACTGGGATCAACCGATTGTGCATGAA AGGTAGGAAGCATAAGGATCTCGGTAACTGTCATCCAATCGGCATGTTGATTGGAACCCCGGCCTGTGATCTTCACCTCACTGGTATGTGG GACACACTGATTGAAAGAGAAAACGCTATTGCCTACTGTTACCCAGGTGCAACGGTTAATGTCGAGGCACTGAGACAGAAAATTATGGAAAG CGGAGGCATCAACAAGATATCAACCGGGTTTACCTATGGCTCATCGATTAATAGCGCAGGCACCACTCGCGCGTGTATGCGAAATGGAGGA AATTCATTTTATGCCGAGTTAAAATGGCTGGTTAGTAAGAGCAAGGGTCAGAACTTCCCTCAGACCACAAATACGTATCGCAATACGGACAC TGCAGAACACCTTATCATGTGGGGTATTCACCACCCTAGCAGTACTCAAGAAAAAAATGACCTGTATGGTACCCAGAGTCTGAGTATAAGCG TGGGTTCCAGTACTTACCGGAACAACTTCGTTCCAGTTGTGGGGGCCCGCCCACAGGTTAACGGGCAATCGGGGCGCATCGATTTTCACTG GACCTTGGTGCAACCTGGAGACAATATTACTTTCAGTCACAACGGCGGATTGATAGCTCCATCCCGTGTCAGTAAATTGATTGGGCGCGGA CTGGGAATCCAGTCTGACGCACCCATAGATAACAACTGTGAGTCTAAGTGTTTCTGGAGGGGCGGTTCCATTAATACTCGGCTGCCATTTCA GAATCTTAGCCCGCGCACTGTGGGTCAATGTCCCAAGTACGTGAACAGGCGTTCTCTAATGCTTGCCACCGGGATGAGAAACGTTCCTGAA CTGATACAAGGTAGGGGACTGTTCGGTGCAATCGCTGGCTTTTTGGAGAATGGCTGGGAGGGAATGGTGGACGGCTGGTATGGATTCCGC CATCAGAACGCCCAGGGAACCGGACAGGCTGCCGACTATAAGTCCACGCAGGCCGCGATTGACCAAATTACCGGAAAACTCAATCGCCTA GTCGAGAAAACAAACACCGAATTCGAGAGTATCGAGTCCGAATTTTCTGAGATCGAGCACCAAATCGGCAACGTAATCAATTGGACGAAGG ATTCCATTACTGACATCTGGACCTACCAAGCCGAATTGTTGGTCGCAATGGAGAACCAGCACACCATAGACATGGCTGACAGCGAAATGCT GAACCTGTACGAGCGGGTTCGGAAACAGCTGCGCCACAACGCAGAGGAGGATGGCAAAGGCTGCTTCGAAATCTATCACGCCTGCGACGA TAGCTGCATGGAATCAATAAGGAATAACACTTATGATCACAGCCAGTATCGGGAAGAGGCCCTGCTGAATCGGCTCAACATCAATCCCGTGA CTCTCTCTTCCGGCTATAAAGACATTATCTTGTGGTTCTCTTTCGGCGCGTCATGCTTCGTGCTGCTGGCAGTGGTAATGGGACTCTTCTTTT TTTGCCTGAAAAACGGGAACATGCGCTGCACAATCTGCATC

3012	ATGTATAAGATTGTGGTGATTATTGCCCTGCTCGGCGCCGTGAAGGGGTTGGACAAGATTTGTCTTGGTCACCACGCCGTGGCTAACGGGA CCATAGTCAAGACTCTGACCAATGAACAGGAAGAGGTAACAAACGCAACCGAAACAGTCGAGTCTACAGGAATCAATCGGTTATGTATGAAA GGCCGAAAACATAAAGACCTCGGCAATTGTCACCCCATAGGGATGCTGATCGGGACTCCAGCTTGCGACCTGCATTTAACAGGCATGTGGG ACACTCTCATCGAGAGAGAGAACGCGATCGCATACTGCTACCCAGGAGCTACCGTGAATGTAGAGGCATTGAGACAGAAAATCATGGAAAG TGCCGCGAATAAATAAGATTTCTACAGGGTTTACGTACGGCAGCTCCATTAACAGTGCCGGAACAACCCGAGCATGTATGAGAAATGGAGGT AATAGCTTTTACGCCGAACTGAAGTGGCTTGTGTCCAAAAGTAAGGGGCAAAATTTTCCTCAGACAACGAACACCTACCGGAACACCGACAC AGCTGAGCACCTGATTATGTGGGGGATACATCATCCTAGCAGCACGCAGGAGAAGAATGATCTATACGGGACCCAATCTCTCAGCATCTCA GTGGGCTCCTCAACTTACCGCAACAACTTCGTGCCTGTCGTCGGCGCCCGCCCTCAGGTCAACGGTCAGTCTGGCCGAATCGACTTTCACT GGACACTGGTGCAGCCTGGAGACAACATCACTTTCTCCCATAACGGGGGACTTATCGCACCGAGCAGAGTGAGCAAGCTGATTGGAAGAG GACTCGGAATACAATCCGATGCCCCGATAGATAATAATTGTGAGAGCAAGTGTTTCTGGCGCGGCGGATCCATCAATACACGCTTACCGTTT CAGAACTTGAGCCCACGGACTGTTGGGCAGTGCCCAAAGTACGTGAACCGCCGCAGTTTAATGCTTGCTACCGGTATGAGAAATGTACCTG AACTCATTCAAGGCAGGGGCCTGTTTGGCGCAATCGCTGGCTTTCTCGAAAATGGATGGGAAGGCATGGTAGACGGTTGGTACGGATTCAG ACACCAGAATGCCCAGGGCACGGGTCAAGCAGCCGATTATAAAAGCACACAAGCAGCAATTGATCAGATTACAGGTAAGCTCAACCGTCTC GTGGAAAAAACTAACACCGAGTTCGAATCCATTGAGAGTGAGTTCAGTGAGATAGAGCACCAGATCGGAAATGTGATTAACTGGACAAAAGA TAGCATTACAGATATCTGGACGTACCAGGCTGAGCTCTTGGTGGCGATGGAAAATCAGCATACCATTGACATGGCCGACTCAGAGATGCTT AACCTGTATGAGAGAGTGCGCAAACAGCTTAGACAGAACGCCGAAGAGGATGGCAAGGGATGTTTCGAGATCTACCACGCTTGTGATGACA GCTGTATGGAGAGTATTAGAAACAACACGTATGACCATTCCCAGTACCGAGAGGAGGCTTTGCTTAATCGTTTGAATATTAACCCCGTCACG CTCAGCAGCGGGTACAAACGATATCATACTCTGGTTCTCTTTCGGAGCATCATGCTTCGTACTTTTGGCTGTGGTGATGGGGCTGTTTTTCTT CTGCTTGAAAAACGGCAACATGAGGTGCACTATCTGTATC
3013	ATGTACAAAATCGTTGTGATCATTGCTCTGTTGGGCGCGGTAAAGGGCCTAGATAAGATTTGTCTGGGCCATCACGCCGTTGCCAACGGCA CTATAGTTAAAACGCTGACAAATGAACAGGAAAAGGTAACTAACGCCACCGAGACTGTGGAGTCCACTGGTCTGAACCGACTCTGCATGAA AGGGCGAAAACATAAAGACCTGGGCAACTGTCACCCAATTGGGATGCTGATCGGGACACCTGCCTGCGATCTCCACCTGACCGGTACATG GGACACAATCATTGAAAGGGAAAACGCCATCGCCTACTGTTATCCAGGCGCTACAGTCAACGAGGAAGCATTGCGGCAGAAAATTATGGAG TCTGGCGGAATCGATAAGATATCCACTGGCTTCACCTACGGATCGTCTATTAACTCTGCGGGTACGACACGCGCCTGTATGAGGAACGGCG GAAATAGCTTTTACGCTGAACTAAAGTGGTTAGTTTCTAAGAGCAAGGGCCAGAATTTCCCACAGACCACAAATACATACAGGAATACAGATA CCGCTGAGCATCTAATCATGTGGGGAATACACCACCCCTCGTCAACTCAGGAGAAGAATGATCTGTATGGCACACAGTCTCTGTCAATTTCC GTGGGCTCAAGCACATATAGGAACAACTTCGTGCCTGTTGTTGGCGCCAGACCCCAAGTCAATGGCCAGTCTGGAAGGATTGACTTCCATT GGACCCTGGTCCAACCCGGGGATAATATCACCTTTTCCCATAATGGAGGGCTGATCGCGCCTAGTCGCGTCAGTAAACTCATCGGGAGAG GGCTGGGAATTCAATCAGATGCCCCCATTGATAACAACTGTGAAAGTAAATGCTTTTGGCGGGGGGGGTTCCATCAATACACGCCTGCCATTT CAGAATCTGAGCCCCAGGACGGTTGGTCAGTGCCCCAAGTACGTGAACAAGAGAAGCCTGATGCTGGCAACTGGGATGCGCAATGTTCCT GAACTGATTCAGGGACGCGGGTTATTTGGAGCTATTGCCGGTTTTCTCGAGAATGGCTGGGAGGGAATGGTCGATGGGTGGTACGGATTTA GACACCAAAACGCTCAGGGAACAGGCCAGGCGGCTGATTACAAAAGCACCCAGGCCGCTATTGATCAGATCACTGGAAAACTGAACCGCC TTGTCGAGAAAACAAACACCGAATTTGAGAGCATAGAGTCCGAATTCTCAGAGATTGAACATCAAATTGGCAACGTGATAAATTGGACTAAA GACTCCATTACCGACATCTGGACCTACCAGGCCGAGCTCTTGGTTGCTATGGAAAATCAGCATACTATCGACATGGCCGACTCTGAAATGCT TAATCTGTACGAACGCGTGAGAAAACAGTTACGCCAGAACGCGGAGGAAGATGGCAAAGGGTGCTTCGAGATCTACCATGCATGCGATGAC AGCTGTATGGAAAGCATCCGTAATAATACCTATGATCACTCTCAGTATAGGGAACAGGCCCTGTTGAATAGATTAAACATTAACCCAGTGACA CTGAGCAGCGGCTATAAGGACATTATTTTATGGTTTTCCTTTGGAGCAAGTTGTTTTGTGCTGCTGGCGGTGGTGATGGGGCTCGTGTTCTT TTGCCTGAAGAATGGTAACATGCGCTGTACTATCTGTATT
3014	ATGTACAAGATAGTCGTGATAATTGCACTCCTCGGCGCAGTTAAGGGCTTGGATAAGATCTGCCTGGGACACCACGCCGTCGCGAACGGCA CTATTGTCAAGACTTTAACGAACGAGCAAGAAGAGGTAACAAATGCAACAGAAACTGTTGAGAGCACGGGCATCAATCGGCTGTGCATGAA GGGTACGAAGCATAAGGACCTGGGAAACTGTCACCCCATCGGCATGCTGATCGGGACCCCCGCATGTGATTTACACCTTACAGGAATGTG GGATACTCTGATCGAAAGAGAGAATGCCATTGCTTACTGTTACCCTGGGGCCACGGTGAACGTGGAGGCCCTTAGGCAGAAAATCATGGAA AGTGGAGGGATCAACAAGATTTCCACCGGTTTCACGTATGGCTCTTCTATTAATTCCGCTGGAACCACTAGGGCCTGTATGCGAAATGGAG GCAACTCCTTCTACGCTGAATTGAAGTGGCTGGTAAGCAAGTCCAAAGGGCAGAATTTCCCACAGACAACAAACACGTACAGAAACACGGA CACCGCTGAGCACCTCATCATGTGGGGCATCCACCACCCTAGCTCTACCCAGGAAAAAAATGATTTATACGGAACGCAGTCTCTTTCAATCT CCGTCGGCTCAAGCACTTATAGAAATAATTTTGTGCCCGTCGTGGGGGCCCGGCCCCAGGTAAACGGCCAATCAGGGAGGATCGACTTTC ATTGGACCCTGGTTCAGCCCGGGGATAACATAACCTTTAGCCATAACGGAGGTCTTATAGCGCCGAGTCGAGTGTCAAAGCTGATCGGGCG CGGTTTGGGCATCCAGTCTGATGCCCCCATCGACAACAACTGCGAGAGCAAATGCTTTTGGAGGGGAGGGTCCATTAATACCCGTCTGCCA TTCCAGAATCTCTCTCCCCGGACCGTCGGCCAGTGCCCCAAATACGTGAATCGTCGTTCTCTCATGCTCGCCACCGGAATGCGCAACGTGC CTGAGTTGATCCAAGGTCGGGGACTGTTCGGGGCCATTGCCGGCTTTCTGGAGAACGGGTGGGAGGGGATGGTGGATGGCTGGTACGGC TTTAGGCACCAGAACGCTCAGGGAACTGGGCAGGCGGCCGACTACAAGTCAACCCAGGCCGCAATAGATCAAATTACAGGCAAGCTGAAC CGACTGGTCGAAAAAACCAACACAGAATTCGAGTCCATAGAATCCGAATTTTCTGAAATAGAACACCAGATTGGGAACGTTATCAATTGGAC TAAAGATTCCATCACCGATATCTGGACATATCAGGCCGAACTGCTTGTGGCCATGGAAAACCAGCATACCATCGATATGGCTGACTCCGAAA TGTTGAATTTGTATGAAAGGGTACGCAAGCAGCTGAGGCAGAACGCTGAGGAGGACGGTAAGGGATGTTTTGAGATCTACCATGCATGCGA TGACAGCTGTATGGAGTCAATACGCAACAACACTTATGACCACTCCCAATATCGGGAGGAAGCACTTCTCAATCGCTTGAATATTAATCCCG TCACTCTGTCTAGTGGTTATAAGGACATTATTTTGTGGTTCAGCTTTGGAGCCTCCTGTTTCGTTCTACTGGCCGTTGTGATGGGACTATTCT TTTTCTGCCTCAAAAACGGCAACATGCGGTGCACCATCTGTATT

3015	ATGTACAAAATAATTGTTATCATAGCCCTCCTGGGTGCCGTCAAAGGGCTGGATAAAATATGCTTAGGGCACCACGCGGTTGCTAACGGCAC CATCGTCAAAACCCTCACAAACGAACAGGAGGAAGTCACCAACGCAACAGAAACAGTCGAATCAACGGGCATCAATAGACTGTGTATGAAA GGGCGCAAGCACAAGGACTTGGGAAACTGCCACCCGATTGGTATGCTAATTGGAACCCCAGCCTGTGACCTACACCTGACCGGGACTTGG GATACTCTTATTGAGCGCGAAAACGCTATTGCGTATTGTTACCCTGGGGCTACCGTCAACGTTGAGGCACTGAGACAGAAAATAATGGAATC AGGTGGCATAGACAAAATTTCTACCGGCTTCACCTACGGGAGCAGCATAAACTCCGCTGGCACTACACGCGCCTGTATGAGGAATGGCGG CAACTCCTTCTACGCCGAGCTGAAGTGGCTGGTTAGTAAGTCCAAGGGACAGAACTTCCCACAGACCACCAACACTTACCGCAACACCGAT ACAGCCGAGCACCTGATTATGTGGGGAATACACCACCCAAGCTCTACCCAGGAAAAAAACGACTTGTACGGCACTCAAAGTCTGTCCATCT CCGTTGGGTCGTCCACTTACCGCAATAATTTCGTCCCCGTGGTTGGAGCCCGCCCTCAGGTGAACGGCCAAAGCGGGCGCATCGATTTCC ATTGGACCCTCGTTCAGCCTGGCGACAACATTACCTTTAGTCACAATGGCGGTCTGATTGCCCCTTCACGCGTGTCCAAGTTGATTGGACG GGGATTGGGCATCCAGTCGGACGCACCCATCGATAACAATTGTGAGAGTAAATGTTTCTGGCGGGGTGGCTCCATTAATACCCGACTGCCA TTCCAAAACCTCTCTCCTCGTACGGTGGGTCAGTGTCCCAAGTATGTGAACAGAAGATCTCTGATGTTAGCAACCGGTATGAGAAACGTCCC GGAGCTGATCCAGGGCAGAGGCCTGTTCGGAGCCATTGCGGGGTTTCTTGAAAACGGATGGGAGGGCATGGTGGACGGATGGTATGGGT TTCGACATCAAAATGCCCAGGGGACCGGGCAGGCAGCTGACTACAAATCTACCCAGGCGGCAATAGATCAGATCACTGGAAAGTTAAACAG ACTCGTGGAGAAAACTAACACTGAGTTCGAATCTATCGAGTCAGAGTTTTCAGAAATTGAACATCAGATCGGAAACGTTATCAATTGGACGA AAGATAGTATCACAGACATTTGGACCTATCAAGCTGAATTGCTTGTGGCCATGGAAAACCAACATACCATCGACATGGCTGACTCGGAAATG TTGAATTTGTACGAGCGGGTGAGGAAGCAACTGAGGCAAAATGCCGAGGAGGACGGCAAAGGCTGCTTTGAGATCTACCACGCATGCGAT GACTCTTGTATGGAAAGTATTCGGAATAATACTTATGACCATTCCCAGTATCGTGAGGAAGCACTGCTCAACCGCCTAAATATTAATCCCGTG ACCCTGTCATCAGGGTATAAGGACATTATCCTTTGGTTTTCCTTCGGCGCTAGCTGTTTCGTGCTGCTAGCGGTAGTGATGGGCCTTTTCTT CTTTTGTTTGAAGAATGGCAATATGCGATGCACAATCTGCATA
3016	ATGTATAAGATCGTGGTCATCATAGCCCTGCTCGGAGCCGTTAAGGGTCTTGACAAGATTTGTCTTGGTCATCATGCTGTTGCTAACGGAAC CATCGTGAAAACTCTGACAAACGAGCAGGAGGAAGTGACAAACGCCACGGAGACAGTGGAATCTACTGGAATCAACCGCTTATGCATGAAA GGCAGGAAACACAAAGACCTGGGGAATTGCCACCCAATTGGAATGCTGATCGGCACACCTGCCTGCGACCTGCATCTCACAGGGATGTGG GATACACTGATCGAAAGAGAGAATGCTATCGCATACTGCTACCCAGGGGCTACTGTCAATGTGCAGGCCCTTCGACAGAAGATTATGCAAT CCGGAGGTATCAACAAAATTAGCACTGGCTTCACCTACGGGTCTTCCATCAATTCTGCCGGCACGACCAGAGCATGTATGAGAAATGGGGG CAACAGTTTTTACGCCGAGTTGAAGTGGCTGGTGAGTAAGAGCAAAGGACAGAACTTCCCGCAGACCACCAACACATACAGGAATACGGAT ACCGCCGAGCACTTAATCATGTGGGGAATTCACCACCCCAGTTCGACACAGGAGAAGAACGATCTCTATGGCACTCAGTCTCTGTCTATTTC TGTGGGATCATCAACATATCGCAATAACTTTGTACCTGTTGTGGGGGCTAGGCCTCAGGTGAACGGCCAATCCGGACGAATCGACTTCCAC TGGACTCTGGTTCAACCAGGTGATAATATTACTTTCTCACACAACGGCGGACTGATTGCACCTAGCCGAGTGAGTAAGCTCATTGGTAGAGG ACTGGGGATTCAGTCCGATGCCCCTATAGACAATAACTGCGAAAGCAAATGCTTCTGGCGCGGCGGAAGCATAAATACAAGATTGCCATTT CAGAACCTGAGTCCTCGGACCGTTGGACAGTGTCCTAAATACGTGAACAGAAGAAGCCTGATGCTTGCCACTGGCATGCGGAATGTGCCC GAGCTCATTCAGGGACGGGGCCTATTCGGCGCAATCGCCGGATTTCTGGAGAATGGTTGGGAAGGGATGGTCGATGGCTGGTACGGCTTT AGACACCAGAATGCACAGGGGACAGGGCAGGCGGCCGATTACAAATCCACTCAGGCTGCTATCGACCAGATCACCGGCAAGTTAAATAGG TTGGTCGAGAAGACGAACACCGAATTCGAATCTATCGAGTCAGAGTTTAGCGAGATTGAGCATCAGATAGGTAATGTGATTAACTGGACTAA GGACAGCATCACTGATATCTGGACTTACCAGGCAGAGCTCCTGGTTGCCATGGAGAACCAGCACACAATAGACATGGCCGACAGTGAAATG CTAAACCTGTACGAACGCGTGCGCAAGCAGCTCCGGCAGAACGCAGAAGAAGATGGCAAGGGGTGTTTTGAGATATATCATGCATGCGAT GACTCCTGTATGGAGAGTATCCGTAACAACACTTACGATCACTCCCAGTACCGTGAGGAGGCTCTGCTGAACAGACTGAACATCAACCCCG TAACTTTAAGTAGTGGATATAAGGATATCATTCTTTGGTTTAGCTTCGGAGCAAGCTGTTTCGTCCTTCTCGCAGTGGTTATGGGCCTGTTCT TTTTTTGCCTAAAGAATGGGAATATGCGGTGCACAATATGTATT
3017	ATGTATAAGATCGTGGTGATTATTGCCCTGCTGGGCGCTGTGAAGGGCTTAGACAAGATTTGCTTAGGTCACCATGCAGTGGCAAATGGAA CAATTGTCAAGACATTGACCAATGAACAGGAGGAGGTAACCAACGCTACAGAAACCGTTGAGTCCACCGGTATTAACCGGCTTTGCATGAA GGGTCGCAAGCACAAGGATCTGGGCAATTGCCACCCCATCGGTATGCTCATTGGAACTCCCGCATGTGATCTGCATCTGACAGGGATGTG GGATACGTTGATAGAGCGTGAGAACGCCATCGCCTACTGTTACCCAGGTGCCACAGTGAACGTGGAAGCCCTGCGTCAAAAGATCATGGAA TCCGGAGGGATCAATAAAATTAGTACAGGCTTCACGTATGGCTCTTCCATCAACTCTGCTGGGACTACTCGTGCGTGCATGCGGAATGGAG GCAATTCATTTTACGCGGAACTAAAATGGCTTGTGAGCAAAAGCAAAGGACAAAACTTCCCGCAGACGACCAACACGTATCGGAACACCGA CACCGCCGAACATCTAATCATGTGGGGCATCCATCATCCCTCATCCACCCAGGAAAAAAATGACTTATACGGAACACAGTCATTGAGCATTT CCGTAGGATCAAGTACATACCGAAATAATTTCGTGCCTGTGGTGGGTGCGGGCCCCCAAGTTAATGGCCAATCAGGCCGTATTGACTTTCA TTGGACCCTCGTTCAGCCTGGCGACAACATCACCTTCTCCCATAACGGGGGACTCATCGCCCCATCAAGAGTGTCGAAGCTGATCGGCCGA GGACTCGGGATACAGTCTGACGCCCCCATCGACAATAATTGTGAATCCAAGTGCTTTTGGAGAGGAGGGTCAATCAACACCCGCCTACCTT TTCAGAATCTTTCACCCAGGACCGTTGGGCAGTGTCCCAAATACGTGAACCGGAGATCTCTCATGCTTGCAACCGGGATGAGAAACGTGCC AGAGCTCATCCAGGGGCGCGGCCTATTTGGGGCCATAGCCGGTTTTCTGGAGAACGGATGGGAGGGAATGGTTGATGGCTGGTACGGCTT CAGGCACCAGAACGCACAGGGCACCGGGCAAGCCGCAGACTACAAATCAACTCAGGCGGCTATCGATCAGATCACAGGTAAATTAAACCG ACTCGTCGAAAAGACAAATACCGAGTTTGAGTCTATCGAATCCGAGTTTTCTGAAATTGAGCATCAAATCGGCAACGTGATTAACTGGACTAA GGACTCCATCACCGATATTTGGACTTATCAAGCAGAACTACTGGTCGCCATGGAAAATCAACATACTATCGATATGGCAGATAGCGAAATGC TTAACCTCTACGAAAGAGTGCGAAAGCAGCTACGCCAAAATGCCGAGGAAGACGGGAAGGGCTGCTTCGAGATTTACCATGCGTGTGATGA TTCCTGCATGGAAAGTATTAGGAACAATACTTATGATCACAGCCAGTATCGTGAGGAAGCTCTGCTTAACAGACTTAACATCAATCCAGTGAC ACTCAGCAGTGGGTACAAAGACATCATCCTCTGGTTCAGCTTTGGAGCTTCTTGCTTCGTTTTGCTTGCTGTAGTCATGGGACTGTTTTTTTT TTGTCTGAAGAACGGTAATATGCGATGCACTATTTGTATA

3018	ATGTATAAGATCGTGGTCATCATCGCTTTGCTGGGGGCTGTTAAGGGGCTCGACAAGATTTGTTTGGGCCACCACGCCGTGGCCAACGGGA CCATTGTGAAGACGCTGACAAATGAACAAGAAGAAGTCACAAATGCCACAGAGACTGTTGAATCCACTGGCATTAATAGACTCTGCATGAAA GGCAGGAAACACAAAGATCTGGGCAACTGTCACCCTATCGGAATGCTGATTGGCACTCCCGCTTGTGATCTCCATCTTACAGGTATGTGGG ATACCCTTATTGAGCGAGAAAATGCCATCGCCTACTGTTATCCAGGTGCTACTGTGAACGTGGAGGCCTTGAGGCAGAAGATTATGGAGTC AGGAGCCATCAATAAGATTTCCACCGCGGTTTACATACGGCTCATCCATCAACTCCGCCGGCACGACTCGAGCTTGCATGAGAAACGGTGGC AACTCCTTTTATGCGGAGCTAAAGTGGCTGGTCAGCAAGTCAAAAGGTCAGAACTTCCCCCAGACTACGAACACCTATCGGAATACAGACAC CGCTGAGCATTTGATCATGTGGGGAATCCACCATCCTAGCTCCACCCAGGAGAAGAATGACCTGTATGGAACACAATCTCTTTCTATCTCCG TGGGCTCTTCTACCTACCGTAACAATTTCGTTCCCGTTGTGGGGGCTCGGCCTCAGGTGAATGGTCAATCCGGAAGAATCGACTTTCACTG GACATTAGTACAGCCCGGCGACAACATTACATTTAGTCATAATGGGGGGCTGATCGCGCCTAGTCGGGTCTCAAAGTTGATCGGGAGAGGC CTCGGAATCCAGAGCGATGCACCTATCGATAACAACTGTGAAAGCAAGTGCTTCTGGCGCGGCGGTTCTATTAATACCAGATTGCCCTTCCA GAACCTGTCTCCCCGAACAGTCGGGCAGTGCCCCAAATACGTGAACAGGAGATCTCTGATGCTGGCCACAGGGATGCGCAACGTGCCAGA GCTGATCCAGGGAAGGGGTCTCTTCGGGGCCATTGCCGGCTTTCTGGAAAATGGCTGGGAAGGGATGGTGGACGGGTGGTATGGCTTCA GGCACCAAAACGCGCAAGGCACTGGCCAGGCGGCGGATTACAAGAGCACGCAAGCTGCTATAGACCAGATCACGGGGAAGCTGAACAGA CTCGTGGAAAAGACTAATACTGAATTTGAATCGATCGAGTCTGAGTTCTCCGAAATCGAGCATCAGATCGGGAATGTCATAAATTGGACCAA GGATTCCATTACAGATATTTGGACCTATCAGGCCGAACTGCTGGTGGCCATGGAGAATCAGCACACTATTGATATGGCCGACAGTGAGATG CTGAACCTGTATGAGCGTGTTAGGAAGCAACTTCGACAGAACGCAGAAGAAGATGGTAAGGGATGTTTCGAGATTTACCATGCCTGTGACG ACAGTTGTATGGAGAGCATTAGGAATAACACTTATGACCATAGCCAATACCGGGAAGAAGCTTTGTTAAATCGCCTGAATATCAATCCCGTC ACCTTATCGTCGGGGTATAAGGATATTATTCTGTGGTTCAGCTTCGGAGCGTCCTGCTTTGTTCTGTTGGCCGTGGTAATGGGCCTGTTTTT CTTTTGTCTTAAGAACGGGAACATGCGCTGCACAATCTGCATC
3019	ATGTACAAGATTGTGGTTATCATCGCACTGTTAGGAGCCGTGAAGGGGTTGCACAAGATCTGTCTGGGGCATCATGCCGTTGCTAATGGAA CAATCGTGAAGACGCTCACCAATGAACAAGAAGAAGTCACAAACGCCACCGAAACCGTGGAGAGCACCGGAATTAACAGACTTTGTATGAA GGGACGCAAACACAAGGATCTGGGGAACTGCCACCCAATCGGCATGCTGATAGGAACCCCTGCCTGTGATCTGCACTTAACTGGCATGTG GGACACACTTATAGAGCGCGAGAACGCGATCGCGTACTGCTACCCCCGAGCAACCGTTAACGTCGAAGCTTTGAGGCAGAAAATTATGGAA AGCGGCGGCATAAATAAAATTTCTACTGGCTTTACGTACGGCAGCTCTATAAACAGTGCTGGCACCACTCGTGCCTGTATGCGGAACGGCG GCAATTCCTTTTACGCCGAATTAAAATGGCTTGTTAGCAAGTCAAAGGGCCAGAACTTTCCGCAGACGACTAATACATACCGAAACACCGAC ACCGCTGAACATCTGATTATGTGGGGTATACATCATCCAAGTTCCACCCAAGAAAAAAATGACTTGTATGGGACCCAGAGCCTAAGCATCTC CGTGGGGAGTTCTACTTACCGAAACAACTTCGTACCTGTGGTGGGGGCCCGACCACAGGTAAATGGACAGTCGGGCCGGATTGATTTCCA CTGGACCTTGGTGCAGCCGGGAGATAACATAACATTCTCTCATAACGGAGGTCTCATAGCCCCAAGTCGTGTGTCAAAGCTCATCGGTAGA GGCCTGGGCATTCAGTCAGACGCACCCATTGATAATAACTGTGAGTCTAAATGCTTTTGGCGCGGCGGGAGTATCAATACTAGACTGCCTTT TCAGAACCTCAGCCCAAGGACCGTGGGGCAGTGTCCCAAGTATGTAAATCGGCGAAGCTTAATGCTGGCTACTGGTATGCGGAATGTGCCT GAACTCATTCAGGGACGGGGCCTCTTCGGAGCAATAGCTGGTTTCCTGGAGAACGGCTGGGAAGGCATGGTGGATGGTTGGTACGGTTTT AGACACCAAAATGCCCAGGGAACGGGACAGGCTGCCGACTATAAGTCAACTCAAGCCGCGATCGACCAGATCACTGGTAAATTAAACCGAC TCGTCGAAAAGACAAACACGGAGTTCGAGTCTATCGAAAGTGAGTTCTCAGAGATTGAACACCAAATCGGGAATGTGATTAACTGGACAAAG GACTCGATAACCGATATCTGGACCTACCAGGCAGAACTTCTGGTGGCCATGGAAAACCAGCACACTATTGACATGGCCGACTCAGAGATGC TGAATCTGTACGAGAGAGTGAGGAAGCAGCTCCGCCAGAATGCGGAGGAGGATGGTAAAGGCTGCTTCGAAATCTACCACGCTTGCGATG ACTCCTGCATGGAATCAATCCGAAATAATACATATGACCACTCACAGTACAGACGAAGAAGCCCTTCTGAACCGCCTGAACATCAACCCCGTG ACGCTGAGTTCAGGATATAAGGATATCATTCTCTGGTTCTCCTTTGGGGCGAGTTGCTTCGTGCTGCTTGCGGTAGTCATGGGACTCTTTTT TTTTTGTCTTAAGAACGGTAATATGCGGTGCACTATTTGTATC
3020	ATGTACAAGATCGTGGTAATTATCGCCCTGCTCGGAGCTGTGAAAGGATTAGACAAAATTTGCCTCGGCCACCATGCTGTAGCTAATGGTAC AATAGTCAAAACCCTAACTAATGAACAGGAAGAGGTGACTAATGCAACGGAAACTGTCGAGAGTACCGGCATTAACCGCCTGTGTATGAAAG GGAGAAAGCACAAGGACTTAGGCAATTGTCATCCAATCGGGATGCTAATTGCCACTCCAGCCTGTGATCTGCATCTGACCGGAATGTGGGA CACCTTAATTGAGAGGGAAAACGCCATAGCCTATTGCTATCCTGGGGCAACTGTTAATGTGGAAGCTCTCCGGCAAAAAATCATGGAATCTG GAGGCATTAATAAGATCTCCACGGGCTTCACATATGGGTCCTCTATCAATTCCGCCGGGACCACGCGCGCATGCATGAGAAACGGAGGGAA TTCTTTCTACGCGGAACTTAAGTGGCTGGTCAGTAAGTCTAAAGGGCAGAATTTTCCTCAGACCACTAATACCTACAGAAATACGGATACAG CTGAGCACCTCATTATGTGGGGAATACACCACCCTTCAAGCACACAGGAAAAAAACGATCTCTACGGGACCCAGAGCCTGTCCATTAGTGT CGGATCTAGCACATACCGGAACAATTTCGTACCCGTTGTGGGAGCACGCCCACAGGTGAATGGGCAGTCAGGCAGAATCGATTTCCATTGG ACTCTCGTGCAGCCCGGCGATAATATAACCTTCAGTCATAATGGCGGATTGATCGCCCCCTCGCGTGTTTCAAAATTAATCGGGAGGGGGC TCGGGATACAAAGCGACGCTCCTATCGACAACAACTGTGAGTCCAAGTGTTTTTGGCGCGGCGGGAGCATTAATACCCGTCTCCCGTTCCA GAACCTGTCTCCCCGGACAGTAGGACAGTGTCCGAAGTATGTCAATCGTAGAAGTCTTATGCTGGCGACGGGAATGCGGAACGTGCCCGA GCTGATTCAAGGCCGGGGTCTGTTCGGAGCAATAGCTGGTTTCCTGGAGAATGGTTGGGAGGGCATGGTTGACGGCTGGTATGGGTTCAG ACATCAGAACGCACAGGGAACCGGCCAGGCTGCTGACTATAAATCCACTCAGGCAGCCATAGATCAGATCACCGGGAAGCTAAACCGGCT GGTTGAGAAAACTAATACGGAGTTCGAGTCTATCGAGTCCGAGTTTAGCGAAATAGAGCATCAGATAGGGAACGTGATAAATTGGACGAAA GACTCTATTACCGACATCTGGACCTACCAGGCGGAACTGCTGGTGGCTATGGAGAATCAGCACACCATCGACATGGCTGATAGCGAAATGT TGAATCTGTACGAGCGGCTCAGAAAACAGTTAAGACAGAACGCAGAGGAAGATGGAAAGGGCTGCTTCGAAATATATCACGCATGTGATGA TTCCTGCATGGAGTCTATTAGAAACAACACTTATGACCATTCGCAGTACCGCGAAGAAGCGCTGCTAAACAGACTCAACATCAATCCTGTGA CCTTGAGTAGCGGATACAAGGACATTATACTGTGGTTCTCCTTTGGTGCCTCCTGTTTTGTCTTGCTTGCCGTAGTTATGGGCCTGTTCTTTT TCTGCTTGAAGAACGGGAACATGCGCTGCACAATTTGTATC

3011	ATGTATAAAGTGGTGGTGATTATCGCTTTACTGGGAGCTGTAAAGGGGCTGGATAAGATTTGTCTGGGACATCACGCAGTCGCGAACGGCA CCATTGTGAAAACACTTACTAATGAACAGGAGGAGGTGACCAACGCCACCGAAACCGTCGAGTCAACAGGTATAAACAGATTGTGCATGAA GGGCCGAAAGCACAAGGACCTGGGAAATTGTCATCCAATTGGAATGTTGATCGGCACTCCAGCCTGCGATCTTCATCTTACCGGAACATGG GACACGCTTATTGAACGGGAAAATGCGATCGCATATTGCTACCCCGGGGCCACAGTGAATGTGGAGGCACTGAGACAGAAAATCATGGAAA GTGGCCGGGATCGACAAGATTAGTACTGGCTTTACATACGGGAGCAGCATCAACTCCGCCGGGACCACCCGAGCCTGCATGAGGAACGGG GGCAACAGTTTTTATGCTGAGCTGAAGTGGCTCGTGTCAAAAAATAAGGGCCAGAATTTCCCTCAAACAACGAATACATATCGCAACACAGA TACAGCGGAACACTTAATCATGTGGGGCATTCACCACCCTAGCTCCATTCAAGAAAAGAACGATCTCTACGGAACACAATCGTTGAGCATCT CAGTGGGGTCTTCCACCTATAGAAACAACTTTGTACCTGTGGTCGGCGCAAGACCTCAAGTCAATGGACAAAGCGGCAGAATCGACTTCCA CTGGACTCTGGTGCAGCCAGGCGATAATATAACCTTCTCTCACAACGGCGGCTTGATCGCTCCCAGCCGGGTATCTAAGTTAATCGGCCGT GGTCTCGGAATCCAGAGCGATGCACCGATTGACAATAACTGTGAAAGTAAATGTTTCTGGCGTGGCGGCTCTATAAACACTAGACTCCCCTT CCAGAACTTGAGTCCTAGAACAGTGGGGCAGTGCCCGAAATACGTCAACCGGAGATCGTTGATGCTCGCAACTGGCATGAGGAACGTCCC CGAGTTAATACAGGGACGGGGACTATTCGGCGCCATAGCCGGGTTCCTCGAGAACGGATGGGAGGGTATGGTTGATGGTTGGTACGGCTT TCGACATCAAAACGCACAAGGTACAGGGCAGGCTGCCGATTACAAGAGCACTCAGGCCGCCATCGACCAGATCACTGGGAAACTCAACAG GTTAGTGGAGAAAACCAATACCGAATTTGAATCCATCGAGAGCGAATTCTCAGAGATTGAGCATCAGATAGGAAATGTGATAAACTGGACCA TGGATAGTATTACAGACATTTGGACGTACCAAGCAGAGCTGCTTGTCGCGATGGAGAATCAACACACCATCGACATGGCGGACTCCGAAAT GCTGAATCTGTATGAGAGGGTTCGGAAACAGCTTAGACAAAATGCAGAGGAGGATGGCAAGGGCTGCTTTGAGATCTACCACGCATGTGAT GACTCCTGTATGGAATCCATCCGGAACAACACGTATGACCACTCTCAGTACAGAGAAGAGGCTCTCCTGAATCGCCTTAATATTAACCCTGT GACACTGTCCTCCGGCTATAAGGATATCATCCTGTCGTTTAGCTTTGGAGCAAGCTGCTTCGTGTTGCTGGCCGTAGTTATCGGCCTGGTCT TCTTCTGTCTTAAGAACGGAAACATGCGGTGCACAATCTGTATT
3012	ATGTATAAGGTCGTCGTTATTATAGCGCTGCTTGGAGCTGTCAAAGGTTTGGATAAAATCTGTCTGGGCCATCACGCAGTCGCCAACGGTAC CATTGTGAAAACGCTCACAAACGAACAGGAAGAGGTCACGAACGCCACAGAGACTGTGGAGTCTACAGGGATTAACCGCCTGTGCATGAAA GGCCGGAAACATAAGGACCTTGGCAACTGCCATCCAATCGGCATGCTGATTGGCACTCCGGCCTGTGATCTGCATTTGACCGGCACATGG GATACCCTGATCGAAAGGGAGAATGCCATCGCTTACTGTTATCCCGGCCGCAACTGTGAATGTGGACGCACTGAGACAGAAGATCATGGAGT CTGGGGGGATCGATAAAATTAGTACCGGGTTTACTTACGGCAGTTCGATTAATAGCGCGGGAACAACGAGGGCGTGTATGAGGAATGGAG GAAACTCCTTCTATGCCGAGCTCAAGTGGCTGGTGAGCAAGAACAAGGGGCAAAACTTTCCCCAAACAACTAACACATACCGGAATACAGA TACTGCTGAACATCTGATAATGTGGGGAATCCACCATCCTTCTTCAATTCAGGAGAAAAACGATCTTTACGGAACTCAAAGTTTAAGCATAAG TGTGGGGTCATCTACGTATAGAAACAACTTTGTGCCCGTGGTTGGCGCAAGGCCGCAGGTGAATGGTCAATCTGGGAGGATAGATTTCCAC TGGACATTGGTGCAGCCGGGTGATAATATCACCTTTTCCCATAATGGGGGATTGATTGCCCCCTCCCGGGTGTCGAAACTTATTGGCCGCG GGCTAGGTATTCAGAGTGACGCCCCCATTGATAATAATTGTGAGTCTAAATGTTTTTGGCGGGGTGGCAGCATTAATACACGTCTGCCATTC CAGAACCTGTCCCCCAGGACTGTCGGGCAATGCCCCAAATACGTGAATAGACGGTCTCTGATGCTGGCCACTGGGATGCGCAATGTACCC GAGCTTATCCAGGGACGGGGCCTTTTTGGGGCCATCGCCGGTTTCCTGGAGAACGGGTGGGAAGGCATGGTTGATGGCTGGTATGGGTTT AGACACCAGAATGCTCAAGGGACTGGGCAGGCGGCCGATTATAAGAGTACACAGGCGGCCATCGACCAGATCACAGGCAAGCTCAACCGT CTGGTGGAGAAAACAAATACAGAATTCGAAAGCATAGAGAGTGAATTTTCTGAGATTGAGCACCAGATTGGTAATGTCATCAATTGGACAAT GGATTCCATCACTGACATCTGGACATACCAAGCCGAGCTATTGGTGGCGATGGAGAATCAGCATACAATAGATATGGCCGACTCTGAAATG CTGAACTTATATGAGAGAGTGCGTAAACAGTTGAGGCAGAATGCTGAAGAAGACGGTAAGGGATGTTTTGAGATATATCATGCCTGTGATGA CAGTTGCATGGAATCCATCAGAAATAACACTTACGACCACAGCCAGTACAGAGAGGAAGCCCTACTGAACAGGCTCAATATCAATCCAGTGA CTCTATCCAGCGGCTACAAGGACATTATCTTATGGTTTTCTTTTGGTGCTTCATGTTTTGTGCTCCTCGCCGTGGTCATGGGGCTCGTCTTCT TCTGTTTAAAGAATGGGAATATGCGGTGCACAATATGTATC
3013	ATGTACAAAATAGTAGTAATCATTGCTTTGCTTGGCGCGGTGAAAGGCTTGGATAAGATTTGCCTGGGACACCATGCTGTCGCGAATGGAAC CATAGTTAAGACTTTGACTAATGAGCAAGAGGAGGTGACCAACGCCACCGAAACGGTGGAGTCCACAGGCATCAACAGGCTCTGTATGAAA GGACGGAAGCATAAGGACCTGCGGCAACTGTCATCCAATTGGAATGCTGATCGGGACGCCCGCGTGTGACCTCCATCTGACAGGGATGTGG GATACTCTTATTGAGAGAGAGAACGCTATCGCCTACTGTTATCCTGGTGCTACAGTGAACGTGGAAGCCCTCAGGCAGAAGATTATGGAGA GCGGTGGGATCAACAAGATTAGTACAGGGTTTACCTATGGGTCATCCATAAACAGTGCGGGCACTACCCGAGCGTGCATGCGCAATGGGG GCAATTCGTTCTATGCCGAACTGAAGTGGTTAGTCTCTAAGTCTAAAGGTCAGAATTTTCCTCAGACAACTAACACATACCGGAATACCGATA CCGCTGAACACCTGATCATGTGGGGAATACATCACCCTAGTAGTACACAAGAGAAAAATGACCTATATGGGACACAGAGCCTGTCAATTAGC GTGGGTAGTTCCACATATAGAAACAACTTTGTTCCTGTCGTGGGTGCCCGGCCTCAGGTAAATGGGCAGAGCGGACGGATAGATTTTCATT GGACTTTGGTCCAGCCCGGGGATAACATTACCTTCTCTCATAACGGGGGGCTGATCGCTCCAAGCCGGGTGTCTAAGCTCATAGGCAGAG GCCTGGGTATCCAATCGGACGCCCCCATCGATAACAACTGTGAAAGTAAATGCTTTTGGAGGGGCGGCAGTATTAATACGCGGCTACCGTT TCAGAATCTGTCTCCACGCACAGTAGGGCAGTGTCCAAAGTACGTCAACCGCAGATCCCTCATGCTGGCCACGGGGATGAGGAACGTGCC CGAGCTGATCCAGGGGAGGGGCCTATTTGGGGCTATCGCCGGCTTCCTGGAGAATGGTTGGGAGGGAATGGTCGATGGCTGGTACGGCT TTCGCCATCAGAACGCCCAAGGAACTGGACAAGCTGCCGACTACAAAAGCACGCAAGCCGCCATCGATCAAATTACCGGAAAGCTGAATCG GCTAGTGGAGAAGACCAACACCGAATTTGAGTCTATAGAGTCAGAGTTTAGCGAAATCGAGCATCAGATTGGCAACGTCATTAATTGGACCA AGGACAGCATAACCGACATATGGACATACCAAGCCGAACTCCTGGTAGCTATGGAAAATCAGCATACCATCGATATGGCAGATTCCGAAATG CTGAACCTGTATGAAAGAGTGAGGAAGCAACTGCGTCAGAATGCCGAGGAAGACGGCAAGGGGTGCTTTGAGATCTACCACGCATGTGAT GATTCTTGCATGGAGAGCATCAGGAATAATACATACGACCACAGCCAGTACCGGGAAGAAGCCCTATTGAACAGGCTTAACATCAATCCCGT CACTCTATCCAGCGGCTATAAAGATATAATCTTATGGTTTTCCTTTGGCGCGTCTTGCTTTGTGCTCCTGGCGGTGGTGATGGGCTTATTCTT CTTTTGTCTCAAGAATGGCAACATGAGGTGCACAATCTGCATT

3014	ATGTACAAGATAGTGGTCATCATCGCACTCCTGGGCGCCGTGAAGGGCCTGGATAAAATTTGCCTGGGGCACCATGCCGTCGTGAACGGC ACGATCGTCAAGACTCTCACTAATGAGCAGGAGGAAGTCACTAATGCTACGGAGACAGTTGAGTCGACCGGATTGAATAGATTGTGCATGA AGGGACGTAACCACAAGGACCTCGGGAACTGCCACCCAATTGGCATGTTGATCGGAACACCCGCCTGCGATCTCCATCTGACCGGTACGT GGGATACACTTATCGAGAGGGAAAACGCAATCGCCTATTGCTATCCCGGTGCAACCGTTAACGAAGAAGCTCTGAGGCAGAAAATCATGGA GTCTGGTGGGATTAACAAGATTAGTACAGGGTTTACATACGGAAGCTCCATCAACAGTGCAGGCACTACCCGTGCCTGCATGAGAAACCGA GGCAACTCTTTTTACGCTGAACTTAAATGGTTAGTAAGCAAGAGTAAAGGCCAGAATTTTCCACAGACGACAAACACATACAGAAACACCGA CACAGCCGAACACCTGATCATGTGGGGCATCCACCACCCATCCAGCACCCAGGAGAAGAACGATCTATATGGAACTCAGAGCCTGTCTATT AGCGTAGGCAGTTCAACCTACCAGAATAACTTCGTGCCAGTGGTGGGAGCTCGGCCTCAGGTTAATGGACAGAGCGGCAGAATCGACTTC CACTGGACTCTCGTGCAACCCGGCGATAACATCACTTTTTCGCATAATGGAGGCTTAATCGCACCCTCCCGAGTCAGCAAGCTCATCGGAA GGGGATTAGGCATACAGAGTGATGCACCCATCGATAACAACTGCGAAAGTAAGTGCTTCTGGCGAGGGGGTTCCATTAATACTAGATTACC ATTTCAAAATCTGAGTCCAAGAACCGTTGGTCAGTGCCCGAAGTATGTCAATAAGCGTTCTCTGATGTTGGCAACTGGGATGAGGAATGTGC CAGAACTGATGCAAGGGCGAGGCCTCTTTGGTGCAATCGCTGGCTTTATAGAGAACGGGTGGGAAGGGATGGTGGATGGCTGGTATGGGT TCCGCCATCAGAATGCACAAGGCACCGGTCAGGCCGCCGACTACAAGTCAACCCAGGCGGCTATCGATCAAATCACGGGGAAACTGAACA GGTTAATCGAAAAAACAAACACAGAGTTCGAGTCAATCGAGTCTGAGTTTTCTGAAATTGAGCACCAAATCGGGAATGTGATTAACTGGACA AAAGACTCTATCACAGACATCTGGACCTACCAGGCCGAACTCCTGGTGGCCATGGAGAATCAACACACCATTGACATGGCGGACTCCGAAA TGCTGAATCTGTATGAGAGGGTTAGGAAACAACTTAGGCAGAATGCTGAGGAAGACGGCAAAGGCTGTTTTGAAATTTACCACGCCTGTGA TGATAGTTGTATGGAGAGCATTCGGAACAATACCTACGATCACTCTCAATATCGCGAGGAAGCTCTGCTTAACAGGCTTAACATTAACCCCG TCACGCTGTCAAGTGGCTACAAAGACATTATTCTCTGGTTCTCCTTCGGTGCATCATGCTTTGTTCTGCTGGCTGTTGTCATGGGCTTAGTGT TCTTCTGCCTCAAGAACGGTAATATGAGATGTACCATCTGCATT
3015	ATGTATAAGGTGGTGGTGATCATCGCCCTCTTGGGTGCTGTTAGAGGGCTGGATAAAATCTGTCTAGGACATCATGCCGTGGCAAACGGTA CTATCGTAAAAACTCTGACAAATGAGCAGGAGGAGGTGACCAACGCTACTGAGACCGTCGAGTCAAAGAGCCTGGGAAAGTTGTGCATGAA AGGGAGGTCATATAATGACCTGGGGAACTGTCACCCTATAGGTATACTTATTGGCACCCCTGCCTGCGATCTCCATCTCACTGGGACATGG GACACCCTGATTGAACGGGAGAACGCTGTTGCTTATTGCTACCCCGGTGCTACAGTTAATGAGGAGGCGTTGCGTCAGAAGATAATGGAAA GCGGGGGAATCTCGAAGATTAGCACTGGCTTTACCTACGGGACCAGTATTAATAGCGCGGGAACCACAAAGCCTGCATGAGAAACGGAG GCAACAGCTTTTACGCCGAATTGAAATGGTTAGTGTCAAAGAACAAAGGACAGAACTTCCCTCAGACCACAAACACGTACAGAAACACCGAT ACTGCTGAGCACTTGATTATCTGGGGCATTCATCACCCATCTAGCACACAAGAGAAAAATGATCTGTATGGAACCCAGTCACTGAGCATCTC AGTCGGCAGTTCCACTTACCAAAATAATTTTGTTCCTGTGGTGGGAGCACGCCCACAGGTGAATGGACAGTCTGGAAGGATTGATTTTCACT GGACCCTGTTGCAGCCAGGAGACAATATAACCTTCAGTCATAACGGGGGCCTCATCGCGCCGTCTCGGGTGTCAAAGTTGATTGGAAGAG GACTCGGGATTCAATCCGAAGCCCCCATCGATAATGGCTGTGAATCTAAATGCTTTTGGAAAGGTGGTTCGATCAATACAAAGCTGCCCTTC CAAAACCTTTCCCCGCGGACAGTTGGACAATGCCCCAAGTACGTGAACAAGCGCAGTCTGATGCTCGCCACCGGGATGAGAAACGTGCCG GAGATTATGCATGGCAGGGGCCTGTTCGGAGCTATCGCTGGATTCATTGAAAACGGGTGGGAGGGCATGGTTGACGGCTGGTATGGATTC AGGCATCAGAACGCACAGGGGACAGGCCAGGCAGCTGACTATAAGAGTACCCAGGCCGCTATCGACCAAATAACAGGTAAGCTTAACCGT CTTATCGAAAAAACTAACACAGAGTTCGAATCAATCGAAAGCGAATTCAGTGAGATTGAACACCAGATTGGTAACATAATTAATTGGACAAAA GACTCTATCACAGATATTTGGACCTACCAGGCAGAACTGCTCGTGGCCATGGAAAACCAGCACACTATCGATATGGCAGATTCCGAAATGTT AAACCTGTATGAGCGAGTGCGAAAACAACTGCGCCAAAATGCAGAGGAAGACGGGAAAGGCTGTTTTGAAATTTACCATGCGTGCGACGAT TCTTGTATGGAATCCATCAGAAACAACACATATGATCACTCTCAGTATAGAGAAGAGGCTCTTCTGAATCGCTTGAATATAAACCCTGTGAAG TTAAGTTCAGGGTATAAAGATATCATCCTTTGGTTCTCTTTCGGGGCTTCTTGTCTGATCTTGCTTGCTGTGGTCATGGGTCTCGTGTTTTTTT GTCTGAAAAACGGCAACATGCGATGTACAATTTGTATT
3016	ATGTACAAGATTGTAGTCATCATCGCCCTACTGGGTGCAGTCAAGGGACTGGACAAAATATGCCTGGGCCATCATGCAGTGGCCAACGGGA CCATTGTTAAGACACTTACAAATGAGAAAGAAGAGGTTACTAATGCCACCGAAACTGTGGAGTCCACCGGCCTAAACCGACTCTGCATGAAG GGAAGCAAACACAAAGATCTGGGAAACTGCCACCCCATAGGGATGCTCATCGGCTCTCCCGCATGTGACCTGCATCTGACAGGAACCTGG GATACCCTGATCGAGAGAGAAAACGCGATCGCCTATTGTTACCCCGGCGCGACAGTGAATGGTGAGGCTCTGCGCCAAAAAATAATGGAAT CAGGAGGAATAGACAAAATTTCAACAGGATTTACATACGAATCCTCCATTAATAGCGCCGGCACAACTAGAGCTTGTATGCGGAATGGGGG CAATTCCTTTTACGCGGAGCTGAAATGGTTGGTATCTAAAAGCAAAGGACAGAACTTCCCTCAGACTACCAACACCTATCGAAATACAGACA CAGCCGAGCACTTAATTATGTGGGGGATACATCACCCCTCTTCTACACAGGAGAAAAATGACTTATACGGAACCCAATCCTTAAGCATTAGT GTGGGCAGCAGCACATATCGAAACAATTTTGTACCAGTGGTAGGAGCTCGCCCTCAGGTGAATGGACAGAGCGGCCGGATTGATTTCCACT GGACACTGGTTCAGCCTGGCGACAATATCACATTCAGCCATAACGGTGGTTTGATTGCTCCTTCCAGAGTCAGTAAATTGATCGGGAGAGG ACTGGGGATTCAGTCAGACGCCCCCATTGATAATAACTGTGAAAGCAAGTGCTTTTGGAGAGGTGGATCCATCAACACACGTCTGCCCTTC CAGAACCTGAGTCCCCGGACAGTCGGTCAGTGTCCGAAGTACGTCAACAAACGGAGTTTAATGCTCGCGACTGGGATGAGGAACGTGCCT GAACTTATGCAAGGCCGGGGCTTATTCGGGGCCATTGCGGGCTTTCTGGAAAACGGGTGGGAAGGGATGGTTGATGGGTGGTACGGGTTC AGACACCAGAATGCTCAGGGCACCGGACAAGCCGCAGATTATAAGTCAACCCAGGCCGCAATAGATCAGATTACCGGCAAACTGAACAGG CTCGTCGAGAAAACCAATACTGAGTTTGAATCTATTGAAAGCGAGTTTTCAGAGATTGAACATCAAATCGGCAATGTGATTAATTGGACCAAA GATTCGATTACAGATATATGGACCTACCAGGCTGAACTTCTTGTCGCCATGGAAAACCAGCATACGATCGATATGGCCGACAGCGAAATGTT AAACCTATACGAGCGCGTGAGGAAGCAGCTGAGGCAGAACGCAGAAGAGGACGGGAAAGGCTGTTTTGAAATATACCACGCCTGTGATGA TTCTTGCATGGAGTCGATCCGTAACAACACATACGACCATTCTCAATATCGAGAGGAAGCATTATTGAACAGGTTAAACATCAATCCCGTTAC TCTCTCCAGCGGATATAAAGACATTATCCTTTGGTTTTCATTCGGAGCATCCTGTTTTGTCTTACTAGCCGTCGTAATGGGTTTAGTCTTCTTT TGCCTTAAGAACGGGAATATGAGGTGTACCATATGTATT

3017	ATGTATAAAATCGTGGTGATCATCGCTCTTCTTGGCGCTGTGAAGGGGCTCGACAAGATCTGCTTAGGTCATCACGCAGTGGCTAATGGTAC GATCGTGAAGACACTGACGAATGAACAGGAAGAAGTCACAAATGCCACGGAGACCGTTGAGAGCACTGGCATCAATAGACTGTGCATGAAA GGACGCAAGCACAAGGACCTGGGCAATTGCCATCCTATCGGCATGCTGATTGGAACGCCGGCGTGTGACCTGCATTTGACCGGGATGTGG GATACCCTAATCGAGAGGGAGAACGCAATTGCCTATTGTTACCCCGGCGCAACCGTGAATGTGGAAGCCCTGCGACAAAAGATCATGGAGA GTCGTGGCATTAATAAGATTTCTACCGGGTTCACTTATGGCAGTAGCATCAACTCTGCCGGCACAACCAGGGCTTGTATGAGGAATGGAGG CAACAGCTTTTATGCTGAACTTAAATGGCTCGTCAGCAAGTCTAAAGGTCAGAACTTCCCTCAGACCACAAATACATACAGAAATACTGACAC AGCCGAGCACCTCATCATGTGGGGCATCCATCATCCATCTTCCACCCAAGAGAAAAACGACCTCTACGGGACACAGAGCCTGTCTATAAGC GTGGGCTCTAGCACTTATAGAAATAATTTCGTTCCAGTGGTGGGCGCACGTCCCCAAGTTAACGGCCAGAGCGGAAGGATCGATTTCCATT GGACTCTGGTGCAGCCTGGGGACAACATTACATTTTCCCACAATGGAGGTCTTATTGCACCCTCACGGGTATCCAAGTTAATCGGCCGCGG ACTTGGCATCCAATCAGATGCACCCATCGACAACAATTGTGAGAGTAAGTGCTTCTGGCGTGGTGGGAGCATCAACACGAGATTACCTTTCC AGAATCTGAGCCCGAGAACAGTAGGGCAATGTCCAAAGTATGTGAACAGGCGCAGCCTCATGCTCGCGACCGGAATGAGAAATGTTCCCG AACTGATACAAGGAAGGGGTTTGTTCGGCGCCATCGCAGGATTTCTGGAAAACGGGTGGGAGGGAATGGTGGACGGTTGGTACGGCTTCC GTCACCAGAACGCCCAGGGCACGGGTCAGGCCGCAGACTATAAAAGTACCCAGGCGGCCATTGATCAGATAACAGGCAAGCTCAACCGGC TGGTAGAAAAAACCAATACAGAGTTTGAGAGCATAGAGTCCGAGTTTAGCGAGATAGAACATCAAATAGGAAATGTCATCAACTGGACCAAG GACAGCATCACGGATATTTGGACTTACCAGGCCGAATTACTGGTTGCTATGGAGAACCAGCACACTATTGATATGGCAGATTCTGAAATGTT AAACTTATATGAACGCGTGAGAAAGCAGTTGAGGCAGAATGCTGAGGAGGACGGGAAGGGCTGCTTTGAAATATACCATGCCTGCGACGAT AGCTGTATGGAGAGCATCAGGAACAATACATACGATCACAGTCAATACCGCGAGGAGGCTTTACTCAACCGGCTCAACATAAACCCCGTCA CTCTCTCTTCCGGCTACAAGGATATTATTCTGTGGTTTTCATTCGGCGCTTCCTGCTTCGTTCTCCTAGCCGTAGTAATGGGCCTCTTCTTCT TCTGCCTGAAAAACGGCAACATGCGGTGTACTATCTGCATT
3018	ATGTACAAGATTGTGGTCATCATCGCCCTGCTGGGCGCCGTCAAGGGGCTCGATAAGATTTGCCTAGGTCACCACGCCGTCGCCAATGGC ACGATTGTTAAAACACTGACCAATGAACAGGAGGAAGTGACTAACGCTACGGAAACAGTGGAATCTACAGGGATTAACAGACTTTGCATGAA GGGGCGCAAGCATAAGGACCTTGGAAATTGCCACCCAATCGGAATGCTTATCGGAACACCTGCTTGTGACCTTCATCTGACCGGCATGTGG GACACCCTGATAGAGAGGGAGAACGCCATCGCCTACTGCTATCCTGGAGCCACCGTCAATGTCGAAGCCTTAAGACAGAAAATAATGGAAA GTGGGGGTATAAACAAAATCAGCACTGGCTTCACTTATGGTTCTTCTATCAATAGCGCAGGGACAACCAGAGCCTGCATGCGCAATGGGGG CAATAGCTTTTATGCCGAACTGAAGTGGCTGGTCTCTAAATCTAAGGGTCAAAATTTTCCTCAGACAACCAACACATACCGCAATACGGACA CCGCGGAGCATCTCATTATGTGGGGGATTCACCACCCTTCAAGCACCCAGGAGAAGAATGACCTCTATGGGACACAGAGCCTGTCAATCTC AGTGGGATCTAGCACTTATCGTAACAATTTCGTTCCAGTAGTTGGTGCACGCCCCCAGGTTAATGGGCAGAGTGGGCGAATTGATTTCCACT GGACGCTCGTGCAGCCCGGCGATAATATCACATTCAGTCACAATGGGGGGCTAATCGCCCCGTCCAGAGTTTCTAAGCTGATCGGCAGAG GATTAGGAATTCAATCCGACGCACCCATTGACAACAATTGTGAATCAAAATGCTTCTGGCGCGGAGGGTCGATAAACACACGACTTCCTTTT CAAAACTTATCCCCAAGAACCGTGGGACAGTGCCCCAAGTACGTCAATAGAAGATCACTGATGCTGGCCACCGGTATGAGGAATGTGCCCG AACTAATCCAGGGCCGCGGGTTGTTCGGCGCTATTGCAGGCTTTTAGAAAACGGCTGGGAAGGGATGGTCGACGGATGGTACGGGTTCA GGCATCAGAATGCGCAGGGCACTGGGCAGGCCGCGGATTATAAGAGTACCCAGGCGGCCATAGATCAAATAACCGGCAAGCTCAACAGGC TGGTCGAGAAGACAAACACAGAGTTTGAGTCAATAGAGTCCGAATTCTCCGAAATCGAACACCAAATAGGCAATGTGATAAACTGGACCAAG GACTCCATCACCGATATCTGGACTTATCAGGCTGAACTGTTAGTGGCGATGGAAAACCAGCACACAATCGATATGGCAGACTCAGAGATGC TTAATCTGTATGAACGAGTTAGAAAGCAGCTCCGGCAGAATGCCGAGGAAGACGGAAAGGGCTGCTTCGAGATCTACCACGCGTGCGATG ATTCCTGTATCGAGTCTATTCGCAATAACACGTATGATCATTCACAGTATAGAGAAGAGGCCCTCCTAAACAGGCTCAACATCAACCCTGTG ACGCTGTCGAGCGGCTATAAGGATATTATACTGTGGTTTAGTTTCGGCGCCAGTTGCTTTGTCCTTCTTGCCGTGGTAATGGGATTATTCTTC TTCTGTCTGAAAAACGGAAACATGAGATGCACAATATGTATC
3019	ATGTATAAAATTGTAGTCATTATTGCCTTGTTAGGTGCAGTGAAGGGCTTAGACAAAATATGCCTAGGTCATCACGCCGTCGCTAATGGTACA ATAGTCAAGACCCTCACCAATGAACAGGAGGAGGTCACGAACGCAACAGAGACCGTGGAGTCCACAGGAATAAACCGACTGTGTATGAAG GGCAGAAAACATAAAGACTTGGGCAACTGTCACCCTATAGGCATGTTGATTGGCACACCAGCTTGTGATCTCCACCTAACCGGGATGTGGG ACACCCTCATAGAACGCGAGAACGCTATCGCCTATTGTTATCCCGGGGCCACGGTTAATGTGGAAGCTCTGCGACAGAAGATCATGGAGAG CGGGGGGATTAATAAGATAAGCACTGGTTTCACCTACGGCTCAAGTATCAATTCAGCTGGCACCACCCGAGCATGTATGAGGAATGGCGGA AATAGCTTTTATGCAGAACTGAAGTGGTTAGTCAGTAAAAGTAAAGGTCAGAACTTTCCACAGACCACCAACACATATCGCAATACTGATACA GCTGAGCACTTAATCATGTGGGGAATTCACCATCCCAGTTCCACACAAGAAAAAAATGATTTATACGGCACTCAGTCCCTTTCAATCAGCGT GGGGTCCTCTACCTACAGGAATAACTTTGTCCCCGTGGTTGGCGCCAGACCCCAGGTAAACGGTCAGTCTGGACGGATCGATTTCCACTGG ACACTGGTGCAGCCAGGTGACAATATTACCTTCAGCCACAATGGGGGCCTGATCGCCCCCTCAAGAGTGTCCAAGCTGATCGGAAGAGGG TTGGGGATCCAGTCCGATGCCCCAATCGACAATAACTGCGAATCCAAATGTTTCTGGCGCGGAGGCTCAATAAATACCAGACTCCCTTTCCA GAACCTTTCACCAAGAACAGTCGGACAATGCCCTAAGTATGTCAATCGCCGATCGCTAATGCTTGCGACAGGGATGAGAAATGTGCCAGAG CTGATTCAAGGGCGCGGGCTGTTCGGCGCTATCGCCGGCTTTTTAGAGAATGGTTGGGAAGGGATGGTGGATGGCTGGTATGGATTTAGG CACCAGAACGCTCAGGGGACAGGGCAGGCTGCCGACTATAAGAGCACACAGGCAGCCATTGACCAAATCACTGGGAAATTGAACCGTCTT GTGGAAAAAACTAACACCGAGTTCGAGTCCATTGAGTCCGAGTTCTCGGAGATAGAACACCAAATCGGGAACGTCATCAATTGGACAAAAG ATTCCATCACCGATATCTGGACTTACCAGGCTGAGTTGCTGGTCGCTATGGAGAACCAGCACACGATCGACATGGCAGACTCTGAGATGTT GAACCTGTATGAGAGAGTGCGAAAACAGCTCAGACAGAACGCTCAAGAAGACGGAAAGGGGTGCTTCGAGATCTATCATGCCTCTGACGA TAGCTGTATGGAGAGCATAAGAAACAATACGTATGATCACTCTCAGTACAGAGAAGAGGCCCTGCTAAATCGGTTAAACATTAATCCAGTGA CCCTGAGCAGCGGCTACAAGGACATCATCCTATGGTTTTCTTTCGGTGCCTCGTGCTTCGTCTTACTCGCTGTCGTCATGGGTCTGTTCTTC TTTTGCCTGAAGAACGGCAATATGCGGTGTACAATCTGTATT

3020	ATGTATAAAGTGGTCGTGATTATTGCTCTGTTAGGCGCCGTGCGCGGCTTGGATAAGATCTGCCTAGGCCATCACGCCGTCGCTAACGGCA CTACCGTAAAGACCCTGACCAACGAGCAGGAAGAAGTCACAAATGCTACCGAGACCGTTGAGAGCACTTCGTTGAACAAGTTATGCATGAA GGGCAGACGCTACAAGGATCTGGGTAATTGCCACCCCATAGGGATGCTCATTGGAACGCCCGTCTGTGACTTGCATTTGACCGGCACATG GGATACCCTCATCGAGCGGGAAAACGCCACAGCATACTGCTATCCTGGGGTGACTATCAACGAGGAAGCATTGAGGCAAAAGATCATGGAA TCAGGCGGGATCAGTAAGATGAGAACCGGGTTCACCTACGGCCTTCCATCAACTCCGCCGGCACCACCCGCAGTTGTATGAGAAACGGC GGCAATTCTTTTTATGCTGAGCTGAAGTGGTTGGTCTCTGGGACCAAGGGCCAAAATTTTCCTCAAACAACGAATACCTACCGGAATACTGA CACGGCTGAACATCTGATCATCTGGGGGATTCACCACCCCTCATCCACCCAAGAGAAGAATGATCTGTATGGAACGCAGAGCCTGTCTATTT CGGTTGGAAGCAGTACATACCAGAATAACTTCGTGCCCGTTATTGGGGCCCGTCCCCAGGTCAACGGTCAGTCGGGCCGGATTGAGTTCC ATTGGACCCTGGTCCGCCCTGGTGACAACATTACTTTTTCGCATAACGGTGGGCTAATCGCACCTGATCGAGTGTCCAAGCTGATCGGCAA AGGAATTGGTATACAGTCTGGTGCGGTAATCGATAAGGACTGTGAAAGCAAATGCTTCTGGAGAGGCGGCAGTATCATCACCGAACTCCCC TTTCAGAATCTCTCTCCACGTACAGTGGGCCAGTGTCCAAAGTATGTTAAAAAGCGGTCTCTGTTACTAGCCACGGGGATGAGGAATGTTCC CGAGGTGGTACAAGGACGGGGCCTCTTCGGGGCGATCGCCGGCTTCATTGAAAACGGCTGGGAGGGTATGGTTGACGGATGGTACGGAT TCCGCCACCAAAACGCCCAGGGAATCGGCCAGGCTGCCGACTACAAATCCACTCAGACCGCCATCGACCAAATTACGGGTAAACTGAACC GCCTAATCGAGAAGACAAATACTGAATTTGAGAGCATTGAGAGTGAGTTCAGCGAAATCGAGCACCAGATCGGAAATGTGATTAATTGGACA AAGGATAGTATTACCGACATCTGGACCTACCAGGCGGAGCTCCTTGTGGCTATGGAAAACCAACACACCATTGACATGGCCGACAGTGAAA TGCTGAATCTCTACGAGCGCGTGAGAAAGCAGCTCCGGCAGAACGCCGAGGAGGACGGCAAAGGATGCTTCGAAATATATCACACATGTG ACAACTCATGCATGGAAAGTATTAGGAACAACACCTACGACCATTCCCAATACAGAGAGGAGGCCCTACTTAACCGCCTTAACATCAATCCC GTGAAACTCTCCTCCGGTTATAAAGATATCATTTTATGCTTCAGCTTCCGGGCCGAGTTGCTTTGTGCTGCTGGCTGTGATTATGGGCCTGGG ATTCTTTTGCCTAAAAAACGGAAACATGCGATGCACTATCTGTATC
3021	ATGTATAAAATTATTGTGATCATTGCACTGCTGGGGGCCGTTAAGGGTCTGGACAAAATTTGTCTAGGACACCACGCTGTGGCAAACGGTAC TATCGTGAAAACATTGACCAACGAACAGGAGGAGGTTACCAACGCCACCGAGACAGTCGAAAGTACCGGAATTAATCGCCTCTGCATGAAA GGCCGAAAGCATAAGGACCTGGGGAATTGCCATCCCATCGGAATGCTGATCGGCACCCCAGCCTGTGACTTGCATCTTACAGGAACCTGG GACACACTAATCGAAAGGGAAAACGCCATTGCTTACTGTTACCCGGGCGCTACAGTTAACGTGGAGGCCCTTAGGCAGAAAATCATGGAGT CCGGCGGGATCGACAAGATCAGTACCGGCTTCACCTATGGGTCAAGCATCAATTCTGCGGGCACTACGAGGGCATGCATGAGGAACGGAG GCAACTCATTCTATGCGGAATTGAAGTGGCTGGTCTCCAAGTCCAAAGGGCAGAACTTCCCTCAAACTACTAACACCTACCGGAATACAGAC ACAGCGGAACACCTCATTATGTGGGGCATCCACCACCCTTCTAGTACTCAAGAGAAGAATGATCTCTACGGCACTCAGTCCCTGAGTATTAG TGTCGGCTCAAGCACCTATAGAAACAATTTCGTGCCCGTTGTGGGGGCCCGCCCTCAGGTTAATGGTCAATCAGGGCGAATTGATTTTCAC TGGACCCTGGTCCAGCCCGGGGACAACATAACATTCAGCCATAATGGGGGTCTGATTGCCCCCAGCCGGGTGAGCAAGTTGATAGGGAGG GGACTTGGCATCCAGAGTGACGCCCCTATTGACAACAACTGCGAAAGCAAATGCTTTTGGCGGGGTGGCTCAATAAACACAAGGCTACCCT TCCAAAATTTGAGTCCAAGGACCGTGGGTCAATGCCCTAAATACGTCAATCGCCGCTCCTTGATGCTCGCCACTGGGATGCGCAACGTGCC TGAGCTTATTCAGGGTCGGGGTCTGTTCGGGGCTATCGCCGGGTTCCTTGAAAATGGCTGGGAGGGTATGGTAGACGGCTGGTACGGATT TAGACACCAGAACGCCCAGGGAACGGGCCAGGCGGCAGACTACAAGTCTACCCAAGCTGCAATCGATCAGATCACTGGCAAGTTAAACAG ACTGGTGGAAAAGACAAATACCGAATTTGAAAGCATTGAATCGGAATTCTCAGAGATCGAGCACCAGATTGGAAATGTGATTAATTGGACGA AAGACAGTATTACCGACATCTGGACTTATCAGGCAGAACTGCTGGTTGCCATGGAAAACCAGCATACCATAGATATGGCAGATTCTGAAATG CTCAATCTTTACGAGCGCGTACGGAAACAGTTAAGACAGAACGCAGAGGAGGACGGGAAAGGTTGCTTTGAAATATACCACGCATGTGACG ATTCTTGCATCGAAAGTATCCCCAATAATACGTATGATCACAGTCAATACCGACGAAGAGGCTCTCTTGAATCGTTTGAATATCAATCCAGTGA CCCTGTCCTCCGGATATAAGGATATAATTCTGTGGTTCAGTTTTGGGGCAAGTTGCTTCGTCCTCCTCGCTGTCGTTATGGGACTGTTTTTTT TTTGTCTAAAGAACGGAAACATGCGCTGCACTATATGCATT
3022	ATGTATAAAATAGTTGTCATCATTGCACTCTTGGGCGCCGTGAAAGGACTGGATAAAATATGTCTCGGCCACCATGCAGTTGCAAACGGGAC AATTGTGAAAACATTGACCAACGAGCAGGAGGAGGTGACCAATGCTACAGAGACGGTGGAATCTACTGGCATCAACCGATTGTGCATGAAA GGTAGCAAACACAAGGATTTGGGTAATTGCCATCCGATTGGGATGCTTATCGGAACCCCAGCTTGCGATCTGCATTTGACCGGAATGTGGG ACACCCTGATTGAGAGGGAAAACGCCATCGCCTACTGCTATCCTGGCGCAACAGTTAACGTGGAGGCACTTCGCCAGAAGATCATGGAAAG CGGGGGGATAAACAAGATCTCTACAGGTTTCACCTATGGTTCTTCCATTAACAGCGCCGGTACCACACGAGCTTGTATGCGAAATGGGGGC AACTCCTTCTACGCTGAATTAAAATGGCTGGTGTCGAAGTCCAAAGGGCAGAACTTCCCTCAGACAACAAACACCTACCGGAACACAGACAC CGCGGAGCACCTGATCATGTGGGGTATACACCACCCCAGCTCTACCCAGGAAAAGAATGATCTCTATGGAACCCAGTCATTAAGTATCAGC GTCGGTTCCTCTACATATCGTAATAATTTCGTGCCTGTGGTTGGCGCCAGGCCACAAGTGAATGGGCAGTCCGGCAGAATTGACTTTCATTG GACTCTGGTGCAGCCAGGCGACAATATCACATTCTCTCACAACGGGGGACTTATCGCCCCCTCCAGGGTGTCAAAACTCATCGGCCGGGG GCTGGGAATCCAAAGCGATGCTCCCATCGACAACAATTGCGAATCAAAATGTTTCTGGCGAGGGGGTTCCATAAATACCAGACTGCCATTC CAGAACCTGTCCCCAAGAACAGTGGGCCAATGTCCCAAGTATGTGAACAGAAGGTCTTTGATGCTCGCAACTGGGATGCGAAATGTTCCAG AGCTGATTCAGGGTCGAGGACTTTTTGGAGCCATCGCTGGCTTCTTAGAGAACGGCTGGGAAGGAATGGTGGACGGCTGGTATGGCTTTA GACACCAGAACGCCCAGGGTACGGGACAGGCTGCAGATTACAAATCTACACAGGCCGCAATCGACCAGATCACAGGCAAGCTCAATCGGC TTGTGGAGAAGACAAACACAGAGTTTGAGAGCATTGAATCGGAATTTAGTGAGATTGAGCACCAGATTGGCAATGTGATTAATTGGACTAAA GACTCTATTACGGATATCTGGACGTATCAGGCAGAGCTGTTGGTGGCAATGGAAAATCAACACACTATCGACATGGCCGATAGTGAGATGC TCAATCTATATGAGAGGGTTCGCAAGCAGCTGCGTCAAAATGCCGAGGAGGATGCCAAGGGATGTTTCGAGATTTATCACGCCTGCGATGA TTCGTGCATGGAGAGCATTCGCAATAACACGTACGATCACTCTCAGTACAGGGAGGAGGCACTTTTAAACCGTCTCAACATCAACCCTGTGA CCCTGAGCTCTGGCTACAAAGACATCATTTTGTGGTTCTCCTTTGGAGCCAGCTGCTTCGTGCTCCTCGCCGTGGTCATGGGCTTGTTTTTC TTCTGTCTGAAGAATGGAAATATGCGATGCACCATCTGTATT

3023	ATGTACAAGGTGGTAGTTATAATCGCACTTCTGGGAGCAGTGAGAGGTCTCGACAAGATTTGCCTCGGGCACCACGCTGTGGCTAATGGCA CAATTGTGAAGACCTTGACAAATGAGCAGGAAGAAGTGACAAATGCGACTGAGACTGTCGAGTCCAAGTCTCTGGGCAAGTTGTGCATGAA GGGCAGATCCTACAACGACTTAGGGAACTGTCATCCGATAGGGATCTTAATCGGGACCCCAGCCTGCGATCTCCACCTGACTGGCACCTG GGATACCTTAATCGAGAGGGAAAACGCAGTGGCCTACTGCTATCCCGGTGCTACCGTGAATGAGGAGGCTCTGAGGCAGAAGATCATGGA GTCAGGGGCGAATTTCGAAAATAAGCACCGGGTTCACTTACGGTACCAGCATCAATTCTGCTGCAACAACTAAGGCATGTATGCGTAATGGC GGAAACTCATTTTACGCAGAGCTCAAGTGGCTAGTGAGCAAGAATAAAGGGCAGAACTTTCCGCAGACCACTAACACTTACAGAAACACAGA TACCGCTGAACATTTGATTATCTGGGGCATTCACCACCCGTCGTCAACACAGGAGAAGAATGATCTGTATGGTACACAGTCACTGTCAATTT CTGTCGGATCTTCCACCTACCAGAATAATTTCGTTCCAGTGGTTGGTGCTAGGCCCCAGGTTAACGGACAATCCGGTCGCATTGACTTCCAT TGGACCCTGCTGCAGCCAGGCGATAACATTACGTTCTCGCATAACGGCGGCCTAATCGCACCTAGCAGAGTGTCAAAGTTGATTGGGAGG GGCCTTGGCATCCAGAGCGAGGCCCCCATCGACAATGGCTGCGAGTCCAAATGCTTCTGGAAAGGGGGGTCAATCAACACCAAGCTGCCC TTTCAGAACCTCAGCCCCCGAACCGTCGGGCAGTGCCCTAAATACGTGAATAAGCGCAGTCTGATGTTAGCTACAGGGATGCGGAATGTAC CCGAAATCATGCACGGGCGAGGTCTGTTCGGGGCAATTGCAGGCTTCATTGAAAATGGGTGGGAAGGCATGGTCGATGGGTGGTATGGGT TTAGACACCAAAACGCACAAGGAACAGGCCAGGCAGCGGACTATAAGAGCACCCAGGCAGCTATTGACCAGATTACCGGCAAGCTGAACA GGCTTATAGAGAAGACTAACACAGAATTCGAGTCCATTGAGTCAGAGTTCAGCGAGATTGAGCACCAGATCGGAAACATCATTAACTGGACT AAGGACTCAATTACAGATATATGGACGTATCAGGCCGAACTGTTGGTGGCAATGGAGAATCAGCATACCATTGATATGGCAGATAGTGAGAT GCTGAATCTTTACGAACGCGTTCGCAAACAGTTGCGGCAGAATGCCGAAGAAGACGGCAAGGGTTGCTTTGAGATATACCACGCTTGTGAC GATTCCTGCATGGAGTCTATCAGAAACAACACGTACGATCACAGTCAGTATCGCGAGGAGGCATTACTCAATCGACTAAATATCAACCCGGT GAAACTGTCTAGCGGATACAAAGACATCATCCTTTGGTTTTCCTTTGGCGCAAGTTGTTTGATTCTGTTGGCCGTGGTTATGGGTTTAGTCTT TTTTTGTCTCAAAAACGGTAATATGCGCTGTACGATCTGCATT
3024	ATGTACAAGATCGTGGTCATCATCGCCTTGTTGGGGGCAGTGAAAGGATTGGATAAAATTTGCCTGGGACACCACGCCGTCGCCAATGGAA CCATAGTCAAGACGCTGACAAACGAGCAGGAGGAAGTAACCAATGCGACTGAAACCGTGGAATCAACCGGAATCAATCGACTGTGTATGAA AGGCAGGAAGCATAAAGACCTGGGCAACTGTCACCCGATCGGGATGTTAATCGGGACACCCGCTTGTGACTTGCACCTGACCGGCATGTG GGACACTCTGATTGAGCGAGAGAATGCAATCGCCTATTGTTACCCCGGAGCCACTGTTAACGTTGAGGCCCTCAGACAGAAGATTATCGAG TCTGGCGGGATCAACAAGATCTCCACAGGGTTTACTTACGGAAGCAGCATCAACTCTGCCGGCACTACTCGCGCTTGCATGCGTAACGGGG GTAACAGCTTCTATGCTGAACTCAAATGGCTGGTGTCTAAAAGTAAGGGCCAGAACTTTCCGCAGACGACTAATACATATAGAAATACCGAT ACAGCCGAGCACCTGATTATGTGGGGGATACACCACCCCAGTTCCACCCAGGAAAAGAACGACCTATATGGAACCCAGTCCCTCTCTATTA GCGTCGGCTCTTCAACTTATCGTAATAATTTTGTGCCGGTCGTCGGAGCAAGACCTCAGGTCAACGGGCAGTCCGGTCGAATAGATTTTCA CTGGACCCTAGTTCAGCCAGGAGATAATATCACATTCAGCCACAACGGAGGGTTGATAGCTCCATCACGAGTGTCCAAGCTGATCGGCAGG GGGCTCGGCATCCAGTCCGACGCCCCAATTGATAACAACTGTGAATCCAAGTGCTTCTGGAGAGGAGGCTCTATCAATACCCGATTGCCGT TCCAAAATCTATCCCCTAGAACAGTCGGGCAGTGCCCTAAATACGTGAATAGAAGAAGCTTAATGTTGGCAACAGGAATGAGGAACGTGCC GGAACTCATCCAGGGGCGCGGCTTATTCGGCGCTATCGCAGGTTTTCTTGAGAATGGATGGGAGGGCATGGTTGATGGTTGGTACGGATT CAGACACCAAAACGCTCAAGGCACCGGCCAAGCTGCCGATTATAAAAGCACTCAAGCTGCTATCGACCAGATTACCGGAAAACTTAATAGG CTGGTTGAGAAAACGAACACCGAATTCGAGAGCATCGAGAGCGAATTTAGTGAGATCGAGCATCAAATAGGCAACGTGATCAATTGGACGA AAGATTCTATCACCGATATTTGGACTTATCAAGCCGAACTTCTAGTGGCAATGGAGAACCAGCACACAATCGACATGGCAGATTCGGAAATG CTGAATTTATACGAGAGGGTGCGTAAACAGTTAAGGCAAAACGCCGAAGAGGACGGTAAAGGTTGCTTTGAAATTTATCACGCGTGCGATG ATTCCTGTATCGAGAGTATTCCGAATAATACATATGATCACTCCCAATATCGCCGAAGAGGCATTCTTAAACCGGCTGAATATAAATCCAGTAA CCCTGTCCTCCGGGTACAAAGACATAATCCTGTGGTTCTCCTTCGGGGCTAGTTGTTTTGTTCTGCTGGCTGTGGTTATGGGATTATTCTTTT TTTGTCTGAAAAACGGGAATATGCGCTGTACAATATGCATA
3025	ATGTACAAAATTGTAGTGATTATCGCTCTTCTAGGCGCCGTAAAGGGGCTTGACAAGATCTGTCTAGGACACCACGCCGTGGCCAATGGAA CCATTGTAAAAACACTGACGAACGAGCAGGAGGAGGTGACAAACGCCACTGAAACAGTTGAGAGCACCGGGATTAACCGTCTCTGTATGAA GGGTACGAAACACAAAGATCTCGGGAATTGTCATCCAATTGGAATGCTGATAGGCACTCCCGCCTGTGACCTTCACTTAACTGGCATGTGG GACACACTGATCGAGAGGGAGAATGCAATTGCGTATTGTTACCCGGGCGCCACTGTTAATGTGGAGGCGCTGAGGCAGAAAATAATGGAG TCCGGCGGAATAAATAAAATCTCTACGGGGTTCACCTACGGATCCTCCATCAACTCTGCAGGCACAACAAGAGCCTGCATGAGAAACGGGG GCAACAGTTTTTACGCCGAGCTGAAATGGTTAGTCAGTAAATCTAAAGGGCAAAATTTCCCTCAGACGACCAACACCTACCGGAACACCGAT ACCGCTGAGCATCTCATCATGTGGGGTATCCATCACCCTTCCAGCACTCAGGAGAAAAACGACCTGTACGGTACTCAATCCCTCTCTATCTC AGTCGGAAGCTCCACATACCGCAATAATTTCGTGCCTGTGGTCGGGGCCCGCCCGCAGGTTAACGGACAGTCAGGACGGATTGACTTCCA CTGGACTCTGGTGCAGCCCCGGTGACAACATTACGTTTAGCCATAACGGTGGTCTCATTGCCCCGTCGCGTGTATCCAAGCTGATTGGCCGG GGGTTAGGGATTCAAAGTGACGCACCCATCGACAACAATTGCGAATCGAAGTGTTTTTGGAGAGGCGGCAGTATTAATACCCGGTTGCCAT TCCAAAACCTCTCCCCAAGAACAGTTGGCCAGTGTCCAAAATATGTCAACCGGCGGTCCCTTATGCTCGCCACCGGCATGCGAAACGTACC CGAACTGATCCAAGGCCGCGGTCTGTTTGGAGCTATCGCAGGCTTCCTGGAGAACGGGTGGGAGGGCATGGTCGACGGATGGTATGGTTT TAGGCACCAGAATGCACAAGGGACGGGTCAAGCCGCCGATTACAAGAGCACACAGGCGGCCATCGACCAAATCACTGGGAAACTGAATAG ACTGGTGGAGAAGACCAACACTGAGTTCGAAAGCATCGAATCAGAATTTAGCGAAATAGAACACCAGATCGGCAATGTGATTAACTGGACAA AGGATAGCATAACAGACATCTGGACATATCAGGCTGAACTGCTTGTCGCGATGGAGAACCAGCACACCATAGATATGGCAGATAGTGAAAT GCTCAATTTATACGAGCGGGTGAGAAAACAGCTTAGACAGAACGCTGAAGAGGACGCCAAGGGGTGCTTTGACATATACCACGCCTGCGA CGACAGCTGCATGGAAAGTATCAGGAACAATACATATGACCACTCACAGTATCGCGAAGAGGCACTTCTCAATAGACTAAACATCAATCCCG TGACACTGAGTAGTGGATATAAAGATATTATATTATGGTTCTCTTTTGGGGCGTCATGTTTCGTGCTTCTGGCCGTGGTGATGGGTCTCTTCT TCTTCTGTCTTAAAAATGGCAATATGCGGTGCACCATTTGCATA

3026	ATGTACAAAATCGTACTGGTGCTGGCATTGCTCGGGGCAGTGCACGGCCTGGACAAGATATGCCTGGGACACCACGCTGTGCCTAACGGT ACCATTGTGAAGACTCTGACGAATGAGAAGGAAGAGGTGACGAACGCAACGGAGACGGTCGAGTCCAAGAGCCTCGATAAGCTGTGTATG AAGAACAGGAACTACAAGGACCTCGGGAATTGTCATCCAATTGGCATGGTGGTGGGCACTCCCGCCTGCGACCTCCATCTGACTGGTACAT GGGATACCCTAATAGAGCGGGACAATAGTATCGCCTACTGTTATCCCGGCGCCACCGTGTCAGAGGAGGCCCTGCGACAGAAAATAATGG AGTCCGGCGGCATTGATAAGATTTCCACTGGGTTTACATACGGTAGCTCCATTAATTCTGCGGGAACTACCAAGGCTTGTATGAGAAATGGC GGTAACTCTTTCTATAGCGAGTTAAAATGGCTGGTGAGCAAGAATAAGGGCCAAAATTTCCCCCAGACTACGAACACCTACAGAAACACCGA CAGTGTTGAGCACCTGATTATATGGGGGATACATCATCCCAGTTCTACTCAGGAGAAAAACGATTTATACGGCACTCAAAGTCTGAGTATCT CTGTGGGGTCTAGCACATACCAGAACAACTTCGTCCCAGTGGTTGGGGCAAGGCCGCAGGTGAATGGACAATCAGGGAGAATCGATTTCC ACTGGACAATGGTACAGCCTGGGGATAATATCACCTTTTCTCATAACGGCGGACTTATAGCCCCCAACCGCGTGAGCAAGCTAAAGGGAAG AGGCCTAGGAATTCAAAGCGGTGCTTCGGTGGACAATGACTGCGAGAGCAAGTGCTTCTGGAAAGGTGGCTCCATCAATACTAAGTTGCCT TTTCAGAACTTGAGCCCGAGAACCGTTGGCCAGTGCCCTAAATATGTGAACAAAAAGTCACTCCTGCTCGCCACAGGGATGAGGAATGTCC CCGAAGTGGCTCAGGGTCGCGGGCTCTTCGGAGCCATCGCCGGATTCATCGAGAACGGCTGGGAAGGGATGGTAGACGGCTGGTATGGC TTTAGGCATCAAAACGCCCAGGGGACAGGGCAAGCCGCTGACTATAAGAGTACCCAGGCAGCCATCGACCAGATCACTGGGAAATTAAATC GGCTGATTGAAAAGACGAACACCGAATTTGAAAGCATCGAGTCAGAGTTCTCCGAGATAGAGCATCAAATTGGGAACGTGATTAACTGGAC GAAGGATAGCATCACCGATATTTGGACATACCAGGCGGAGCTGCTGGTGGCCATGGAGAATCAACATACAATCGACATGGCCGACTCTGAG ATGTTGAATCTCTACGAAAGGGTGAGAAAGCAGTTAAGGCAGAATGCCGAGGAGGATGGGAAAGGATGCTTCGAGATCTATCACAAATGCG ATGACAATTGTATGGAATCTATTCGTAACAATACTTACGACCACACTCAGTATCGTGAGGAAGCGCTGTTGAATCGGCTCAATATCAATCCAG TCAAGCTGTCATCTGGCTACAAAGATGTGATATTATGGTTTTCTTTTGGCGCATCTTGCTTCGTGCTGCTGGCTGTCATTATGGGACTTGTAT TTTTTTGTTTGAAGAATGGAAACATGCGCTGCACAATCTGCATC
3027	ATGTATAAGATAGTGGTCATAATTGCCCTGCTGGGGGCGGTGAAGGGCCTGGACAAGATTTGTCTGGGTCATCATGCTGTGGCAAACGGTA CGATCGTTAAGACCCTGACGAATGAACAAGAGGAGGTGACTAACGCTACCGAAACCGTGGAGTCCACCGGCATAAACCGCCTTTGCATGAA GGGGAGGAAGCACAAGGACCTCGGTAACTGTCACCCAATCGGCATGTTAATTGGCACACCAGCTTGTGATCTGCACCTGACCGGAATGTG GGATACACTCATTGAAAGAGAAAACGCTATCGCGTATTGCTACCCTGGAGCAACTGTTAATGTCGAGGCTTTGCGCCAAAAAATAATGGAAA GTGGCGGTATCAATAAAATCTCAACAGGGTTTACCTACGGGAGCAGTATCAACTCAGCCGGAACAACCCGGGCCTGCATGCGCAATGGCG GCAACTCTTTTTACGCTGAACTGAAATGGTTGGTGTCAAAATCAAAGGGTCAAAACTTTCCTCAAACTACAAATACATATCGGAACACCGATA CAGCTGAGCACTTAATAATGTGGGGGATCCACCACCCTTCTTCAACTCAGGAAAAGAATGATCTGTATGGCACTCAGTCACTCTCCATTAGC GTAGGGTCATCTACCTATAGAAATAATTTCGTACCAGTGGTAGGTGCAAGGCCCCAGGTGAACGGCCAGAGTGGCAGAATAGACTTCCACT GGACTCTTGTTCAGCCAGGGGATAACATTACATTCTCGCACAACGGCGGCCTGATAGCCCCATCTAGGGTGTCTAAGCTAATTGGCAGAGG TCTGGGCATCCAATCCGACGCACCAATTGACAACAATTGCGAAAGTAAGTGTTTCTGGAGAGGAGGGTCAATTAATACAAGATTGCCGTTTC AAAATCTGTCCCCGAGAACAGTGGGACAGTGCCCAAAATATGTAAATAGACGGTCACTGATGTTAGCTACCGGCATGCGAAACGTCCCCGA GCTGATTCAGGGTAGAGGCCTGTTCGGAGCTATTGCAGGGTTTTTGGAAAATGGGTGGGAGGGCATGGTGGACGGATGGTACGGGTTCCG ACACCAGAATGCCCAGGGAACCGGCCAGGCGGCTGACTACAAGTCTACACAGGCAGCAATTGACCAAATCACCGGAAAGTTGAACCGCCT TGTAGAGAAAACAAACACGGAATTTGAATCAATTGAATCAGAGTTTTCCGAGATAGAGCACCAGATCGGGAATGTCATAAATTGGACAAAGG ATTCTATTACGGATATTTGGACCTACCAAGCAGAACTGCTTGTGGCTATGGAAAATCAGCACACCATAGATATGGCCGACAGTGAGATGCTG AACCTGTACGAGCGGGTAAGAAAGCAGTTGCGCCAGAACGCAGAGGAAGACGGGAAAGGCTGCTTCGAAATCTATCACGCCTGTGACGAC TCATGCATGGAAAGTATCAGGAATAATACATACGATCATTCCCAATACCGTGACGAAGCACTACTCAACCGGCTGAATATCAACCCCGTTAC ACTCTCCTCAGGTTATAAAGACATAATCCTGTGGTTCTCCTTCGGCGCTAGCTGCTTTGTCCTGTTGGCCGTAGTAATGGGGCTGTTCTTCTT TTGCCTTAAAAACGGGAACATGCGATGTACTATCTGCATT
3028	ATGTACAAGGTCGTAGTGATTATCGCCCTCCTTGGCGCCGTTAAAGGCCTAGACAAGATCTGTCTGGGACATCACGCCGTCGCTAATGGCA CAATAGTGAAAACGCTCACTAATGAGCAGGAGGAAGTAACCAACGCCACAGAAACGGTCGAATCCACCGGCATAAATCGGCTTTGCATGAA GGGGCGTAAACATAAGGACCTGGGGAACTGTCACCCAATAGGGATGCTGATCGGAACACCTGCTTGTGACCTCCACCTGACAGGCACCTG GGATACGTTGATCGAGCGGGAGAACGCTATCGCATACTGCTACCCCGGCGCTACCGTCAATGTGGAAGCACTGCGTCAGAAGATAATGGA ATCAGGAGGCATTGATAAGATCTCAACAGGGTTTACCTACGGCTCTAGTATCAATAGCGCAGGTACCACCAGGGCATGCATGCGCAATGGT GGCAACAGTTTCTACGCAGAGCTGAAGTGGCTGGTGTCTAAAAATAAAGGCCAGAACTTCCCTCAGACCACAAACACTTACCGCAATACAGA TACTGCAGAGCATCTGATTATGTGGGGGATCCACCACCCTTCCTCCATTCAAGAGAAGAACGACCTGTACGGCACTCAGAGCTTATCAATAT CGGTGGGTTCCTCCACTTACCGTAACAATTTCGTCCCCGTAGTCGGGGCTAGGCCTCAGGTGAATGGTCAGTCAGGGAGGATAGACTTTCA CTGGACGCTGGTTCAGCCAGGCGACAATATAACCTTCTCACACAACGGGGGTCTGATTGCTCCTTCTAGGGTGTCAAAGCTGATCGGGAGA GGACTGGGTATACAAAGCGATGCCCCCATCGATAATAACTGTGAAAGTAAGTGTTTTTGGCGCGGAGGGAGCATTAACACGCGGCTGCCTT TTCAGAACCTGTCCCCCCGGACTGTGGGCCAGTGTCCCAAGTACGTGAACCGCAGGTCACTCATGCTAGCCACGGGTATGAGAAATGTGC CAGAGTTAATTCAGGGCCGCGGTCTGTTTGGGGCTATTGCAGGTTTTCTGGAGAACGGGTGGGAGGGCATGGTGGACGGTTGGTACGGCT TTAGGCATCAGAACGCACAGGGTACTGGACAGGCAGCCGATTACAAGAGCACTCAGGCAGCGATAGACCAGATAACTGGCAAGCTCAACC GTTTAGTGGAGAAGACCAACACCGAATTTGAAAGTATTGAGTCCGAGTTCTCAGAGATTGAGCATCAGATCGGCAATGTCATTAATTGGACA ATGGATTCGATCACCGATATATGGACCTATCAGGCCGAGTTGCTGGTCGCCATGGAAAATCAACACACTATAGATATGGCAGATTCCGAAAT GCTGAATCTCTATGAGAGGGTGAGAAAACAACTGAGGCAAAATGCGGAGGAAGACGGGAAAGGTTGCTTCGAAATTTACCATGCTTGCGAT GATAGTTGCATGGAGTCCATCCGGAATAACACGTATGATCACAGCCAATACCGGGAAGAAGCCCTGCTTAACCGGCTTAACATAAATCCTGT GACTCTCTCATCAGGATATAAGGATATCATCCTCTGGTTCTCTTTCGGGGCCAGCTGCTTTGTTTTGTTAGCCGTGGTGATGGGACTCGTGT TCTTCTGCTTAAAAAACGGCAATATGCGTTGCACCATCTGCATC

3029	ATGTATAAGATCGTCGTTATTATTGCCCTGCTTGGTGCTGTCAAGGGGTTGGACAAAATTTGCTTGGGCCACCACGCAGTGGCTAACGGCA CTATAGTCAAAACCCTGACGAACGAGCAGGAGGAGGTGACAAACGCCACTGAGACCGTGGAATCTACGGGAATCAACAGGCTCTGCATGA AGGGGCGCAAACACAAAGACCTCGGGAACTGCCACCCTATCGGGATGCTCATAGGTACTCCAGCGTGTGACCTGCACTTAACAGGCACTT GGGATACCCTTATTGAACGGGAAAATGCTATCGCTTACTGCTACCCGGGCGCGACCGCCAACGTGGAAGCATTACGCCAAAAGATAATGGA ATCAGGCGGCATAGATAAAATCTCCACCGGATTTACATACGGGTCTACCATAAATTCTGCCGGCACCACTAGAGCCTGTATCAGAAACGGC GGAAATTCATTCTACGCTGAGCTGAAATGGCTGGTCTCAAAGTCTAAGGGGCAGAATTTCCCCCAGACCACGAATACGTACCGGAACACAG ATACCGCCGAGCATCTGATCATGTGGGGAATCCATCACCCCAGTTCAATCCAGGAGAAGAATGACTTATACGGTACACAATCCCTGTCCATA TCAGTAGGTAGCTCTACATATCGGAACAATTTCGTGCCTGTCGTCGGGGCTAGGCCTCAGGTAAACGGCCAGTCCGGTAGAATCGATTTTC ACTGGACCCTCGTCCAGCCTGGGGACAATATAACGTTTAGTCACAACGGAGGTCTCATCGCCCCCAGCCGGGTATCTAAGCTTATTGGCAG GGGCCTGGGTATTCAGAGCGACGCCCCAATTGACAATAATTGCGAAAGCAAGTGCTTCTGGAGGGGTGGAAGTATTAACACAAGACTGCCT TTCCAAAACCTGAGCCCTAGAACTGTGGGACAGTGCCCAAAATACGTCAACAGGAGATCACTTATGCTGGCGACAGGAATGCGCAATGTAC CTGAACTTATTCAAGGGCGAGGACTGTTTGGTGCCATAGCTGGGTTCCTGGAAAACGGTTGGGAGGGAATGGTGGACGGCTGGTACGGGT TTAGACACCAAAACGCACAAGGAACCGGCCAGGCTGCCGATTATAAGTCCACACAAGCAGCCATCGATCAAATTACCGGTAAGCTGAATCG GTTAGTTGAAAAAACCAATACCGAGTTTGAATCCATCGAGAGTGAGTTCTCAGAGATCGAGCACCAGATAGGCAATGTGATCAACTGGACAA TGGATTCAATTACCGATATCTGGACCTACCAGGCTGAGCTGCTGGTGGCCATGGAAAATCAGCATACAATTGATATGGCCGACAGTGAGAT GCTGAACTTGTATGAAAGAGTTAGAAAACAGCTGAGGCAAAATGCCGAAGAGGATGGCAAGGGATGTTTTGAAATCTATCATGCGTGTGAT GATAGCTGTATGGAATCCATCAGGAACAATACATACGATCACTCACAGTATAGAGAAGAAGCACTCCTGAACAGATTAAATATCAATCCAGTG ACCTTATCATCTGGCTACAAGGATATAATTCTTTGGTTTTCGTTTGGGCCTTCCTGCTTCGTGTTGCTGGCCGTGGTCATGGGTCTAGTGTTT TTTTGCTTGAAAAACGGAAATATGCGGTGTACCATCTGTATC
3030	ATGTACAAGATTGTGGTTATAATTGCGCTTCTGGGAGCTGTGAAGGGATTGGACAAGATCTGCCTTGGGCACCATGCCGTTGCAAATGGTA CCATTGTGAAGACGCTGACCAATGAGCAGGAAGAGGTGACCAATGCTACAGAGACAGTGGAGTCCACTGGCATCAACCGACTCTGTATGAA AGGTAGAAAACACAAAGATCTCGGCAATTGCCATCCTATAGGTATGCTGATTGGTACGCCTGCCTGTGACCTCCATCTGACTGGCATGTGG GACACGCTGATCGAAAGGGAGAATGCCATTGCGTACTGTTATCCTGGCGCTACCGTGAACGTGGAAGCCTTACGGCAGAAGATCATGGAAT CCGGCGGGATCAATAAGATTAGCACCGGATTCACTTACGGATCTTCTATCAACAGCGCTGGCACCACTAGGGCATGCATGAGAAACGGCGG GAATAGTTTTTACGCTGAGCTTAAGTGGTTGGTTTCTAAGAGTAAGGGACAGAACTTCCCTCAGACCACTAATACCTACAGGAACACCGACA CCGCAGAGCATTTAATCATGTGGGGAATCCACCATCCATCCTCCACTCAGGAAAAGAATGACTTATACGGAACACAGAGCCTTTCCATATCT GTCGGTTCCTCCACATATAGAAATAATTTTGTGCCCGTGGTCGGTGCACGACCTCAGGTGAATGGCCAGTCAGGAAGAATCGATTTTCACTG GACCCTGGTACAGCCTGGCGACAATATCACCTTCAGCCATAACGGCGGTCTGATCGCCCCTAGTCGGGTGAGTAAATTGATAGGTCGGGG ACTGGGAATCCAGAGTGATGCTCCTATTGATAATAATTGCGAGTCAAAATGTTTTTGGCGCGGCGGCTCTATTAACACCAGATTACCTTTCCA GAACCTGAGCCCTCGGACTGTTGGACAGTGTCCTAAATATGTGAATCGCAGGTCTCTGATGCTGGCTACAGGCATGAGGAACGTGCCTGAG CTCATCCAGGGAAGGGGGCTGTTCGGGGCCATCGCTGGATTCCTGGAAAACGGATGGGAGGGAATGGTCGATGGTTGGTACGGTTTTCGG CATCAGAATGCTCAAGGGACGGGCCAGGCGGCTGATTATAAGTCAACGCAGGCGGCCATAGACCAGATCACCGGAAAGCTCAACCGCCTG GTGGAGAAGACAAATACAGAGTTTGAGTCTATAGAGTCCGAGTTTTCCGAAATCGAACACCAGATCGGGAATGTGATAAACTGGACCAAAGA TTCTATCACTGACATTTGGACTTACCAGGCAGAACTATTGGTGGCCATGGAAAATCAGCACACCATCGACATGGCCGACTCTGAAATGCTGA ACTTGTATGAGAGAGTGCGCAAGCAGCTGCGACAAAATGCCGAAGAAGACGGAAAGGGCTGTTTTGAAATCTATCATGCTTGTGATGACTC TTGCATGGAATCTATTAGGAACAATACCTATGATCATTCTCAGTACCGGGAACAAGCATTGCTGAACCGCTTGAATATAAACCCACTGACGC TGTCATCTGGTTATAAAGACATTATCCTGTGGTTTAGCTTTGGTGCCTCATGCTTTGTACTCCTAGCCGTAGTCATGGGCCTGTTTTTCTTCT GCCTGAAGAACGGGAACATGCGATGCACTATTTGCATC
3031	ATGTACAAGATCGTGGTGATCATAGCCTTACTGGGGGCCGTAAAAGGTCTGGATAAGATTTGTCTGGGCCACCACGCTGTCGCCAATGGTA CCATTGTGAAGACACTAACCAACGAGCAGGAGGAGGTCACAAACGCTACCGAGACGGTTGAATCTACTGGTATAAACCGCCTTTGCATGAA GGGACGCAAACATAAGGATCTGGGGAACTGTCACCCCATTGGTATGCTGATCGGAACCCCTGCATGTGACCTACACCTGACCGCAATGTG GGATACCTTAATAGAGCGAGAGAACGCGATCGCCTACTGTTATCCGGGGGCTACGGTGAACGTTGAGGCTCTGAGGCAGAAGATAATGGA AAGTGGGGGCATCAACAAGATTAGTACCGGATTCACTTATGGTAGTAGCATTAATAGTGCTGGGACTACCAGAGCATGTATGAGAAATGGC GGGAACAGCTTCTATGCCGAGCTGAAGTGGCTAGTGAGCAAGAGCAAGGGCCAAAACTTTCCCCAAACGACCAACACATACCGCAATACCG ATACAGCAGAGCATCTCATAATGTGGGGAATCCACCACCCTAGTTCAACCCAGGAGAAGAACGATCTTTATGGAACTCAGTCCCTGTCTATC AGCGTGGGGAGTTCTACATATCGCAATAACTTTGTGCCAGTCGTGGGGGCCCGACCACAGGTAAACGGTCAGTCTGGTCGAATTGACTTTC ATTGGACCTTGGTCCAGCCGGGCGATAATATCACATTTTCCCATAATGGAGGCCTGATAGCCCCTTCCAGAGTTAGCAAATTGATAGGCCGA GGACTCGGGATCCAGAGCGATGCACCCATCGACAACAACTGCGAGAGCAAATGCTTCTGGCGGGGCGGAAGCATTAACACTAGATTACCC TTCCAGAACCTTAGTCCTCGCACCGTGGGCCAGTGCCCTAAATATGTGAACCGGCGCTCACTGATGCTGGCCACAGGCATGAGGAATGTCC CCGAGCTGATCCAGGGCAGAGGACTCTTCGGTGCTATCGCAGGATTTCTTGAGAACGGATGGGAAGGAATGGTAGACGGATGGTACGGGT TCCGACATCAGAACGCTCAGGGCACTGGCCAAGCAGCCGACTACAAATCCACACAGGCGGCCATCGATCAGATAACAGGCAAACTGAATC GGCTAGTAGAGAAGACCAACACCGAATTCGAATCAATCGAGTCTGAATTCAGCGAGATAGAGCACCAAATCGGGAATGTGATTAACTGGAC AAAGGATTCCATAACCGACATTTGGACCTACCAAGCCGAACTGCTGGTCGCGATGGAAAACCAACATACCATCGACATGGCAGACTCCGAA ATGCTGAATCTGTACGAACGTGTGCGGAAACAGCTGAGACACAATGCTGAAGAAGACGGGAAAGGCTGTTTTGAAATCTACCACGCCTGCG ACGACTCATGCATGGAAAGCATCAGGAATAACACTTATGACCATTCCCAGTACCGGGAAGAAGCACTCCTGAACCGGTTGAACATTAATCCG GTTACCTTGAGCTCTGGATATAAAGACATCATCCTATGGTTCAGCTTCGGCGCATCCTGCTTTGTTCTGCTTGCAGTGGTCATGGGGCTGTT CTTTTTTTGCCTGAAGAATGGCAACATGAGATGTACCATTTGCATC

3032	ATGTATAAGATAGTAGTCATCATAGCTCTCTTGGGCGCCGTGAAAGGGCTGGACAAGATCTGTCTGGGTCATCATGCCGTGGCCAACGGAA CAATCGTGAAAACCTTGACTAATGAGCAGGAGGAGGTGACCAATGCGACCGAGACCGTTGAAAGTACGGGCATTAATAGGCTCTGTATGAA GGGTAGGAAACATAAGGACCTTGGGAACTGTCACCCCATTGGAATGCTGATTGGGACTCCTGCATGTGACTTACACTTAACAGGCATGTGG GACACTCTGATTGAGAGGGAGAACGCCATTGCCTATTGTTATCCCGGCGCCACCGTGAACGTGGAGGCACTGCGGCAGAAGATAATGGAA AGCGGAGGGATTAATAAAATCAGCACCGGATTCACTTATGGTAGTTCTATTAATTCCGCCGGGACAACAAGGGCATGTATGACGAACGGAG GTAATTCCTTCTACGCGGAGCTGAAGTGGCTGGTCTCTAAGTCCAAGGGGCAGAACTTTCCCCAGACTACAAACACCTACCGAAATACAGAT ACCGCTGAGCACCTGATCATGTGGGGAATTCATCACCCTTCATCAACACAAGAGAAGAACGACCTGTACGGCACACAGTCACTGTCAATCT CAGTAGGGAGCTCGACATATAGAAATAATTTCGTCCCGGTGGTGGGCGCTCGGCCGCAGGTTAATGGCCAGAGTGGTAGAATCGACTTTCA CTGGACGCTGGTCCAGCCAGGCGATAATATCACGTTCTCACACAATGGAGGCCTGATAGCCCCAAGCCGCGTGTCCAAGCTGATAGGGCG AGGACTTGGTATCCAGAGTGATGCCCCCATCGATAACAACTGCGAATCCAAGTGTTTCTGGCGTGGCGGCTCAATTAACACCCGGCTTCCA TTCCAGAACCTTAGCCCCCGCACTGTCGGACAATGCCCAAAGTATGTCAACAGGCGGAGCCTGATGCTGGCCACCGGCATGAGAAATGTG CCTGAGCTCATTCAGGGGCGGGGCCTTTTTGGGGCAATCGCTGGCTTTCTCGAGAATGGTTGGGAGGGAATGGTTGACGGCTGGTATGGC TTTAGACATCAGAATGCCCAGGGAACGGGCCAGGCTGCTGACTACAAGTCCACCCAGGCCGCCATCGACCAAATTACCGGGAAGCTGAAT AGACTCGTCGAAAAGACCAACACTGAATTTGAGAGTATCGAGTCTGAGTTTTCCGAGATCGAACATCAAATCGGAAACGTTATCAACTGGAC TAAGGACTCAATCACCGACATTTGGACATATCAAGCGGAGCTCCTGGTGGCCATGGAAAATCAGCACACGATCGACATGGCCGACAGTGAG ATGCTGAATCTGTATGAGCGAGTCCGTAAGCAGCTTAGGCAGAACGCTGAGGAGGATGGAAAAGGATGTTTCGAGATATATCACGCCTGTG ATGACTCCTGTATGGAATCCATCAGGAACAACACATACGATCATAGTCAGTACAGGGAAGAAGCCTTGCTAAATAGGTTGAACATCAACCCT GTCACTCTGAGTTCTGGATACAAGGACATCATATTATGGTTTTCCTTTGGCGCCTCCTGCTTCGTCCTGCTAGCAGTGGTGATGGGCCTGTT TTTTTTTTGCCTTAAAAATGGCAATATGCGGTGCACCATTTGCATA
3033	ATGTATAAGATCGTTGTCATCATTGCTCTCCTCGGAGCCGTGAAAGGGCTCGATAAGATCTGTTTAGGACACCATGCCGTGGCAAACGGTAC TATAGTCAAGACCCTAACAAATGAGAAAGAGGAGGTGACGAATGCCACAGAAACAGTAGAATCTACAGGACTGAACCGGCTGTGCATGAAG GGCCGCAAACACAAGGATTTAGGTAATTGTCACCCAATAGGCATGTTGATTGGAAGTCCGGCATGTGACCTGCACTTGACGGGCACATGGG ATACCCTGATAGAAAGGGAGAACGCTATCGCCTACTGCTATCCGGGAGCCACGGTCAACGGGGAAGCGCTGCGGCAGAAGATTATGGAGT CGGGCGGTATAGATAAGATATCCACAGGCTTTACCTATGAATCCTCTATTAACTCAGCTGGTACAACTCGCGCTTGCATGAGGAACGGGGG GAACAGTTTCTACGCCGAACTCAAGTGGCTTGTTTCTAAGTCTAAGGGCCAGAATTTCCCACAGACTACCAATACCTATAGAAACACCGACA CGGCTGAGCACCTTATTATGTGGGGTATTCACCACCCCTCAAGCACTCAGGAGAAGAACGACCTCTATGGTACCCAGTCCCTTTCCATCTCC GTTGGTAGTTCTACCTACCGCAACAATTTTGTTCCCGTGGTCGGTGCAAGACCACAGGTTAACGGCCAGAGTGGGCGCATCGATTTCCACT GGACGTTGGTCCAGCCTGGGGATAACATCACCTTTTCCCACAACGGGGGATTAATCGCACCCTCTCGGGTCTCTAAGCTGATCGGACGAG GACTGGGAATCCAAAGTGACGCACCAATAGATAATAATTGCGAGAGCAAATGCTTCTGGCGCGGTGGGTCAATCAATACACGGCTGCCGTT TCAGAATCTATCACCAAGGACTGTGGGACAGTGCCCCAAATATGTAAATAAGCGGTCCCTGATGCTCGCCACTGGTATGCGGAACGTGCCT GAGCTAATGCAAGGAAGGGGCCTGTTTGGGGCCATTGCCGGCTTTCTCGAAAACGGGTGGGAGGGCATGGTGGACGGATGGTACGGTTT CCGGCACCAGAATGCTCAAGGCACCGGTCAGGCTGCCGATTATAAAAGCACCCAAGCCGCTATCGATCAGATCACTGGCAAGTTAAACAGG CTGGTAGAAAAAACTAATACTGAATTCGAGTCCATTGAGTCCGAGTTCTCTGAGATTGAGCATCAGATCGGGAATGTTATTAACTGGACGAA AGACAGTATTACGGATATCTGGACATATCAAGCCGAGCTCCTGGTCGCTATGGAAAACCAACACACTATCGACATGGCCGACAGCGAAATG TTGAATCTTTATGAAAGAGTGAGGAAGCAGCTGCGTCAGAACGCAGAGGAGGATGGGAAAGGATGTTTTGAGATCTACCACGCCTGTGACG ATTCCTGCATGGACAGTATTAGAAACAACACATACGACCACAGCCAGTACCGGGAAGAGGCCCTATTGAACAGGCTTAATATTAATCCAGTA ACCCTGTCGAGTGGGTATAAAGATATTATACTGTGGTTCTCCTTTGGCGCTTCTTGCTTTGTCCTGCTCGCGGTAGTTATGGGTTTGGTGTTT TTTTGCCTCAAGAACGGAAATATGCGTTGCACCATCTGTATC
3034	ATGTATAAGATCGTGGTCATCATCGCCCTGCTTGGAGCTGTAAAAGGTCTCGACAAGATTTGTCTTGGACATCATGCTGTGGCAAATGGCAC CATTGTAAAGACCCTGACTAATGAGCAAGAAGAAGTAACCAACGCAACTGAGACCGTAGAGTCAACCGGCATTAATCGGCTCTGCATGAAG GGAAGCAAGCACAAAGATCTCGGCAATTGTCACCCAATAGGCATGTTGATTGGAACCCCTGCTTGTGACCTGCATCTCACGGGGATGTGGG ATACTTTGATTGAAAGGGAAAACGCAATTGCGTATTGTTACCCGGGAGCCACCGTGAACGTCGAGGCCCTCCGACAAAAGATCATGGAATC AGGAGGGATCAACAAAATAAGCACTGGGTTTACCTATGGAAGTTCGATCAACTCAGCCGGGACAACGCGCGCTTGCATGAGGAATGGCGG AAATTCCTTCTACGCTGAACTCAAATGGCTCGTGTCGAAATCAAAGGGTCAGAACTTCCCACAGACTACTAATACCTACAGAAATACTGACAC AGCGGAGCATTTGATCATGTGGGGCATACACCACCCTTCTTCCACACAGGAGAAGAATGACCTTTACGGCACGCAGAGTCTGTCCATCAGC GTAGGAAGCTCAACCTACAGAAATAATTTCGTTCCCGTGGTGGGGGCTCGACCACAGGTTAACGGGCAGTCCGGTCGTATTGATTTTCATT GGACATTGGTGCAGCCCGGCGATAACATTACTTTTAGCCACAACGGAGGCCTAATTGCCCCTAGCAGAGTCTCCAAACTGATCGGGAGGG GACTGGGGATACAGTCAGATGCCCCTATCGACAACAATTGTGAGAGCAAGTGTTTCTGGCGAGGCGGAAGCATTAACACCCGGTTACCCTT TCAGAACCTATCTCCAAGAACTGTGGGTCAGTGCCCAAAATACGTGAACAGACGGTCTCTGATGTTGGCTACTGGTATGCGCAACGTTCCG GAACTAATTCAAGGCAGAGGACTGTTCGGCGCCATCGCAGGGTTCCTCGAGAACGGGTGGGAGGGGATGGTCGATGGGTGGTACGGCTT TAGGCATCAGAACGCTCAGGGGACTGGTCAGGCCGCCGATTACAAAAGCACGCAAGCTGCTATAGATCAGATCACAGGGAAACTGAATCG CCTCGTTGAGAAAACTAACACCGAATTTGAATCAATAGAATCCGAATTCAGTGAAATCGAGCACCAGATAGGGAACGTAATCAACTGGACCA AAGACAGTATTACAGACATCTGGACATACCAGGCGGAACTTTTAGTCGCTATGGAGAATCAGCACACAATAGACATGGCCGACAGCGAAAT GTTGAACCTGTACGAACGAGTAAGGAAGCAGCTGAGGCAAAACGCAGAAGAGGACGGAAAGGGCTGCTTTGAGATATACCACGCATGTGA TGATAGCTGTATGGAGTCAATTCGGAACAATACATATGATCACTCTCAATACCGTGAGGAAGCTCTGCTCAATCGTCTGAATATTAACCCAGT GACTCTTTCCAGTGGGTACAAAGACATCATTCTTTGGTTTTCCTTCGGAGCTTCCTGCTTTGTGCTGCTGGCAGTCGTTATGGGATTGTTCTT TTTCTGCCTGAAGAACGGGAACATGCGCTGTACTATATGTATC

3035	ATGTACAAGGTAGTGGTAATAATTGCTCTGCTGGGAGCCGTGAGAGGTCTGGATAAAATCTGTTTGGGTCACCACGCCGTCGCCAACGGGA CGATCGTCAAAACACTAACAAATGAGCAAGAAGAAGTGACCAACGCAACAGAGACAGTGGAGAGTAAGAGCCTGGGAAAGCTTTGCATGAA GGGCCGCTCATATAACGACCTGGGCAACTGTCACCCCATCGGTATTCTGATTGGGACTCCAGCTTGCGACTTGCATCTGACCGGCACCTGG GACACACTCATCGAGAGAGAAAACGCAGTGGCTTACTGCTACCCTGGCGCTACCGTCAATGAAGAAGCCCTGCGACAGAAGATTATGGAGA GCGGTGGGATCAGCAAAATCTCCACCGGTTTCACCTATGGAACCTCCATTAACAGCGCAGGCACAACAAAAGCTTGCATGAGGAATGGCGG CAACTCGTTCTATGCTGAACTGAAATGGCTGGTGTCCAAAAATAAGGGTCAGAATTTCCCTCAGACTACCAACACTTACAGGAACACGGATA CAGCAGAGCACCTGATTATATGGGGGATTCACCACCCTTCGTCTACCCAAGAGAAAAATGACCTGTACGGAACTCAGAGCCTTTCAATATCC GTGGGGTCCAGTACATATCAAAATAATTTTGTCCCCGTGGTTGGCGCCCGTCCACAGGTTAACGGTCAGTCAGGGCGGATCGATTTTCATT GGACCTTACTTCAGCCCGGAGACAATATCACTTTTTCTCACAACGGAGGACTCATCGCCCCCTCCCGTGTGTCCAAGCTCATAGGACGGGG CCTGGGTATCCAGAGCGAGGCCCCCATCGATAACGGCTGCGAGAGCAAATGTTTTTGGAAGGGAGGGAGTATAAACACTAAGCTCCCATTC CAAAATTTAAGTCCACGTACTGTGGGCCAGTGTCCCAAGTATGTGAACAAAAGAAGCTTGATGCTGGCAACTGGGATGCGGAACGTGCCTG AGATCATGCACGGTCGGGGCCTGTTTGGCGCTATCGCAGGGTTTATTGAAAATGGCTGGGAGGGCATGGTTGATGGGTGGTACGGGTTTA GGCATCAGAATGCCCAGGGCACTGGTCAGGCGGCCGATTACAAGAGTACTCAGGCTGCCATCGACCAGATCACCGGCAAGCTGAACCGGT TGATCGAAAAGACAAACACTGAATTCGAGAGTATCGAAAGCGAATTTTCTGAGATCGAGCACCAAATCGGGAATATAATTAATTGGACAAAG GACTCGATCACTGACATATGGACATACCAAGCCGAGCTGCTGGTGGCCATGGAAAACCAGCATACCATAGACATGGCCGATTCAGAAATGC TTAATCTTTACGAGCGTGTTAGGAAGCAACTGAGGCAGAATGCCGAAGAGGACGGTAAGGGGTGCTTTGAGATTTATCACGCATGTGACGA CAGCTGTATGGAGTCCATTAGAAATAATACCTACGATCACAGTCAGTATAGAGAAGAGGCATTGCTTAATAGGCTGAACATAAATCCCGTCA AGCTGTCTTCGGGGTACAAGGACATTATCCTGTGGTTCAGCTTTGGCGCATCGTGTCTGATATTACTGGCCGTCGTGATGGGATTAGTGTTT TTCTGCTTAAAGAACGGTAACATGCGCTGTACTATCTGTATC
3036	ATGTACAAGATCGTGGTCATTATCGCTCTGTTAGGGGCAGTAAAAGGCCTGGACAAAATCTGCCTTGGCCATCATGCTGTGGCTAATGGTAC AATTGTCAAAACCCTCACAAATGAACAGGAGGAGGTGACAAACGCCACAGAAACTGTGGAGAGTACAGGGATCAACCGTCTCTGCATGAAG GGAAGAAAGCATAAAGACCTTGGAAACTGTCATCCCATAGGCATGCTGATCGGCACTCCCGCCTGTGACCTGCACCTGACCGGTATGTGGG ACACCCTGATCGAAAGGCAGAATGCCATTGCCTACTGTTATCCTGGCGCAACAGTGAACGTAGAAGCCTTAAGACACAAGATCATGGAATC CGGAGGCATAAACAAAATTAGTACTGGATTCACCTACGGGAGCTCCATTAACTCTGCAGGCACCACTCGGGCATGCATGCGCAACGGCGGT AACAGCTTCTATGCCGAGCTGAAGTGGTTAGTGTCCAAGTCGAAAGGCCAGAACTTTCCCCAGACGACAAACACATATCGGAACACCGATA CTGCAGAACACCTGATCATGTGGGGGATTCACCACCCCAGTTCCACCCAGGAGAAGAACGATCTGTACGGTACTCAGAGCCTGTCCATAAG TGTTGGATCCTCTACATATCGGAATAATTTCGTGCCCGTGGTGGGCGCAAGACCACAGGTGAACGGCCAGTCTGGGAGAATTGACTTTCAT TGGACTCTGGTGCAACCAGGGGACAATATTACCTTCTCACATAATGGCGGCCTGATCGCACCTAGCAGGGTGTCCAAGCTGATAGGACGCG GACTCGGAATTCAGTCGGATGCACCTATCGATAATAACTGCGAGTCTAAGTGTTTTTGGAGAGGCGGTTCAATCAATACTAGGCTTCCTTTT CAGAACCTGAGCCCCAGAACAGTGGGGCAATGCCCTAAATATGTGAATAGGCGGTCTCTGATGCTGGCCACTGGGATGAGAAATGTGCCA GAGCTGATTCAGGGTAGAGGCCTGTTCGGCGCAATCGCAGGGTTCCTCGAAAACGGCTGGGAAGGTATGGTGGACGGATGGTACGGGTT CCGACACCAAAACGCCCAGGGGACGGGCCAGGCCGCCGATTACAAATCAACTCAAGCCGCGATAGATCAGATAACCGGAAAGCTGAACAG ACTCGTCGAGAAGACCAATACCGAGTTTGAGTCCATCGAAAGCGAGTTCTCAGAGATTGAGCATCAGATAGGGAACGTAATTAACTGGACG AAAGATAGCATCACGGACATCTGGACATATCAGGCGGAACTACTGGTGGCAATGGAAAATCAGCATACCATCGACATGGCCGACAGTGAGA TGTTAAATCTCTATGAGCGGGTGCGGAAACAGCTGAGGCAAAACGCTGAAGAAGACGGAAAAGGGTGCTTTGAGATCTACCATGCCTGCGA CGATAGCTGCATGGAGTCAATTCGGAACAATACTTATGACCACTCCCAGTATCGGCAGGAGGCCCTGCTCAATCGACTGAACATAAACCCA GTGACTCTGAGTTCCGGATACAAAGATATTATTCTTTGGTTCAGCTTCGGAGCTTCTTGTTTTGTCCTCCTAGCAGTGGTAATGGGCCTCTTT TTCTTTTGCCTGAAGAACGGAAACATGAGGTGCACAATTTGTATT
3037	ATGTACAAAATAGTGGTGATAATCGCCCTTCTCGGCGCAGTGAAAGGGCTCGACAAGATATGTCTGGGACATCACGCTGTGGCTAATGGCA CTATCGTGAAGACGCTCACAAATGAACAGGAGGAGGTTACAAATGCCACTGAGACAGTTGAAAGCACAGGAATCAATAGATTGTGTATGAAA GGCAGAAAGCATAAGGACTTGGGGAACTGTCACCCTATTGGCATGCTTATCGGCACCCCTGCTTGTGATCTACATTTGACAGGCATGTGGG ACACCCTTATTGAGCGCGAAAACGCAATCGCGTACTGTTATCCTGGTGCTACTGTGAACGTAGAAGCTCTGAGGCAGAAGATAATGGAATCT GGCGGGATAAATAAAATCTCAACCGGCTTCACATACGGGAGTAGCATTAATAGCGCCGGTACTACCCGGGCCTGTATGAGGAATGGAGGAA ACTCTTTTTACGCAGAGCTCAAGTGGCTAGTATCTAAGTCCAAGGGACAGAATTTTCCCCAGACTACTAATACTTACCGCAATACGGATACC GCAGAACACTTAATTATGTGGGGCATACATCACCCATCCTCTACCCAAGAAAAGAACGACCTTTACGGCACACAGTCCTTGAGTATATCTGT GGGGAGTAGCACCTATAGGAACAATTTCGTTCCGGTCGTGGGCGCTCGCCCCCAGGTGAACGGCCAATCAGGACGGATCGACTTCCACTG GACCCTGGTCCAGCCCGGCGATAACATCACATTTTCCCATAACGGAGGACTTATAGCACCCTCCCGGGTGTCTAAACTTATTGGACGGGGG CTGGGAATCCAGTCCGACGCCCCCATCGATAACAACTGTGAGTCTAAATGTTTTTGGCGCGGCGGATCTATCAACACCAGGCTTCCCTTCC AGAATCTGTCCCCCCGGACAGTGGGCCAGTGCCCAAAGTACGTTAACAGAAGGTCTTTAATGCTCGCCACCGGCATGCGAAACGTGCCTG AACTGATCCAGGGAAGAGGACTTTTCGGAGCCATCGCCGGCTTTTTAGAGAACGGATGGGAAGGGATGGTAGACGGGTGGTATGGATTCA GGCACCAAAACGCTCAGGGGACCGGGCAAGCAGCAGACTACAAAAGTACACAAGCCGCTATCGACCAAATCACGGGTAAACTGAACCGGC TTGTGGAAAAGACAAACACCGAATTTGAATCGATAGAGAGCGAATTCAGCGAGATCGAGCACCAAATAGGAAATGTGATCAACTGGACAAA GGATTCGATCACAGACATCTGGACGTACCAAGCCGAGCTCCTCGTAGCAATGGAGAACCAGCATACCATCGACATGGCCGATTCCGAGATG CTTAATCTTTATGAACGGGTCCGCAAACAGCTGCGCCAAAATGCCGAAGAAGATGGTAAAGGATGTTTTGAAATCTATCATGCTTGTGATGA TTCCTGTATGGAGTCAATCCGGAATAATACATACGACCACAGCCAGTACCGGGAAGAAGCCCTGCTGAATCGCTTAAATATCAACCCCGTTA CCCTTTCCTCTGGATATAAGGATATTATCCTGTGGTTTTCATTTGGGGCATCCTGTTTCGTCCTTCTGGCAGTGGTCATGGGACTTTTTTTTTT CTGTCTGAAGAACGGAAACATGCGGTGTACGATCTGTATA

3038	ATGTATAAAGTCGTGGTTATCATCGCCCTTCTGGGAGCCGTCAGAGGCCTGGACAAGATATGTTTGGGGCATCATGCGGTAGCTAATGGAA CCACTGTCAAAACACTTACTAATGAACAGGAAGAGGTTACGAACGCCACCGAAACAGTGGAATCTACCAGCCTGAATAAATTATGCATGAAG GGACGACGCTACAAGGATCTGGGGAATTGCCACCCCATCGGAATGCTGATTGGAACACCAGTCTGTGACCTCCACCTTACAGGGACGTGG GATACCTTGATTGAGCGGGAGAATGCAACAGCTTACTGTTATCCCGGAGTAACTATTAATGAGGAGGCTCTCCGTCAAAAGATAATGGAGTC AGGAGCTATTAGTAAAATGCGCACTGCCTTTACATACGGGCCCTCTATAAACAGCGCAGGAACCACTCGTTCATGTATGCGCAATGGGGCC AACTCGTTCTACGCCGAGCTGAAGTGGCTGGTGAGTGGCACCAAAGGTCAGAATTTTCCCCAGACTACAAACACCTACCGGAACACAGACA CTGCCGAGCATCTGATCATCTGGGGGATACATCACCCTTCCTCTACTCAAGAAAAGAACGATCTGTATGGCACACAAAGCCTGTCTATATCC GTGGGTTCTTCTACCTATCAAAACAACTTTGTTCCAGTTATAGGCGCCCGGCCCCAGGTGAACGGGCAGTCAGGCCGCATCGAGTTTCACT GGACTCTGGTCCGGCCCGGAGATAATATTACATTCAGTCATAACGGCGGTTTGATCGCCCCAGACCGAGTGAGCAAGCTCATCGGGAAGG GGATTGGCATACAATCCGGTGCTGTGATTGACAAGGATTGCGAGAGTAAGTGCTTTTGGCGAGGCGGATCTATTATTACCGAGCTGCCATTT CAGAACCTTTCTCCTCGAACCGTTGGGCAGTGCCCCAAATATGTGAAAAAGAGGTCACTGCTCCTCGCCACTGGCATGCGAAATGTTCCTG AGGTGGTGCAGGGCCGTGGGTTGTTCGGTGCCATCGCAGGGTTCATCGAGAATGGATGGGAAGGTATGGTCGACGGATGGTACGGATTC CGCCATCAGAATGCACAGGGTATCGGACAGGCCGCCGATTATAAGAGCACCCAGACGGCTATAGACCAAATCACCGGCAAGCTTAACCGG CTCATAGAAAAGACCAACACAGAATTTGAGTCCATCGAGTCAGAATTCAGCGAAATTGAACATCAGATTGGCAACGTCATCAATTGGACCAA GGACTCCATCACTGATATCTGGACATACCAGGCAGAACTTTTGGTTGCTATGGAGAACCAACACACCATAGATATGGCCGATTCCGAGATGT TGAATCTGTACGAGAGGGTCCGTAAGCAGCTGAGGCAGAACGCTGAGGAGGACGGGAAGGGGTGCTTTGAGATCTACCACACCTGTGACA ATAGCTGCATGGAGTCGATCCGGAATAACACTTATGACCATAGTCAATACAGAGAGGAAGCCCTGCTCAATCGTTTAAATATTAATCCAGTAA AGTTATCTTCAGGGTACAAGGATATTATTCTCTGGTTCTCTTTTGGGGCTAGCTGTTTTGTGCTCCTGGCCGTTATAATGGGACTCGGGTTCT TCTGCCTAAAAAATGGCAACATGCGCTGTACCATTTGCATT
3039	ATGTACAAGATTATCGTGATCATCGCACTGCTGGGCGCCGTTAAAGGGCTGGACAAAATATGCTTGGGACACCACGCCGTAGCTAATGGAA CAATCGTCAAAACCCTGACTAATGAGCAAGAGGAGGTCACAAATGCAACGGAAACAGTCGAATCCACTGGCATTAATAGGTTGTGCATGAA GGGCCGAAAGCACAAAGATCTCGGAAATTGTCATCCCATCGGGATGCTGATTGGCACCCCTGCCTGTGATTTGCACTTAACAGGGACTTGG GACACTCTGATAGAGCGTGAAAACGCAATTGCATATTGCTACCCAGGGGCAACGGTCAACGTGGAGGCTCTCCGACAGAAAATTATGGAAA GCGGAGGGATCGATAAGATTTCGACAGGCTTCACGTACGGCTCCTCAATCAACTCTGCCGGTACGACGCGCGCGTGTATGCGGAATGGAG GGAACAGCTTTTACGCAGAACTGAAGTGGCTCGTAAGCAAAAGTAAAGGGCAAAATTTCCCTCAGACTACAAATACCTACCGCAACACAGAC ACAGCGGAGCACCTGATTATGTGGGGAATTCACCATCCCTCAAGTACCCAGGAGAAGAACGACTTATATGGCACGCAGAGTCTTTCCATTTC TGTGGGGTCTAGCACCTATAGGAACAATTTCGTCCCAGTTGTCGGCGCTAGGCCTCAGGTGAATGGACAAAGCGGCCGCATAGATTTCCAC TGGACCCTGGTCCAGCCCGGCGATAATATCACATTTTCTCACAACGGAGGACTGATTGCTCCTAGCAGAGTATCAAAACTGATTGGGAGAG GCCTTGGCATTCAGTCTGACGCCCCCATTGACAATAATTGTGAGTCTAAGTGTTTTTGGCGGGGAGGAAGCATCAATACCAGACTTCCCTTT CAAAACCTGAGTCCGCGAACTGTCGGACAGTGCCCCAAATACGTTAATAGGCGCTCCCTAATGCTTGCTACAGGTATGAGGAATGTACCAG AGTTAATTCAGGGGCGCGGCCTTTTTGGCGCCATCGCCGGCTTCCTGGAGAACGGCTGGGAGGGCATGGTGGATGGGTGGTATGGCTTC CGCCATCAGAACGCACAGGGGACGGGCCAGGCTGCAGATTATAAGTCAACCCAGGCTGCAATAGACCAAATTACTGGGAAGCTGAATAGG CTAGTGGAGAAGACCAACACTGAGTTTGAATCAATTGAGAGCGAATTTAGTGAGATCGAGCACCAGATCGGTAACGTGATTAACTGGACGA AAGATTCAATTACTGACATTTGGACATATCAGGCTGAGTTGCTGGTGGCGATGGAGAACCAGCACACCATAGATATGGCGGATAGCGAGAT GCTTAATCTGTACGAACGGGTGCGCAAGCAGCTCCGCCAGAACGCTGAAGAAGATGGCAAGGGGTGCTTTGAGATTTACCACGCTTGCGA TGATTCATGCATGGAGTCCATTCGGAATAATACATACGACCACACTCAGTACAGGGAAGAAGCCCTACTGAACCGGTTGAACATTAACCCTG TGACCCTCAGTTCAGGGTACAAAGACATAATACTTTGGTTCTCCTTCGGGGCCTCATGTTTCGTGCTGCTGGCAGTCGTTATGGGCCTATTC TTCTTCTGCTTGAAGAATGGAAATATGCGTTGCACCATCTGCATC
3040	ATGTACAAAATCGTCGTCATCATTGCACTTCTGGGAGCCGTGAAGGGGCTGGATAAGATCTGCCTCGGCCATCATGCCGTCGTGAACGGTA CCATAGTGAAGACATTGACGAACGAGCAAGAGGAAGTCACGAACGCCACCGAGACGGTGGAATCCACGGGACTGAATCGTTTGTGCATGA AAGGCAGGAACCATAAGGATCTGGGAAATTGCCATCCCATAGGCATGCTTATTGGCACGCCAGCCTGTGACCTACACTTAACCGGCACTTG GGACACTTTGATCGAGCGCGAGAATGCCATCGCCTACTGTTATCCAGGAGCCACCGTAAACGAAGAGGCTCTGAGACAGAAGATCATGGA GAGTGGCGGTATAAACAAAATCAGCACCGGCTTCACCTACGGGAGCTCTATCAACTCAGCCGGTACTACTAGAGCTTGCATGCGCAACGGC GGAAACTCCTTCTACGCAGAACTTAAGTGGCTCGTCAGCAAGAGCAAGGGCCAGAACTTTCCTCAGACAACCAACACCTACCGGAATACCG ATACTGCCGAGCACTTGATAATGTGGGGCATCCACCACCCTTCCTCTACTCAGGAAAAGAACGATCTGTATGGTACCCAGTCTCTGTCCATA TCAGTGGGTTCTTCAACATACCAGAACAATTTCGTCCCTGTGGTAGGAGCAAGGCCCCAGGTAAATGGCCAGAGCGGCCGCATCGACTTTC ATTGGACTCTGGTGCAGCCAGGGGACAATATTACCTTCAGTCACAACGGCGGCCTGATCGCACCATCCAGAGTGTCTAAACTGATAGGCCG AGGATTGGGAATTCAGTCAGACGCACCCATAGATAACAATTGCGAATCGAAGTGTTTCTGGCGCGGCGGCTCAATAAACACCCGACTTCCA TTCCAAAACCTCTCTCCCAGAACTGTGGGGCAGTGCCCCAAATACGTGAATAAGAGGTCGCTCATGTTAGCCACTGGTATGCGAAACGTCC CGGAACTGATGCAGGGACGAGGCCTTTTTGGGGCTATAGCAGGCTTCATCGAGAATGGATGGGAAGGCATGGTTGATGGGTGGTATGGCT TCCGCCATCAGAATGCTCAGGGGACAGGGCAAGCGGCCGATTACAAATCAACCCAGGCCGCTATAGATCAAATCACTGGCAAGCTGAATC GCCTCATCGAAAAAACAAATACGGAGTTCGAGTCCATTGAAAGCGAGTTTTCCGAGATTGAACACCAAATTGGCAACGTCATAAACTGGACT AAGGACTCTATTACCGACATATGGACTTACCAGGCAGAGCTTCTGGTCGCTATGGAGAACCAGCACACCATCGATATGGCCGATTCCGAAA TGTTAAACCTCTACGAAAGAGTGCGCAAGCAGCTCAGACAGAACGCCGAAGAAGATGGCAAAGGGTGCTTTGAAATCTACCATGCCTGCGA CGATAGTTGTATGGAGTCGATCCGCAATAACACTTACGATCACAGTCAATATCGGGAGGAGGCACTGTTAAACAGGCTGAACATTAACCCG GTAACCCTCTCTAGTGGTTACAAAGACATCATACTTTGGTTTTCGTTTGGGGCATCTTGTTTTGTGCTCTTGGCCGTAGTGATGGGACTTGTT TTCTTTTGTCTCAAGAACGGCAATATGCGGTGTACTATTTGCATC

3041	ATGTATAAAATCGTCGTGATCATCGCCCTGTTAGGTGCGGTCAAGGGCTTGGACAAGATTTGTCTGGGACATCACGCAGTCGCCAACGGAA CCATCGTTAAAACGCTGACTAACGAGCAGGAAAAAGTTACTAATGCCACAGAGACCGTTGAGTCAACCGGCCTCAATAGGCTTTGCATGAAA GGACGGAAGCATAAAGACCTGGGCAACTGCCATCCTATTGGGATGTTAATCGGTACCCCCGCATGCGACCTGCATCTGACTGGAACCTGG GACACCATTATTGAGCGCGAAAATGCCATCGCATATTGTTACCCCGGAGCTACTGTGAATGAAGAGGCCCTGAGGCAGAAGATCATGGAGT CCGGTGGAATTGACAAGATCTCCACTGGATTTACCTACGGGAGTTCTATCAATAGTCCAGGTACCACAAGAGCCTGTATGAGAAATGGGGG CAACAGCTTTTACGCCGAATTAAAATGGCTGGTGAGCAAGAGTAAGGGACAAAATTTTCCTCAGACTACAAACACCTACAGAAATACAGATA CGGCTGAGCACCTTATCATGTGGGGCATCCACCATCCGTCATCCACCCAAGAGAAGAATGATCTTTACGGAACCCAGAGCTTGAGCATCTC CGTGGGTTCATCCACCTACCGGAACAACTTCGTGCCAGTGGTGGGGGCTAGGCCACAGGTGAACGGCCAGTCAGGCCGCATCGACTTCCA CTGGACACTCGTGCAGCCTGGAGATAACATCACTTTCTCCCACAATGGAGGACTGATAGCCCCATCTCGCGTTAGCAAGCTGATTGGCCGG GGTCTCGGAATTCAATCTGACGCCCCCATAGACAATAACTGTGAAAGCAAGTGCTTCTGGCGTGGGGGGTCCATTAATACCCGTCTACCTTT TCAGAATCTGTCCCCCCGCACCGTCGGACAATGCCCTAAGTACGTGAATAAAAGGTCCCTTATGCTAGCTACGGGGATGCGGAATGTTCCG GAGCTAATCCAGGGGCGCGGGCTGTTCGGAGCCATCGCAGGTTTTCTCGAGAATGGCTGGGAGGGAATGGTGGACGGTTGGTACGGATT CCGCCACCAGAATGCCCAAGGCACAGGACAGGCAGCTGACTACAAATCAACGCAAGCTGCCATCGACCAGATAACAGGAAAACTGAACAG ACTCGTCGAGAAGACCAATACAGAGTTCGAATCCATCGAGTCTGAATTTAGCGAAATCGAGCACCAGATCGGAAACGTTATCAACTGGACAA AAGACTCCATCACCGATATCTGGACTTACCAGGCTGAACTTCTTGTTGCCATGGAGAATCAGCATACTATTGATATGGCCGACTCAGAGATG CTCAACCTCTATGAACGGGTCCGGAAACAATTACGACAGAACGCGGAGGAGGACGGCAAAGGTTGCTTTGAGATATATCACGCGTGCGAC GATTCGTGCATGGAATCTATCCGCAATAATACTTACGACCACTCTCAGTACCGGGAGGAAGCACTCCTGAATCGGTTGAACATCAACCCCGT GACACTTTCTTCCGGCTATAACGATATAATCCTGTGGTTTTCCTTTGGAGCATCCTGTTTTGTACTCCTCGCCGTAGTGATCGGGCTGGTCTT TTTCTGCCTTAAGAACGGGAACATGAGGTGTACTATATGTATT
3042	ATGTACAAGGTGGTGGTCATAATCGCCTTGCTAGGCGCCGTGAGAGGGCTCGACAAAATTTGCTTGGGGCATCATGCAGTCGCTAATGGAA CCATCGTGAAAACACTCACCAACGAGCAAGAAGAGGTTACTAACGCCACCGAGACCGTAGAGAGTAAATCCCTGGGGAAGCTGTGTATGAA GGGCCGGTCTTATAATGATCTGGGCAATTGCCATCCCATCGGAATTCTGATCGGGACCCCTGCATGTGATCTGCATCTAACTGGGACGTGG GATACTCTGATCGAGCGCGAGAACGCAGTTGCTTACTGTTATCCGGGCGCCACCGTGAACGAGGAGGCACTCCGACAGAAAATTATGGAAT CGGGTGGAATTAGCAAGATATCTACTGGGTTCACCTATGGCACTAGCATCAACTCCGCTGGGACAACCAAAGCGTGTATGAGAAACGGGGG GAATTCTTTCTACGCCGAACTGAAATGGCTGGTGTCCAAAAATAAAGGGCAGAATTTTCCTCAGACCACCAACACATACCGGAACACCGACA CCGCCGAGCACCTGATCATCTGGGGCATTCATCACCCGAGCTCGACTCAAGAGAAGAACGATCTCTACGGGACTCAGTCGCTGAGTATCAG CGTGGGAAGCAGTACTTACCAGAACAATTTTGTACCCGTGGTGGGAGCCCGGCCTCAAGTCAACGGCCAGTCCGGCAGGATTGACTTCCA CTGGACCTTACTGCAGCCAGGGGACAACATCACCTTTTCACACAATGGTGGACTCATCGCACCCTCCAGAGTTAGCAAACTGATTGGAAGA GGGCTTGGTATTCAGTCGGAAGCCCCTATTGATAATGGTTGTGAGAGTAAGTGCTTCTGGAAAGGAGGCAGCATAAATACCAAACTCCCCTT CCAGAACCTCTCTCCCAGAACCGTCGGCCAGTGCCCTAAGTACGTGAATAAACGCTCGTTGATGCTCGCGACAGGAATGAGGAACGTGCCT GAAATTATGCATGGTCGCGGACTGTTCGGAGCTATCGCCGGGTTCATCGAGAATGGCTGGGAAGGGATGGTGGACGGTTGGTACGGGTTT CGACACCAGAACGCACAGGGCACTGGCCAGGCAGCAGATTATAAGTCCACTCAGGCCGCAATCGACCAGATTACCGGCAAATTAAATAGAT TGATCGAGAAAACGAATACTGAGTTCGAGAGCATCGAATCGGAGTTCTCTGAAATCGAGCATCAGATTGGAAACATTATAAATTGGACCAAG GACAGCATTACAGATATTTGGACGTATCAGGCAGAGCTGCTGGTGGCCATGGAAAACCAGCATACGATCGACATGGCCGATTCCGAAATGC TCAACCTGTATGAAAGGGTGAGGAAACAGCTGAGACAAAACGCAGAAGAAGACGGCAAAGGTTGCTTCGAAATTTACCACGCTTGCGACGA TTCATGCATGGAGAGCATTCGGAATAATACCTATGATCATAGCCAGTACAGAGAGGAGGCCCTGCTAAATACACTGAATATAAATCCCGTAA AACTGAGCTCTGGCTATAAGGACATTATTCTGTGGTTTAGTTTCGGAGCATCCTGCCTCATCCTCCTGGCAGTGGTCATGGGCCTCGTGTTC TTTTGTCTGAAGAACGGAAATATGCGATGTACTATTTGTATC
3043	ATGTACAAAATAGTCGTGATCATCGCACTGCTGGGGGCCGTGAAAGGTCTCGATAAGATTTGCCTTGGACACCATGCCGTGGCTAACGGGA CTATTGTGAAGACACTTACCAATGAGCAGGAGGAAGTTACCAACGCTACAGAAACTGTCGAGTCGACTGGTATTAATAGGCTGTGTATGAAG GGACGTAAGCATAAACACCTCGGGAATTGCCACCCTATAGGCATGCTGATAGGCACCCCAGCATGTGACCTGCACCTCACACGCATGTGG GACACATTAATCGAAAGGGAGAATGCCATAGCCTATTGCTACCCAGGTGCAACAGTTAATGTTGAAGCATTGCGGCAGAAAATCATGGAGTC CGGCGGCATTAACAAGATCTCTACGGGATTTACGTACGGCTCAAGCATTAATAGTGCCGGGACTACTCGAGCATGCATGAGGAATGGAGGG AACAGTTTCTACGCCGAGCTGAAGTGGCTCGTGTCTAAAAGTAAAGGGCAGAACTTCCCTCAGACCACTAATACCTACAGAAACACAGACAC CGCAGAACACTTAATAATGTGGGGCATACATCACCCCAGCAGCACTCAGGAGAAAAACGACCTGTACGGTACCCAGTCCCTGTCCATTTCC GTGGGAAGTTCTACGTATCGGAATAACTTCGTCCCGGTGGTGGGAGCCAGGCCGCAGGTTAACGGACAAAGCGGGCGAATTGATTTCCAC TGGACACTGGTGCAGCCTGGCGACAATATAACTTTCTCCCATAATGGCGGACTGATCGCTCCATCCCGGGTCAGCAAGCTAATCGGTAGAG GGCTGGGGATCCAATCAGACGCACCTATTGATAACAACTGCGAGAGTAAGTGCTTCTGGCGAGGCGGTAGCATAAACACACGTTTGCCATT TCAGAATCTTAGTCCTCGTACTGTGGGCCAATGTCCAAAGTATGTCAACAGGCGCAGTTTAATGTTAGCTACTGGAATGCGCAACGTACCTG AGCTCATTCAAGGTCGAGGCCTGTTCGGAGCCATTGCTGGGTTCCTTGAGAATGGGTGGGAGGGCATGGTCGACGGATGGTACGGATTCC GCCACCAGAACGCCCAGGGCACAGGCCAGGCAGCCGATTATAAAAGTACCCAGGCTGCCATCGACCAGATCACTGGGAAACTCAACCGTC TGGTCGAAAAGACTAACACCGAATTTGAGTCTATTGAATCAGAATTCAGCGAAATTGAGCACCAGATCGGCAATGTGATCAATTGGACGAAA GACTCTATTACCGACATTTGGACCTACCAGGCCGAGTTACTGGTGGCCATGGAGAACCAGCACACCATCGACATGGCCGATAGTGAAATGC TTAACCTCTACGAAAGAGTCCGAAAACAACTCCGCCAGAACGCGGAGGAGGATGGAAAGGCCTGCTTCGAAATCTATCATGCCTGCGACGA CTCCTGTATGGAAAGCATTCGCAACAACACCTACGACCACAGTCAGTACAGGGAGGAGGCGCTGCTTAATAGGCTGAACATCAATCCGGTA ACACTGTCATCGGGATACAAGGACATCATACTCTGGTTCTCATTCGGCGCGTCCTGTTTCGTACTGCTCGCAGTCGTAATGGGGCTCTTCTT CTTCTGCTTAAAGAACGGCAACATGCGCTGCACTATCTGCATT

3044	ATGTATAAAATTGTGGTCATAATTGCACTGCTCGGCGCAGTCAAGGGTCTGGATAAGATTTGCCTTGGTCACCATGCCGTAGCCAACGGAAC CATTGTGAAGACCCTTACAAATGAACAAGAAGAAGTGACTAATGCAACTGAGACCGTGGAATCTACAGGTATCAATCGCCTGTGTATGAAGG GGCGAAAACATAAAGATCTCGGCAATTGTCACCCCATTGGGATGCTGATTGGCACCCCGGCATGTGACCTACATTTGACCGGAATGTGGGA TACACTAATCGAACGGGAGAATGCTATTGCGTACTGTTACCCTGGCGCTACCGTCAATGTGGAGGCCCTGCGCCAAAAAATCATGGAATCT GGGGGAATCAATAAGATCTCCACTGGCTTCACTTATGGATCCAGCATTAACAGCGCTGGCACAACTCGCGCCTGTATGCGCAATGGGGGCA ATTCCTTCTATGCAGAGCTTAAGTGGTTAGTGAGCAAGTCGAAAGGCCAAAACTTCCCTCAAACAACCAATACATATAGGAACACCGATACC GCTGAACATCTCATTATGTGGGGAATACATCATCCCAGCTCCACACAAGAGAAGAATGATTTGTACGGGACCCAATCCTTGTCTATTAGCGT AGGCTCCTCCACATACCGGAACAACTTCGTCCCAGTCGTGGGGGCACGGCCTCAGGTTAACGGACAGTCCGGCCGTATTGACTTCCACTG GACTCTGGTGCAACCAGGGGACAACATCACTTTCTCTCATAACGGGGGTCTGATCGCACCTAGTAGGGTGTCCAAACTTATAGGAAGGGGT CTCGGCATCCAAAGCGACGCCCCCATCGATAATAATTGCGAAAGCAAGTGTTTTTGGAGGGGAGGGAGCATCAATACTCGGTTACCATTTC AGAATTTAAGTCCAAGAACAGTAGGACAATGCCCTAAGTATGTTAATAGACGTTCCCTCATGTTAGCAACTGGAATGAGGAACGTTCCGGAG CTGATTCAGGGAAGAGGACTCTTCGGGGCAATCGCCGGTTTTTTGGAGAACGGATGGGAGGGCATGGTGGATGGTTGGTACGGATTCCGC CATCAGAACGCCCAGGGCACCGGGCAAGCTGCAGACTATAAATCAACACAAGCAGCAATTGATCAAATCACCGGCAAGTTGAACAGGCTGG TCGAGAAAACCAACACTGAATTCGAGTCCATCGAGAGCGAGTTCTCTGAAATAGAGCATCAGATTGGAAACGTTATCAACTGGACCAAGGAT AGTATCACGGACATTTGGACTTATCAAGCCGAGCTGCTGGTCGCCATGGAGAATCAACATACTATCGACATGGCGGATAGTGAAATGCTCAA CCTGTACGAGCGCGTGCGCAAGCAACTGCGCCAGAACGCTGAAGAGGACGGTAAAGGTTGCTTTGAAATTTATCACGCTTGTGATGACTCC TGCATGGAGTCGATCCGCAATAATACGTACGATCACAGCCAGTACCGGGAGGAGGCTTTGCTGAACCGGCTGAACATCAACCCTGTAACCT TGTCTTCTGGCTACAAGGATATTATACTCTGGTTCAGCTTCGGCGCCAGCTGCTTCGTCCTGCTGGCCGTGGTTATGGGACTCTTCTTTTTC TGTCTGAAAAATGGTAATATGCGGTGCACAATTTGTATC
3045	ATGTACAAGATTGTGGTAATCATCGCTTTACTTGGGGCAGTGAAAGGATTGGACAAGATTTGTTTGGGGCACCATGCAGTGGCCAACGGCA CCATTGTTAAAACACTGACAAATGAGCAGGAAGAGGTTACCAATGCAACGGAAACCGTGGAATCTACTGGAATCAACAGACTGTGCATGAAG GGCCGCAAGCATAAAGACTTGGGGAATTGCCACCCAATTGGAATGCTTATTGGAACACCAGCTTGCGATTTACATCTGACTGGCATGTGGG ACACCCTCATCGAGCGGGAAAATGCTATCGCTTATTGCTACCCGGGGGCCACAGTGAATGTGGAAGCACTACGGCAAAAAATCATGGAATC AGGCGGAATCAACAAAATCTCAACCGGCTTCACCTATGGGTCATCCATCAATAGTGCTGGCACCACACGCGCCTGCATGCGAAATGGTGGA AACTCATTCTATGCGGAACTGAAATGGCTAGTGTCAAAAAGCAAAGGGCAGAATTTTCCACAGACTACGAACACTTATAGAAATACTGACACA GCTGAGCATCTTATTATGTGGGGCATCCATCATCCAAGCAGTACACAGGAAAAGAATGACCTATACGGTACACAGTCCCTGTCTATTTCCGT GGGATCCTCCACATACAGAAACAACTTTGTCCCTGTGGTGGGAGCAAGGCCTCAGGTCAACGGACAGTCTGGCCGCATCGATTTCCATTGG ACCCTGGTTCAACCGGGCGATAACATCACCTTCAGCCACAATGGAGGCCTGATCGCCCCTAGCAGGGTGTCAAAGCTTATAGGCAGGGGT CTTGGGATCCAATCCGACGCCCCCATAGACAATAACTGCGAGTCGAAATGTTTCTGGCGGGGTGGTTCCATTAATACTCGTCTCCCCTTCCA GAATCTTTCTCCCAGAACTGTTGGGCAATGCCCCAAGTACGTAAATCGTAGAAGTCTTATGCTCGCAACCGGTATGCGCAACGTGCCTGAG CTCATTCAGGGCCGGGGCCTCTTTGGGGCTATTGCCGGCTTCCTTGAAAACGGCTGGGAGGGTATGGTGGATGGCTGGTACGGATTCCGC CACCAGAATGCCCAAGGCACAGGTCAGGCTGCCGACTATAAGTCAACGCAGGCTGCCATCGATCAGATCACAGGCAAGTTGAACAGGCTA GTGGAGAAGACCAACACAGAGTTTGAGAGTATTGAGAGCGAGTTTTCCGAAATCGAACACCAGATAGGCAACGTTATCAACTGGACAAAGG ACAGTATTACAGACATTTGGACATATCAGGCTGAACTGTTGGTGGCCATGGAGAACCAGCACACTATCGATATGGCAGACTCCGAGATGCT GAATTTGTACGAAAGGGTCCGGAAGCAGCTCCGCCAGAACGCCGAAGAGGATGGGAAAGGCTGCTTTGAAATTTACCACGCTTGCGACGA CTCCTGTATGGAATCCATTCGAAATAATACATATGATCACAGCCAGTACCGGGAACAAGCTTTACTTAATCGATTGAATATCAATCCAGTTAC CCTTTCCTCCGGCTACAAGGACATCATTCTGTGGTTCTCCTTCGGTGCCTCGTGCTTCGTCCTGCTTGCAGTTGTTATGGGGCTGTTTTTTTT CTGTCTGAAGAACGGGAATATGCGGTGCACTATTTGCATC
3046	ATGTATAAAATCGTGGTTATCATCGCCCTGTTGGGCGCAGTCAAAGGCTTAGACAAAATTTGCCTGGGCCACCATGCCGTGGCCAATGGCA CAATTGTAAAAACTTTGACTAATGAGCAGGAGGAAGTCACTAACGCCACGGAAACCGTGGAGTCCACTGGAATCAACCGCCTGTGCATGAA GGGACGGAAGCATAAGGACCTCGCTAATTGTCATCCGATAGGTATGCTCATCGGCACCCCCGCTTGCGACTTACATTTGACAGGAATGTGG GACACTCTGATTGAGAGGGAGAACGCCATTGCCTACTGTTACCCCGGCGCAACAGTCAATGTGGAGGCGCTGCGTCAAAAGATAATGGAG AGTGGAGGCATCAACAAAATTTCAACAGGGTTCACATACGGGAGCTCAATTAACTCTGCGGGCACGACCAGGGCTTGCATGAGAAACGGCG GTAATTCCTTTTATGCGGAGTTGAAATGGCTGGTCTCCAAGTCCAAGGGTCAGAATTTCCCACAAACCACAAATACTTACCGAAATACGGAC ACTGCCGAGCATCTGATAATGTGGGGAATACACCATCCTAGTTCTACGCAAGAGAAAAATGACCTCTATGGGACGCAATCCTTGAGTATCAG CGTTGGCTCCTCAACCTACCGGAACAACTTCGTCCCAGTTGTAGGAGCTCGACCTCAGGTGAATGGACAGTCAGGTCGTATCGATTTTCATT GGACCCTCGTGCAACCTGGGGACAACATAACCTTCTCCCACAACGGCGGGCTGATAGCACCCAGTCGTGTCTCCAAATTGATCGGGAGGG GGCTGGGCATCCAATCAGATGCACCAATTGATAATAATTGTGAGAGCAAGTGCTTCTGGCGTGGGGGTAGCATCAACACTCGCCTCCCCTT TCAGAACTTGTCACCCCGGACCGTCGGTCAGTGTCCCAAGTATGTTAATCGGCGTTCATTAATGCTGGCAACCGGTATGCGCAACGTCCCC GAGCTAATCCAAGGAAGAGGGCTTTTTGGGGCTATAGCAGGATTTTTAGAGAACGGTTGGGAGGGCATGGTGGACGGATGGTACGGCTTT CGGCACCAGAATGCGCAGGGGACAGGGCAAGCTGCCGATTATAAGTCCACTCAAGCTGCTATTGATCAGATTACAGGCAAGCTTAATCGCC TGGTCGAGAAGACTAACACAGAGTTCGAGAGCATTGAGTCAGAATTCTCTGAAATTGAGCACCAAATCGGGAATGTAATAAATTGGACAAAG GACAGTATTACCGACATCTGGACTTATCAAGCCGAGCTCCTGGTTGCTATGGAGAATCAGCACACAATCGACATGGCCGATAGCGAGATGC TAAACCTGTACGAAAGGGTGCGCAAGCAGCTGAGGCAAAACGCTGAGCAAGATGGCAAGGGCTGTTTCGAGATTTATCACGCGTGTGATG ATTCATGCATGGAATCAATTAGGAACAACACTTACGATCATTCTCAGTACCGGGAGGAGGCACTCCTGAACAGGCTCAACATAAACCCTGTC ACCCTGTCTTCAGGCTATAAGGACATTATACTATGGTTTTCTTTTGGCGCCTCCTGTTTTGTGCTCCTCGCCGTTGTAATGGGCCTTTTTTTTT TTTGCCTGAAGAACGGTAATATGCGGTGTACTATTTGCATT

3047	ATGTACAAGATTGTAGTGATTATCGCACTGCTAGGCGCGGTGAAGGGTCTCGACAAGATTTGCCTGGGCCATCACGCGGTGGCCAACGGG ACTATCGTGAAGACTCTGACCAATGAACAAGAGGAAGTTACTAACGCCACCGAAACTGTGGAGTCCACAGGCATCAATCGCTTGTGTATGAA GGGAAGGAAGCATAAAGACCTGGGTAATTGTCACCCCATAGGGATGCTTATCGGTACACCAGCTTGTGATCTGCATTTAACGGGGATGTGG GACACACTTATAGAACGTGAGAATGCCATTGCCTATTGTTACCCCGGAGCTACCGTAAACGTGGAGGCCCTGCGCCAGAAGATCATGGAGT CACGGGGCATCAATAAGATCAGTACCGGCTTTACCTATGGTAGCTCTATCAACTCCGCCGGCACTACCAGGGCCTGTATGAGGAATGGGG GAAACAGCTTCTATGCTGAGCTGAAGTGGCTAGTTAGCAAGAGCAAGGGCCAGAACTTCCCTCAAACTACAAATACGTATCGGAACACCGA CACAGCGGAGCACCTGATAATGTGGGGAATCCACCACCCATCGTCCACCCAAGAAAAGAACGATCTGTACGGTACTCAGTCCCTGAGTATT TCCGTGGGCTCCAGTACGTACCGAAATAATTTCGTCCCAGTGGTCGGGGCAAGGCCGCAAGTGAATGGCCAGTCAGGACGGATAGACTTT CATTGGACACTGGTGCAACCCGGCGATAACATCACCTTCAGCCATAACGGGGGGCTGATCGCCCCGAGCCGAGTTAGCAAACTTATCGGC CGGGGCCTGGGCATCCAGAGTGATGCACCCATCGACAATAATTGCGAGAGCAAATGTTTTTGGAGGGGTGGATCGATAAACACACGCCTC CCGTTCCAAAATCTCAGTCCTAGAACAGTGGGGCAGTGTCCTAAGTATGTGAATCGCCGGAGCCTTATGTTGGCCACTGGGATGCGAAATG TGCCCGAACTAATTCAAGGGCGAGGACTGTTCGGCGCCATTGCCGGCTTCCTGGAGAATGGCTGGGAAGGCATGGTTGATGGTTGGTATG GCTTTCGCCATCAGAATGCACAGGGGACAGGCCAGGCCGCGGATTACAAATCGACCCAAGCTGCAATTGATCAGATCACAGGGAAACTGA ATCGACTTGTGGAGAAAACTAATACCGAATTTGAATCTATCGAGTCCGAATTTAGCGAAATCGAGCACCAGATCGGTAACGTGATCAATTGG ACTAAGGACTCCATTACAGACATATGGACATATCAGGCCGAGTTGTTGGTGGCAATGGAGAACCAGCATACTATCGACATGGCCGATTCTGA GATGCTAAATCTGTATGAACGCGTCCGGAAACAGCTCAGACAGAATGCTGAGGAAGATGGGAAGGGATGCTTCGAAATATACCACGCCTGC GACGACAGCTGTATGGAGTCAATTCGCAACAACACTTACGATCACTCCCAATACCGGGAGGAGGCCCTTCTTAACAGACTCAACATTAATCC TGTGACTCTGTCATCCGGATACAAAGACATAATACTCTGGTTTAGCTTTGGCGCAAGTTGTTTTGTGCTGCTCGCAGTGGTGATGGGGTTGT TTTTTTTCTGCCTGAAAAATGGTAATATGCGATGCACTATCTGCATC
3048	ATGTACAAAATTGTGGTGATAATTGCATTGCTCGGAGCAGTCAAAGGGCTGGACAAAATATGTTTGGGGCACCACGCCGTGGTTAATGGTA CCATCGTTAAGACTCTGACCAACGAACAGGAAGAGGTGACCAATGCTACCGAGACCGTGGAAAGCACTGGGTTGAACAGGCTGTGCATGA AAGGAAGGAATCACAAGGACCTGGGAAACTGTCATCCAATCGGAATGCTCATTGGGACTCCCGCATGTGATCTGCATCTGACGGGCACCTG GGATACCTTGATAGAGAGGGAGAACGCCATAGCATACTGTTATCCCGGAGCCACCGTGAACGAGGAGGCACTGAGCCAGAAAATTATGGA ATCTGGAGGGATCAATAAAATTAGCACAGGGTTTACCTATGGCTCAAGTATCAATAGCGCCGGGACTACACGGGCCTGCATGCGAAATGGA GGGAACTCCTTCTACGCTGAACTAAAGTGGCTGGTCAGTAAGTCCAAAGGGCAGAATTTCCCGCAGACAACCAACACTTACCGCAATACCG ATACGGCTGAACACTTAATCATGTGGGGCATCCATCATCCCTCTAGCACACAAGAGAAAAACGACCTGTATGGAACACAAAGCCTCAGCATC TCCGTGGGTTCATCTACATATCAGAACAACTTCGTGCCCGTCGTTGGCGCCCGACCTCAGGTGAATGGTCAATCGGGCCGGATTGACTTTC ACTGGACCCTCGTCCAGCCAGGTGATAACATCACCTTCAGCCACAACGGCGGACTCATCGCCCCATCTCGTGTGTCCAAGCTCATTGGCAG AGGGCTTGGCATCCAGTCTGATGCACCCATCGATAACAATTGCGAGAGCAAGTGTTTCTGGCGCGGTGGATCAATAAATACGCGGCTCCCC TTTCAGAATCTTTCTCCCAGAACCGTAGGCCAGTGTCCAAAGTATGTAAACAAAAGATCATTGATGCTGGCCACAGGAATGAGGAATGTGCC TGAGCTGATGCAGGGCAGAGGCCTGTTCGGTGCGATCGCCGGCTTCATCGAAAATGGCTGGGAGGGGATGGTGGACGGATGGTATGGCT TCCGCCACCAGAACGCTCAGGGGACCGGACAAGCTGCGGATTATAAGAGCACTCAGGCGGCAATCGACCAGATCACCGGAAAACTCAACC GTCTCATCGAGAAGACAAACACCGAATTTGAGTCTATAGAATCTGAGTTCAGTGAAATCGAACACCAAATCGGCAACGTCATAAATTGGACT AAGGACTCCATCACCGATATCTGGACCTATCAAGCCGAGCTGCTGGTTGCCATGGAGAATCAGCACACGATCGACATGGCTGATTCTGAAA TGCTGAACCTCTATGAAAGAGTTCGGAAACAGCTCAGACAAAACGCCGAAGAGGACGGCAAAGGATGTTTTGAGATCTATCATGCCTGCGA TGATAGTTGTATGGAGAGCATCAGGAATAATACATCACCACTCCCAGTATCGCCAGGAACCACTGCTCAATCGGTTAAATATAAATCCAG TGACGCTTAGTAGTGGGTATAAGGACATAATCCTCTGGTTCTCTTTCGGCGCCAGCTGTTTCGTTCTGCTGGCCGTGGTCATGGGACTTGTG TTCTTTTGTCTTAAAAATGGGAACATGAGATGCACAATATGTATT
3049	ATGTATAAAATAGTGGTGATAATCGCGCTACTGGGCGCAGTGAAGGGGCTCGATAAGATTTGCTTGGGTCATCATGCTGTTGCAAACGGCA CCATCGTTAAAACACTGACCAACGAGCAGGAGGAGGTAACGAACGCCACGGAGACGGTGGAGTCTACCGGGATCAACCGTCTGTGTATGA AGGGTCGCAAGCATAAGGATTTAGGAAATTGTCATCCTATCGGAATGCTTATAGGTACCCCGGCCTGCGATCTCCATCTGACAGGCATGTG GGATACTCTCATCGAAAGAGAGAATGCCATCGCGTATTGTTACCCTGGGGCCACAGTAAATGTCGAAGCATTGAGACAGAAAATCATGGAG TCCGGAGGCATTAATAAGATTTCCACCGGCTTCACCTACGGATCTTCCATCAATTCCGCAGGCACAACCAGAGCCTGCATGCGGAATGGAG GGAATTCCTTTTACGCCGAATTGAAATGGCTTGTGTCAAAGTCAAAGGGACAGAACTTTCCCCAAACTACGAACACATATAGAAATACAGATA CCGCAGAACACTTAATTATGTGGGGCATCCACCACCCGTCAAGTACTCAGGAGAAGAATGACCTTTACGGCACCCAGTCTCTTTCAATCTCA GTCGGGAGCTCCACATATAGAAACAACTTTGTACCTGTGGTCGGTGCAGGACCGCAGGTGAATGGCCAATCAGGAAGGATCGATTTTCACT GGACACTGGTCCAGCCTGGTGACAACATCACGTTCTCACACAATGGGGGCCTAATAGCCCCCTCCCGCGTGAGCAAGCTAATTGGCCGCG GATTGGGTATCCAGAGCGACGCCCCTATAGACAATAACTGTGAGAGCAAGTGCTTCTGGCGCGGAGGGTCAATTAACACCCGGCTCCCCTT CCAGAACCTGTCCCCCCGGACGGTGGGCCAGTGCCCAAAATACGTCAATCGGCGCTCCCTGATGTTGGCCACAGGCATGCGGAATGTTCC TGAACTGATACAGGGTCGGGGGTTGTTCGGAGCAATTGCCGGGTTTCTTGAAAACGGCTGGGAAGGTATGGTGGATGGTTGGTACGGGTT CAGGCACCAGAACGCTCAGGGTACCGGCCAGGCGGCTGACTACAAATCAACCCAGGCAGCAATCGACCAGATTACAGGAAAGCTGAATCG CCTTGTGGAGAAAACCAATACCGAATTTGAGTCCATCGAGTCTGAGTTCAGCGAGATAGAGCATCAGATTGGCAACGTCATCAACTGGACG AAGGACTCCATTACTGATATATGGACTTATCAGGCTGAACTGCTGGTAGCCATGGAGAACCAGCACACAATTGACATGGCTGACTCCGAGAT GCTGAATCTATACGAGCGCGTGAGGAAGCAGCTCAGACAGAACGCTGAGGAGGACGGTAAGGGTTGCTTCGAGATTTACCACGCCTGTGA TGACTCCTGTATGGAGTCAATCAGGAACAATACCTATGACCACTCTCAATATCGGGAGGAAGCGCTTCTCAACCGTCTTAACATTAATCCCG TGACACTGTCTTCAGGTTACAAGGATATCATCCTGTGGTTCAGCTTTGGCGCCAGCTGTTTCGTGCTACTAGCAGTGGTGATGGGCCTGTTT TTTTTCTGCCTGAAAAATGGGAACATGAGGTGCACGATTTGCATC

3050	ATGTATAAAATCGTTGTAATTATTGCACTGCTTGGCGCCGTGAAGGGACTCGACAAGATCTGCCTGGGCCACCACGCTGTAGCCAACGGCA CCATTGTGAAGACCCTGACAAATGAGCAGGAGGAGGTTACCAATGCCACTGAGACCGTGGAATCCACAGGCATTAACCGCCTGTGTATGAA GGGAAGGAAGCACAAGGACCTTGGCAATTGTCACCCAATCGGCATGCTCATCGGCACACCAGCATGCGACTTGCACCTGACCGGAATGTG GGACACGCTCATAGAAAGAGAGAATGCCATTGCCTATTGCTACCCAGGAGCCACCGTTAACGTGGAGGCACTGAGGCAGAAGATCATGGA GAGCGGACGCATCAATAAAATTAGTACGGGCGTTTACATATGGCTCCTCTATAAATTCCGCCGGCACCACCCGGGCCTGCATGCGGAATGGC GGAAATAGTTTCTACGCTGAGTTGAAATGGCTGGTGAGCAAGAGTAAGGGGCAGAACTTTCCGCAAACTACCAATACGTATAGGAACACTG ACACCGCCGAGCACCTAATCATGTGGGGCATACATCATCCCAGCTCCACGCAGGAGAAGAACGACCTGTATGGAACTCAAAGCCTATCTAT CTCCGTGGGATCCAGTACATACCGGAATAACTTTGTGCCAGTAGTGGGCGCCAGACCGCAGGTGAACGGGCAATCCGGCAGGATCGACTT CCACTGGACACTGGTGCAACCAGGCGACAACATTACTTTCAGCCACAATGGTGGTTTAATAGCCCCCTCAAGGGTTTCCAAATTGATTGGGC GGGGGCTTGGTATTCAATCTGACGCTCCAATAGATAACAATTGTGAAAGTAAGTGTTTCTGGCGCGGCGGCTCTATTAATACTCGCCTTCCA TTCCAAAATCTCTCACCACGTACCGTTGGCCAATGTCCCAAATATGTGAACCGTAGGTCCTTGATGCTCGCAACGGGAATGAGGAACGTTCC GGAGCTGATCCAGGGCCGAGGACTCTTTGGTGCTATTGCCGGATTTCTGGAGAACGGGTGGGAGGGCATGGTGGATGGATGGTACGGCTT CCGACATCAGAATGCACAAGGCACCGGGCAGGCCGCCGATTACAAGTCCACGCAAGCTGCCATTGACCAGATTACGGGGAAGTTGAACCG GCTCGTCGAGAAGACCAATACTGAATTCGAATCCATTGAGAGCGAATTTTCTGAAATTGAACATCAGATTGGTAACGTCATCAATTGGACTAA GGACTCAATAACAGACATCTGGACTTATCAGGCTGAACTCCTCGTCGCGATGGAAAATCAGCACACCATCGATATGGCCGACTCTGAGATG CTGAACTTGTACGAGCGGGTGAGAAAACAGCTTCGACAGAATGCCGAGGAAGATGGCAAGGGCTGCTTCGAGATCTATCACGCTTGCGAT GACTCATGCATGGAGTCTATCCGGAACAACACTTATGACCACTCCCAATACCGCGAGGAGGCCCTGCTGAATAGGCTGAACATCAACCCAG TGACCTTGAGCTCAGCCTATAAGGACATAATCCTGTGGTTTAGCTTTGGGGCCTCGTGCTTTGTCCTGCTGGCCGTAGTTATGCGCCTTTTC TTCTTTTGTCTTAAGAATGGGAATATGAGGTGTACAATCTGTATC
3051	ATGTACAAGATCGTGGTTATCATCGCACTCCTCGGGGCTGTGAAGGGTCTCGACAAAATCTGCCTTGGTCATCACGCAGTTGCCAACCGAA CTATTGTTAAGACCTTGACTAATGAGCAGGAGGAGGTGACAAACGCCACAGAAACCGTGGAAAGTACGGGGATCAACCGTTTATGCATGAA GGGAAGGAAACACAAAGATCTCGGTAACTGCCATCCGATCGGTATGCTCATTGGCACTCCAGCTTGCGACTTGCACTTGACCGGTATGTGG GACACATTAATTGAACGCGAGAACGCCATCGCCTATTGCTACCCCGGAGCCACTGTGAACGTGGAAGCACTCCGACAGAAAATAATGGAAA GTGGGGGGATAAATAAGATCTCAACCGGATTTACATACGGTTCCTCAATTAATAGCGCCGGGACCACTAGAGCATGCATGCGAAATGGAGG TAATTCTTTTTACGCTGAGCTTAAATGGCTGGTGTCTAAAAGCAAGGGCCAGAATTTTCCACAGACGACGAACACTTATAGGAATACAGACAC AGCCGAACACCTGATCATGTGGGGAATCCACCACCCATCTAGCACACAGGAGAAAAACGACCTATATGGTACCCAAAGCCTGTCAATTAGC GTCGGCAGTAGTACATACCGCAATAATTTCGTTCCCGTTGTCGGAGCCAGGCCGCAAGTGAACGGGCAAAGTGGCAGAATTGATTTCCACT GGACACTGGTCCAGCCCGGGGATAACATTACATTCAGCCACAACGGCGGGCTTATCGCCCCCTCCCGTGTGAGTAAGCTGATCGGCCGGG GCTTAGGTATCCAGTCCGATGCTCCCATCGACAACAATTGTGAATCGAAGTGCTTTTGGCGGGGCGGCTCCATCAATACCCGGCTGCCTTT CCAGAACCTGAGCCCCAGGACCGTTGGACAGTGTCCCAAATACGTTAACCGGCGGAGTCTGATGCTTGCAACCGGGATGCGCAATGTGCC CGAACTGATCCAGGGGCGAGGACTGTTCGGGGCCATCGCCGGATTCCTGGAGAATGGCTGGGAGGGTATGGTGGACGGCTGGTACGGCT TTAGGCACCAAAATGCTCAAGGAACAGGGCAGGCTGCTGACTACAAATCAACCCAGGCCGCTATTGATCAAATTACTGGAAAGCTGAATAG ATTGGTGGAGAAAACTAATACTGAATTCGAGTCCATCGAAAGTGAGTTCAGCGAGATCGAGCACCAGATTGGTAACGTTATTAACTGGACCA AAGACAGCATCACCGACATTTGGACCTACCAGGCCGAGCTTCTAGTAGCCATGGAAAATCAACACACGATCGATATGGCTGATTCAGAGAT GCTGAATCTGTATGAGCGGGTCCGGAAACAGCTGAGACAGAATGCAGAGGAAGACGGTAAGGGGTGTTTTGAGATTTACCATGCCTGCGA TGATTCTTGCATGGAATCTATCAGGAACAACACCTATGACCACTCCCAATATCGCGAGGAGGCCCTGCTGAACAGGTTAAACATCAACCCCG TCACCCTGTCTTCAGGATATAAAGACATCATCTTATGGTTCTCCTTCGGAGCGTCCTGTTTTGTGCTGTTGGCAGTGGTCATGGGGCTTTTCT TTTTCTGTCTAAAAAACGGTAACATGCGGTGCACCATATGCATC
3052	ATGTATAAAATTGTAGTTATTATCGCCCTGTTGGGCGCTGTGAAGGGCTTGGACAAGATCTGCCTGGGACACCATGCAGTCGCAAATGGAA CAATTGTCAAGACACTGACTAATGAGCAGGAAGAGGTTACCAACGCCACAGAGACCGTAGAGTCCACCGGGATTAATCGGCTATGCATGAA AGGAAGAAAGCACAAACATCTCGGGAACTGTCATCCAATTGGTATGCTGATCGGAACCCCCGCCTGTGATCTCCATCTTACTGGCACTTGG GACACCTTGATTGAGAGGGAAAACGCCATTGCTTATTGTTACCCAGGAGCTACCGCCAACGTCGAGGCCCTAAGGCAGAAGATCATGGAGT CTGGAGGAATTGATAAGATCTCAACGGGGTTTACATACGGGTCTAGTATCAATAGCGCAGGCACTACCCGCGCTTGTATGAGGAACGGCGG CAACTCGTTTTACGCCGAGTTGAAATGGCTGGTCAGTAAATCAAAGGGGCAGAATTTTCCACAGACCACAAACACTTATAGGAATACCGATA CAGCTGAACACCTCATCATGTGGGGCATCCATCACCCCAGCTCCATTCAAGAGAAGAACGACTTGTATGGCACACAGTCATTGAGCATCAG TGTGGGCAGCAGCACATACAGAAACAACTTCGTGCCCGTGGTTGGGGCAAGACCTCAGGTAAACGGCCAGTCCGGTAGGATTGACTTCCA TTGGACGCTCGTGCAACCGGGAGATAACATCACCTTCAGCCACAATGGGGGACTCATCGCACCCTCTCGAGTTAGCAAACTGATCGGGCG AGGCTTAGGCATACAATCCGACGCGCCTATCGATAACAATTGCGAATCCAAATGTTTCTGGCGCGGCGGTAGCATTAATACTCGTCTGCCAT TCCAGAACCTTAGCCCCAGGACAGTGGGACAGTGTCCAAAATACGTGAATCGCAGAAGTCTCATGCTGGCTACCGGCATGAGGAACGTGC CAGAGCTTATACAGGGGCGCGGGCTTGTTCGGAGCGATCGCTGGCTTCTTGGAGAATGGCTGGGAGGGCATGGTCGACGGATGGTATGGTT TTCGACATCAAAACGCCCAGGGCACGGGGCAGGCCGCCGATTACAAGTCCACTCAGGCAGCAATCGACCAGATCACCGGTAAGCTTAATA GACTGGTCGAGAAGACAAATACCGAGTTCGAATCTATTGAATCCGAGTTCAGCGAAATCGAACACCAAATAGGGAACGTCATCAACTGGAC CATGGACAGCATCACCGATATATGGACATATCAGGCTGAGTTGCTTGTGGCCATGGAAAATCAGCACACAATCGATATGGCTGATTCCGAAA TGCTGAACCTGTACCAACGGGTAAGGAAGCAGCTGCGGCAGAACGCAGAAGACGGATGGGAAGGGCTGCTTCGAAATTTACCACGCTTGTG ATGACTCTTGTATGGAGTCTATTCGCAACAACACCTACGACCATTCTCAGTATCGTGAGGAGGCCCTTCTCAATCGTCTGAACATCAATCCG GTGACCCTATCTTCAGGATATAAGGATATTATACTCTGGTTCTCCTTCGGGGCAAGCTGCTTCGTCTTGCTGGCCGTCGTCATGGGGCTGGT TTTCTTCTGTTTAAAAAATGGAAATATGAGGTGCACCATTTGTATC

3053	ATGTACAAGATTGTAGTTATTATCGCTTTACTGGGAGCGGTCAAAGGTCTGGACAAGATATGCCTTGGACACCACGCAGTGGCTAACGGCA CAATCGTCAAAACTCTGACAAACGAGCAGGAGGAGGTCACTAATGCGACCGAAACGGTGGAGAGCACCGGAATTAATCGACTATGTATGAA GGGCAGGAAGCACAAGGATCTCGGGAACTGCCATCCCATCGGCATGCTGATAGGCACGCCAGCCTGCGATCTTCATCTGACCGGAATGTG GGATACCCTGATCGAGAGAGAAAACGCCATTGCATACTGTTACCCGGGAGCAACCGTCAATGTGGAGGCGCTGAGACAGAAAATTATGGAA AGCGGAGGGATCAATAAAATCAGTACCGGGTTTACTTATGGCTCATCAATCAATTCAGCAGGCACTACTAGAGCTTGCATCAGGAATGGAG GGAATAGTTTCTACGCGGAGTTGAAATGGCTCGTGAGTAAATCTAAGGGGCAGAATTTTCCCCAGACTACAAATACGTATAGAAACACCGAT ACTGCCGAGCACTTAATCATGTGGGGCATCCATCATCCAAGCAGTACCCAGGAGAAAAACGACCTCTATGGGACTCAGTCCTTGAGCATTA GCGTCGGATCTTCCACATACAGAAACAATTTCGTTCCAGTCGTCGGAGCTCGACCTCAGGTTAACGGGCAATCCGGGCGAATAGACTTTCA TTGGACTCTGGTGCAGCCCCGGGGACAACATAACCTTCAGTCACAATGGAGGGCTGATAGCCCCCAGTCGCGTGAGTAAATTGATTGGCAG GGGCCTAGGAATCCAATCTGACGCACCCATCGACAACAACTGCGAGAGCAAATGCTTTTGGCGCGGTGGGTCAATAAACACTCGCCTCCCT TTTCAGAATTTGTCACCTCGGACAGTGGGACAGTGCCCTAAGTATGTGAATCGCCGATCCCTGATGCTAGCCACCGGCATGCGCAATGTGC CCGAGCTGATTCAGGGAAGGGCCTGTTCGGCGCCATCGCCGGCTTCTTGGAAAATGGCTGGGAGGGGATGGTTGACGGCTGGTATGGC TTCCGGCATCAGAACGCACAGGGCACCGGCCAGGCGGCCGACTATAAGTCCACACAAGCAGCCATCGATCAAATCACGGGAAAGCTCAAT AGGCTGGTCGAGAAAACAAACACAGAATTCGAAAGTATCGAATCAGAGTTCTCCGAGATAGAGCACCAGATCGGAAATGTGATCAATTGGA CAAAGGACAGCATCACTGACATATGGACGTACCAGGCAGAGCTGCTTGTCGCCATGGAGAACCAGCATACGATTGATATGGCAGACTCTGA AATGCTGAACCTCTACGAGAGAGTACGGAAGCAACTACGTCAGAATGCAGAGGAGGATGGGAAAGGATGCTTTGAAATCTATCATGCATGT GACGATTCTTGCATGGAGTCAATACGCAACAATACGTACGACCACAGCCAGTATCGGGAGGAAGCGCTGCTTAACAGACTGAATATAAATC CGGTGACACTGTCATCTGGCTACAAGGACATCATCTTATGGTTTTCGTTTGGTGCGACTTGTTTTGTCTTACTAGCCGTCGTTATGGGTTTAT TCTTTTTTTTGTTTGAAGAATGGGAACATGAGGTGTACAATTTGCATT
3054	ATGTACAAGATCGTGGTCATAATCGCCCTTCTCGGTGCTGTTAAGGGACTAGATAAGATATGCCTGGGCCACCATGCCGTAGCTAACGGTA CAATAGTGAAGACATTAACTAATGAGCAAGAAGAGGTGACGAACGCTACTGAGACTGTTGAGAGCACAGGCATTAACCGGCTGTGTATGAA GGGCCGCAAGCACAAGGATCTGGGAAATTGCCACCCAATTGGCATGCTGATCGGAACACCAGCGTGCGATTTGCACCTAACCGGTATGTG GGATACTCTGATTGAGCGTGAAAATGCTATTGCCTACTGTTATCCCGGCGCTACAGTCAACGTGGAAGCGCTGCGCCAGAAGATCATGGAA TCTGGAGGAATTAATAAGATCTCAACTGGTTTCACCTACGGTAGCTCGATTAACAGCGCAGGAACAACTCGTGCCTGCATGAGAAATGGAG GCAATAGCTTTTATGCTGAACTGAAATGGCTCGTGTCTAAGTCAAAGGGCCAGAATTTCCCTCAAACAACTAACACCTACAGGAATACTGAC ACTGCGGAACATCTGATCATGTGGGGCATTCACCACCCCTCTTCAACACAGGAGAAAAACGATCTCTACGGCACTCAGTCGCTGTCCATCT CTGTGGGGAGCTCTACCTATCGTAACAATTTCGTCCCGGTTGTTGGCGCCCGGCCACAAGTGAACGGGCAGAGCGGCCGGATCGACTTCC ACTGGACCCTCGTACAGCCTGGGGACAACATCACCTTCTCACACAACGGAGGGCTAATCGCCCCCTCAAGGGTCAGCAAGTTGATAGGCA GGGGCTTGGGAATTCAGTCGGATGCCCCAATTGATAACAACTGCGAGAGCAAATGCTTTTGGCGCGGAGGATCCATCAATACAAGGCTCCC CTTCCAAAACCTCAGCCCCAGAACCGTTGGGCAGTGTCCTAAATATGTGAACAGGAGATCTCTGATGCTGGCCACTGGAATGAGAAATGTG CCTGAGCTGATCCAGGGCCGAGGACTGTTTGGAGCGATCGCCGGATTTCTTGAAAACGGCTGGGAAGGCATGGTGGATGGCTGGTACGGA TTTAGACATCAGAATGCCCAAGGGACCGGCCAAGCTGCAGACTACAAAAGTACCCAGGCCGCGATAGATCAGATTACGGGAAAGCTGAACC GTCTGGTGGAAAAAACTAATACGGAGTTTGAGTCCATAGAGAGTGAGTTTTCTGAAATTGAACACCAGATTGGGAACGTGATCAACTGGACC AAGGACTCAATCACTGATATATGGACCTACCAGGCCGAGCTCCTAGTGGCAATGGAAAACCAGCACACAATAGACATGGCCGACAGCGAAA TGTTGAATCTCTACGAAAGAGTGAGGAAGCAGCTTCGGCAGAATGCTGAGGAGGACGGAAAGGGATGTTTCGAGATCTACCACGCCTGTG ACCATAGTTGCATGGAAAGCATCCGTAACAATACCTACGATCACAGTCAATATACCGAAGAAGCATTGCTGAATAGGTTGAATATAAATCCC GTTACTCTTTCCTCGGGGTATAAGGATATTATCCTTTGGTTTTCATTTGGTGCCTCGTGTTTCGTCCTGCTGGCAGTGGTGATGGGCCTCTTT TTCTTCTGTTTGAAGAACGGGAATATGCGCTGTACCATCTGCATA
3055	ATGTACAAAGTAGTCGTAATCATCGCCCTTTTGGGAGCCGTCAAGGGCCTGGACAAGATATGCCTGGGACATCACGCTGTCGCCAACGGGA CGATTGTCAAAACACTGACAAACGAGCAGGAAGAGGTTACCAACGCGACTGAGACTGTTGAGTCAACAGGTATAAACCGACTGTGTATGAA AGGACCGAAACACAAAGATCTGGGGAATTGCCACCCCATCGGGATGCTGATCGGTACCCCTGCATGCGACCTGCACCTCACTGGTACTTG GGATACTCTCATAGAACGGGAGAACGCAATCGCATACTGCTATCCTGGAGCCACCGTTAATGTTGAGGCCTTGAGACAAAAAATTATGGAGA GTGGGGGAATCGACAAAATCTCCACCGGCTTTACCTATGGCAGCTCCATCAATTCAGCCGGTACCACGCGTGCCTGTATGCGCAACGGAG GTAATTCATTCTACGCCGAGCTAAAATGGCTGGTGTCCAAAAATAAAGGGCAAAACTTCCCGCAAACAACAAACACTTACAGAAACACAGAC ACTGCAGAACATCTGATAATGTGGGGAATCCACCATCCGTCATCTATACAGGAGAAAAACGATCTGTATGGCACTCAGTCACTCAGTATCTC CGTCGGCTCATCAACGTACAGGAATAATTTTGTGCCCGTCGTCGGCGCACGTCCACAAGTCAACGGTCAGTCCGGGAGAATCGACTTTCAT TGGACGCTTGTTCAACCGGGCGATAACATTACATTTTCACATAACGGAGGGCTCATCGCTCCAAGTAGAGTCTCTAAACTGATCGGACGTGG CCTGGGAATTCAGTCAGATGCGCCCATCGACAACAATTGCGAGTCGAAGTGCTTCTGGCGGGGGGGGTCCATAAATACCCGCCTCCCCTTT CAAAACCTCTCACCCCGTACCGTGGGCCAGTGCCCTAAATATGTGAATCGGCGATCCCTGATGCTGGCAACTGGAATGCGCAACGTACCAG AGCTGATTCAGGGCAGAGGACTCTTCGGGGCCATTGCAGGGTTCCTGGAAAATGGTTGGGAAGGAATGGTTGACGGATGGTATGGCTTCC GCCACCAGAATGCACAAGGGACAGGTCAGGCTGCAGATTATAAGTCCACGCAGGCCGCTATTGATCAGATTACGGGAAAACTCAACAGACT GGTGGAAAAAACCAACACTGAGTTTGAGAGCATTGAGTCCGAGTTTTCCGAGATTGAACATCAAATCGGTAATGTTATCAATTGGACTATGG ATAGTATTACGGATATTTGGACTTACCAGGCCGAATTGTTAGTGGCTATGGAAAACCAACACACTATTGACATGGCAGACTCAGAGATGCTC AATCTTTATGAGCGCGTGAGGAAGCAGCTCAGACAGAACGCGGAAGAAGACGGCAAGGGGTGCTTTGAGATTTACCACGCCTGCGACGAC TCCTGCATGGAAAGCATCCGCAACAACACCTACGACCACAGCCAATATCGCGAGGAGGCCCTCCTTAATCGGTTGAACATCAATCCCGTAA CACTGTCCTCCGGGTACAAAGACATCATCCTGTGGTTCAGTTTTGGCGCGAGCTGTTTCGTCTTGTTGGCTGTGGTGATGGGCCTCGTATTT TTTTGCCTTAAAAACGGGAACATGCGGTGTACAATCTGTATT

3056	ATGTACAAAATAATTGTGATCATTGCCCTCCTGGGGGCTGTGAAGGGCCTTGACAAGATCTGCCTAGGACACCATGCAGTTGCAAACGGAA CAATTGTGAAGACCCTAACCAACGAACAAGAGGAGGTGACAAACGCAACTGAGACCGTGGAGTCAACCGGCATAAATAGGCTTTGCATGAA GGGAAGGAAACACAAAGATCTCGGCAACTGTCATCCAATCGGCATGCTTATCGGGACGCCTGCATGTGATCTGCACCTAACTGGCACCTGG GACACACTGATTGAAAGGGAGAATGCTATCGCTTACTGTTACCCTGGTGCTACTGTGAACGTTGAGGCTTTGCGGCAGAAAATCATGGAGT CACGGGGCATTGACAAAATCTCGACAGGCTTTACCTACGGCTCAAGCATTAACTCCGCTGGGACCACTAGAGCATGTATGAGAAATGGAGG TAACTCATTCTATGCCGAGCTCAAATGGCTCGTGTCAAAAAGCAAGGGGCAGAACTTCCCGCAGACGACCAATACTTATCGGAATACAGACA CTGCCGAACATCTTATTATGTGGGGAATTCATCATCCATCCTCAACTCAGGAAAAAAACGACTTGTACGGGACTCAATCCCTCTCCATTTCCG TGGGCAGTAGCACATATAGGAATAATTTCGTTCCTGTCGTTGGGGCGAGGCCACAAGTGAATGGCCAGAGCGGCCGAATCGATTTTCATTG GACCCTTGTACAACCAGGGGACAATATCACTTTCTCACACAACGGAGGGTTGATTGCCCCATCCCGCGTTAGTAAGCTTATCGGCCGCGGC CTGGGTATACAGTCCGATGCACCCATCGATAATAATTGTGAGTCTAAGTGTTTTTGGCGAGGTGGCTCCATCAATACGCGGCTACCTTTTCA GAACCTGTCGCCTCGCACCGTAGGGCAGTGTCCAAAGTATGTCAATAGACGGTCTTTGATGTTGGCCACCGGAATGAGAAACGTTCCAGAG CTGATTCAGGGGCGCGGGCTGTTCGGCGCCATCGCCGGCTTCCTTGAGAATGGGTGGGAGGGCATGGTTGATGGATGGTACGGGTTTCG GCACCAGAACGCCCAGGGGACCGGCCAAGCCGCAGATTACAAGAGCACCCAGGCTGCCATCGACCAGATTACCGGCAAGCTCAATAGGC TCGTGGAAAAAACTAACACCGAGTTTGAGTCTATCGAATCGGAGTTTAGCGAGATAGAGCATCAGATTGGAAATGTCATAAATTGGACAAAA GACAGTATCACAGATATCTGGACATACCAGGCTGAGCTGCTGGTCGCTATGGAAAATCAGCACACTATTGATATGGCCGATTCAGAAATGTT GAACCTGTATGAGCGGGTGCGTAAGCAGCTCCGACAGAATGCTGAGGAAGATGGGAAGGGCTGCTTCGAGATATACCACGCTTGTGATGA CTCTTGCATGGAGAGCATCCGGAACAACACGTATGACCATAGTCAGTATCGGGAAGAGGCACTGCTCAACAGGCTGAATATCAATCCCGTG ACCCTCTCATCTCGATACAAGGACATCATCCTGTGGTTCAGCTTTGGGCGCATCTTGTTTCGTACTTCTGGCTGTCGTCATGGGCCTGTTTTT CTTCTGCTTAAAGAACGGAAATATGAGATGTACCATATGCATA
3057	ATGTATAAAGTGGTAGTCATCATCGCGTTACTGGGGGCAGTGCGAGGCCTTGATAAAATTTGTCTGGGACATCATGCCGTCGCCAACGGCA CCATCGTCAAGACACTAACTAACGAGCAGGAGGAGGTCACCAATGCCACCGAGACCGTCGAAAGTAAATCGTTGGGAAAGCTATGCATGAA GGGGAGGTCCTACAACGACCTGGGTAATTGCCATCCTATTGGGATCCTCATCGGTACCCCCGCATGTGACCTGCACCTGACAGGCACATG GGATACTCTGATTGAAAGAGAGAACGCTGTAGCGTACTGCTACCCCGGCGCAACTGTCAACGAAGAAGCGCTGAGACAGAAGATCATGGAA TCCGGGGGCATTTCAAAGATCTCTACCGGTTTCACGTATGGAACCAGTATCAACAGCGCAGGCACAACAAAGGCCTGCATGCGCAATGGCG GCAATTCCTTCTACGCCGAGCTTAAATGGCTCGTCAGTAAGAATAAAGGCCAGAATTTTCCCCAGACCACCAACACTTACCGGAACACTGAC ACGGCCGAGCATCTCATTATTTGGGGGATTCATCATCCAAGCTCGACTCAAGAAAAGAATGATCTGTATGGCACTCAGTCTCTGTCTATCTC AGTTGGCAGCTCCACGTACCAGAATAATTTCGTGCCTGTCGTTGGGGCTAGACCCCAAGTGAATGGACAATCAGGTAGGATAGATTTTCACT GGACCTTATTACAGCCTGGAGATAACATAACATTCAGTCACAATGGAGGATTAATAGCTCCATCCCGCGTGTCTAAACTGATCGGCAGAGGC TTGGGCATCCAGTCTGAAGCCCCTATTGACAATGGTTGCGAGAGCAAATGCTTTTGGAAAGGCGGAAGCATTAACACAAAGTTGCCTTTCCA GAACCTGTCTCCCCGCACTGTGGGCCAGTGTCCCAAGTACGTCAATAAACGATCTCTGATGCTGGCTACCGGTATGCGCAATGTTCCCGAA ATCATGCATGGGAGAGGGCTGTTCGGGGCTATAGCAGGATTTATCGAAAATGGATGGGAAGGGATGGTGGATGGATGGTACGGTTTCAGA CACCAAAATGCCCAGGGGACGGGGCAGGCTGCTGATTACAAAAGTACGCAGGCCGCCATTGATCAGATTACCGGCAAGCTTAATCGGCTG ATTGAAAAAACGAACACCGAGTTTGAGAGCATCGAGTCCGAGTTTAGCGAGATTGAGCATCAAATTGGCAACATCATCAACTGGACAAAAGA CAGCATCACTGATATCTGGACTTACCAAGCTGAACTGCTCGTGGCCATGGAGAATCAGCACACCATTGACATGGCAGATTCCGAGATGCTA AACCTGTACGAACGCGTCCGCAAGCAGCTCCGGCAGAACGCAGAGGAAGATGGAAAGGGGTGCTTCGAGATTTACCACGCTTGCGATGAC AGCTGTATGGAATCCATTCGAAACAATACATACGATCACTCACAGTATCGCGAGGAGGCTCTGCTCAACCGTTTGAATATTAATCCACTGAA GCTATCTAGCGGCTATAAGGACATAATCCTGTGGTTCAGCTTCGGCGCCTCTTGCCTAATCTTACTCGCCGTCGTGATGGGACTGGTGTTTT TTTGTCTCAAGAACGGCAACATGAGGTGCACCATATGCATC
3058	ATGTATAAGATCGTGGTCATAATCGCCCTGCTGGGCGCCGTCAAAGGGCTGGACAAAATATGCCTCGGACACCATGCTGTCGCCAATGGGA CGATTGTCAAAACGCTCACCAACGAACAGGAGGAAGTTACTAACGCCACAGAAACGGTAGAATCCACAGGGATCAATCGATTGTGTATGAA GGGTCGTAAGCACAAAGACTTGGGAAACTGTCACCCCATCGGCATGCTGATCGGCACGCCTGCCTGCGATCTACATCTGACAGGCATGTG GGACACACTAATAGAGAGGGAAAACGCAATCGCCTACTGTTACCCCGGTGCCACAGTAAACGTGGAAGCTCTGCGGCAGAAGATAATGGA GTCTGGAGGAATCAATAAAATCAGCACCGGTTTCACATATGGCTCCAGTATCAACTCCGCAGGCACTACCAGGGCTTGCATGCGCAATGGG GGAAATTCTTTCTACGCGGAGCTAAAATGGTTAGTCTCTAAGTCCAAAGGACAGAACTTTCCACAGACAACAAACACCTACAGGAATACAGA TACCGCTGAGCACCTGATCATGTGGGGTATTCATCATCCCAGTTCTACTCAGGAAAAGAACGATCTGTATGGGACACAGTCGCTGAGTATAT CTGTCGGTAGTAGTACATACCGTAATAATTTCGTGCCCGTTGTAGGCGCTAGACCTCAGGTGAATGGGCAATCCGGCCGGATTGATTTCCA CTGGACTCTCGTGCAGCCTGGGGATAATATCACGTTTAGCCACAATGGAGGTCTAATCGCCCCCTCCCGGGTTTCTAAACTGATTGGCCGG GGACTGGGAATCCAATCTGACGCCCCAATAGACAACAACTGCGAATCCAAGTGTTTCTGGCGGGGCGGCTCCATAAATACGCGACTTCCCT TCCAGAACTTAAGTCCCCGGACTGTGGGTCAGTGTCCAAAATACGTGAATCGCCGGAGTCTTATGCTCGCCACAGGCATGCGTAACGTCCC GGAACTTATACAGGGCCGGGGACTGTTTGGAGCTATCGCAGGATTCCTGGAGAACGGATGGGAGGGAATGGTGGACGGATGGTATGGTTT CCGGCACCAAAACGCTCAAGGAACGGGCCAGGCAGCAGATTATAAGTCAACACAGGCCGCGATCGACCAAATTACTGGCAAACTCAATAG GCTGGTCGAGAAAACCAACACTGAGTTTGAGAGCATTGAATCCGAATTCAGCGAAATCGAACACCAGATCGGCAACGTGATAAATTGGACG AAAGATTCCATTACCGACATTTGGACATACCAGGCCGAGTTGCTGGTCGCTATGGAAAATCAGCATACGATTGATATGGCCGATAGTGAGAT GCTTAACCTGTATGAACGCGTACGCAAACAGCTGCGCCAAAATGCTGAAGAGGACGCTAAAGGCTGCTTTGAGATTTATCACGCTTGCGAC GATTCATGCATGGAAAGTATCAGAAACAATACATATGACCACAGCCAGTACAGGGAGGAAGCGCTGCTCAATCGACTAAACATCAATCCTGT GACCCTGTCGAGTGGCTATAAAGACATAATTCTATGGTTTTCATTCGGAGCATCTTGCTTCGTCCTGCTCGCAGTTGTGATGGGACTCTTCTT CTTTTGTCTGAAAAATGGTAATATGCGGTGTACCATCTGCATT

3059	ATGTACAAGATCGTGGTGATCATTGCTCTGCTTGGAGCCGTCAAGGGCCTGGATAAGATTTGCCTGGGACATCACGCTGTCGCAAATGGCA CCATCGTCAAGACTTTAACAAACGAGCAGGAAGAGGTGACTAATGCCACTGAGACAGTAGAGAGCACTGGGATAAATAGGTTGTGCATGAA GGGGAGAAAGCACAAGGACCTCGGAAACTGCCACCCAATTGGTATGCTAATCGGCACTCCAGCTTGCGACTTGCATCTGACTGGCATGTG GGATACGCTGATAGAACGTGAGAATGCCATCGCCTACTGTTATCCTGGGGCTACTGTGAACGTCGAGGCTTTAAGACAAAAGATCATGGAA AGTGGCGCCATCAACAAGATAAGTACCGGGTTTACCTATGGGAGTAGTATTAATTCAGCAGCCACAACTCGGGCCTGCATGCGGAATGGGG GGAACAGCTTCTACGCCGAGCTAAAGTGGCTTGTGAGCAAGTCCAAAGGGCAGAATTTCCCGCAGACCACCAATACTTACCGCAACACCGA TACTGCAGAACACTTAATAATGTGGGGCATCCACCATCCTAGTAGCACGCAAGAGAAGAATGACTTGTACGGTACCCAGTCGCTAAGTATCA GCGTGGGCAGTTCGACATATAGAAATAACTTTGTCCCTGTTGTGGGCGCGCGCCCTCAGGTAAATGGACAGTCTGGTCGGATCGACTTTCA TTGGACCCTGGTGCAGCCAGGGGACAACATTACCTTTTCACATAATGGCGGTCTCATCGCACCCAGTCGGGTCTCAAAGCTGATCGGGCG GGGGCTGGGCATTCAGTCAGACGCTCCCATAGATAATAATTGCGAGTCCAAGTGTTTCTGGAGAGGGGGCTCCATCAATACTAGGCTGCCT TTTCAGAACCTCTCGCCCCGGACTGTGGGGCAGTGCCCCAAATACGTAAACCGAAGATCCCTAATGCTGGCTACTGGGATGCGGAACGTAC CTGAACTGATCCAGGGACGTGGGCTATTCGGAGCCATAGCTGGATTTCTGGAAAATGGCTGGGAGGGAATGGTTGATGGGTGGTACGGGT TTCGGCACCAGAATGCCCAGGGCACTGGGCAGGCTGCCGACTACAAATCAACTCAGGCCGCTATTGATCAAATTACAGGTAAGCTCAACCG TCTAGTTGAGAAAACAAATACTGAGTTCGAAAGCATTGAGTCTGAGTTTTCCGAGATTGAGCATCAGATCGGCAACGTTATTAATTGGACAAA GGACAGCATCACTGACATTTGGACATACCAGGCTGAACTTCTGGTTGCAATGGAAAATCAGCACACTATCGACATGGCTGATTCCGAGATGC TAAATCTCTACGAACGGGTCAGAAAGCAACTGCGGCAGAATGCCGAGGAAGATGGAAAAGGCTGTTTCGAAATTTACCATGCTTGCGACGA CTCCTGTATGGAGTCGATCAGGAACAACACCTACGATCACAGTCAGTATCGTGAGGAGGCACTTTTGAACCGTCTGAACATCAACCCCGTG ACCCTTTCCTCCGCGTATAAAGATATCATCCTCTGGTTCTCCTTCGGAGCATCTTGCTTCGTTCTCCTGGCCGTAGTGATGGGCCTGTTTTTT TTCTGCCTGAAGAATGGGAACATGAGATGCACCATCTGCATT
3060	ATGTATAAAATCGTTGTTATTATCGCCTTATTGGGCGCTGTGAAAGGGCTGGATAAAATCTGCTTAGGGCATCATGCCGTTGCTAATGGAAC GATTGTCAAAACCCTGACAAATGAGCAGGAAGAGGTTACAAACGCTACTGAGACAGTGGAAAGCACCGGGATAAACCGGTTATGTATGAAA GGTCGGAAGCACAAAGACCTAGGAAACTGCCACCCAATCGGCATGCTGATTGGTACCCCAGCATGTGATCTGCACCTAACAGGAACCTGG GACACACTCATCGAGCGGGAAAACGCCATTGCCTATTGTTATCCTGGCGCCACGCCAAACGTGGAGGCGCTGCGCCAGAAGATTATGGAG TCCGGAGGTATCGACAAAATCTCTACCGGCTTTACGTACGGGAGTTCAATTAATTCAGCCGGAACAACCAGGGCCTGTATGCGCAATGGGG GGAATTCATTTTATGCCGAATTGAAGTGGCTCGTCTCCAAGAGCAAAGGACAGAACTTTCCTCAGACAACGAATACTTACAGAAACACCGAT ACCGCTGAGCACTTAATTATGTGGGGAATCCACCATCCTAGTTCAATCCAGGAAAAAAACGATCTGTACGGGACACAGAGTCTCTCTATATC TGTTGGATCTTCCACCTATAGGAACAATTTTGTTCCCGTGGTGGGCGCCAGGCCACAGGTGAATGGCCAGTCCGGGAGAATAGATTTTCAC TGGACCCTTGTCCAGCCTGGGGATAACATTACTTTCAGCCACAACGGAGGCCTCATCGCGCCAAGTAGGGTTTCCAAATTGATAGGGAGAG GCCTGGGGATTCAGTCCGACGCCCCCATAGATAATAATTGCGAATCAAAGTGTTTTTGGAGGGGGGGGTCCATCAACACAAGACTCCCCTT TCAGAATCTGTCACCCCGAACCGTTGGCCAGTGTCCGAAATATGTGAACCGGAGGTCCTTAATGCTGGCCACTGGAATGCGGAATGTGCCA GAGCTTATCCAAGGAAGGGGACTGTTTGGAGCAATCGCCGGCTTCCTAGAAAATGGCTGGGAGGGTATGGTGGACGGCTGGTATGGGTTT AGACATCAGAATGCCCAAGGAACCGGTCAGGCTGCGGACTATAAGTCCACCCAGGCCGCTATCGACCAGATCACAGGTAAGCTCAACAGG CTGGTCGAGAAAACAAATACAGAGTTCGAGTCCATAGAGTCCGAGTTCAGCGAAATAGAACACCAGATTGGAAATGTGATTAACTGGACGAT GGATTCCATCACAGACATCTGGACCTACCAGGCAGAGTTACTGGTAGCTATGGAAAATCAGCATACAATCGACATGGCCGACTCTGAAATG CTAAATCTTTACGAGAGAGTTCGTAAACAGCTTCGCCAGAACGCCGAGGAAGATGGGAAAGGGTGCTTTGAAATATATCACGCCTGTGATG ATAGCTGTATGGAATCAATTAGAAATAACACATATGACCATTCTCAGTACAGAGAAGACGCTCTGCTCAATAGGCTCAATATCAATCCAGTGA CCCTATCCTCAGGATACAAGGACATCATACTCTGGTTCAGTTTTGGCGCCTCTTGTTTCGTTCTGTTGGCCGTCGTCATGGGCCTCGTGTTT TTTTGTTTAAAGAACGGCAATATGCGATGCACTATTTGCATC
3061	ATGTATAAGATTGTAGTGATCATTGCTCTGCTCGGGGCCGTGAAGGGCCTGGATAAGATCTGCCTGGGTCACCATGCCGTCGCAAACGGCA CGATTGTGAAAACACTGACGAATGAGCAGGAGGAGGTTACTAACGCTACAGAGACGGTCGAGTCAACTGGAATTAACCGGCTTTGCATGAA AGGACCAAAACACAAAGATCTTGGGAACTGCCACCCAATTGGGATGCTGATTGGAACCCCAGCCTGTGATCTGCACTTGACAGGCATGTGG GACACTCTGATCGAGAGGGAGAACGCAATAGCCTATTGCTACCCTGGCGCAACCGTGAACGTGGAGGCCTTGCGCCAGAAGATAATGGAG TCTGGGGGTATCAACAAGATCAGTACAGGTTTCACTTACGGCAGTTCTATTAACTCTGCCGGAACCACAAGAGCATGCATGCGTAACGGAG GCAATTCCTTCTATGCCGAACTTAAATGGCTTGTCTCCAAGTCTAAGGGGCAGAACTTCCCCCAGACAACGAACACTTACCGCAATACGGAC ACAGCAGAACACTTAATCATGTGGGGGATCCATCATCCCAGCAGTACACAGGAGAAAAATGACTTATACGGAACTCAGAGCTTAAGTATCTC CGTGGGCAGTAGTACTTACAGAAACAATTTCGTCCCTGTCGTAGGCGCTAGGCCTCAGGTGAATGGCCAGTCTGGACGCATCGACTTCCAC TGGACTCTGGTTCAACCAGGGGACAACATCACTTTCTCGCACAACGGAGGGTTGATCGCACCCAGTCGGGTGAGTAAACTTATCGGGAGAG GCCTCGGAATACAATCCGATGCTCCCATCGATAATAATTGCGAGAGCAAATGTTTTTGGCGCGGAGGGAGCATTAATACACGCCTGCCTTTT CAGAACTTGTCACCACGAACCGTGGGCCAATGCCCCAAGTACGTCAACAGAAGATCATTGATGCTGGCTACTGGCATGAGGAATGTGCCTG AATTAATTCAAGGAAGAGGACTGTTCGGCGCCATCGCAGGTTTTCTTGAGAATGGGTGGGAGGGGATGGTGGATGGTTGGTATGGCTTTCG CCACCAGAATGCGCAGGGCACAGGGCAGGCTGCTGATTACAAGTCAACCCAGGCAGCCATCGACCAGATTACTGGCAAGCTGAATAGGCT CGTTGAAAAAACCAACACAGAGTTCGAGTCAATTGAGAGCGAGTTTTCTGAAATAGAGCATCAGATAGGCAACGTGATAAATTGGACCAAAG ACAGTATCACAGACATTTGGACTTATCAAGCAGAGTTGTTGGTAGCCATGGAGAATCAACACACGATCGACATGGCCGATAGCGAAATGCTA AACCTGTATGAGCGCGTTCGGAAGCAGCTGCGCCAGAACGCAGAGGAGGACGGGAAGGGCTGTTTCGAAATTTATCACGCTTGTGACGAC AGTTGCATGGAATCCATCCGCAACAATACGTATGATCACTCCCAGTATAGAGAAGAGGCCCTCTTGAATAGACTGAACATCAACCCAGTGAC TCTGAGCTCAGGTTACAAGGACATTATTCTCTGGTTTTCTTTCGGAGCTTCCTGCTTTGTCCTCCTCGCTGTCGTAATGGGACTCTTTTTCTTT TGCCTGAAGAACGGGAATATGAGATGCACCATTTGCATC

3062	ATGTATAAGATTGTCGTTATCATTGCACTGCTTGGTGCAGTTAAAGGGCTGGATAAGATTTGTCTAGGGCACCATGCCGTGGCCAACGGGAC CATAGTGAAAACACTGACCAATGAACAGGAGGAAGTCACCAATGCTACTGAAACTGTTGAATCCACCGGCATTAACCGGCTGTGTATGAAG GGCCGCAAACACAAGGACCTGGGCAATTGTCACCCCATCGGCATGTTGATTGGCACCCCTGCATGTGACTTGCACCTCACTGGCATGTGG GACACACTGATCGAACGAGAGAATGCAATTGCCTACTGCTACCCTGGTGCCACAGTGAACGTGGAGGCTCTTCGACAAAAGATCATGGAAA GCGGGGGCATCAATAAGATTTCTACAGGGTTCACATACGGTTCATCTATTAACAGTCCAGGAACGACCCGGGCCTGCATGCGGAACGGCG GCAATTCATTCTACGCCGAATTGAAGTGGTTGGTGTCGAAGAGTAAGGGCCAGAATTTCCCGCAGACAACAAACACATATCGCAATACGGAT ACGGCCGAACACTTAATCATGTGGGGAATCCATCACCCTTCTTCTACCCAAGAAAAGAACGATCTATACGGCACCCAGTCCCTGTCAATTTC AGTGGGTTCTTCGACATACAGGAACAATTTCGTCCCTGTTGTCGGCGCACGGCCCCAGGTAAATGGGCAGAGTGGGAGAATCGACTTCCAC TGGACTCTGGTGCAGCCTGGGGATAACATTACGTTCAGTCACAATGGCGGCCTCATAGCCCCATCACGGGTGAGCAAACTAATAGGACGAG GCCTGGGAATACAGTCCGACGCCCCAATCGATAACAACTGCGAGTCAAAATGTTTCTGGCGAGGAGGCAGCATCAATACAAGACTCCCTTT TCAGAATCTTAGTCCAAGGACAGTCGGACAGTGCCCAAAATACGTGAATAGACGCAGCTTGATGTTGGCAACGGGGATGCGCAATGTTCCT GAGCTCATACAGGGTAGAGGCCTATTTGGGGCCATCGCAGGCTTCCTGGAAAACGGATGGGAGGGCATGGTGGACGGATGGTACGGATTT CGTCACCAGAATGCTCAAGGCACAGGTCAGGCTGCCGACTACAAATCTACCCAGGCCGCCATCGACCAGATTACCGGAAAACTGAACAGG TTGGTGGAAAAAACGAATACAGAATTCGAGAGTATTGAGAGTGAGTTCTCCGAGATCGAGCACCAAATCGGCAATGTGATTAATTGGACAAA AGACTCCATTACCGACATCTGGACTTACCAGGCTGAGTTGCTCGTGGCGATGGAAAACCAGCATACCATTGATATGGCCGATAGTGAAATG CTGAATCTGTACGAACGCGTCAGAAAACAGCTACGACAGAACGCGGAGGAGGATGGGAAGGGCTGCTTCGAAATCTACCACGCTTGCGAT GATTCCTGCATGGAGTCCATTCGGAATAACACGTACGACCATAGCCAATACAGGGAGGAGGCGCTCCTAAATAGGTTGAACATAAATCCAG TTACCCTTTCTTCCGGTTACAAGGATATTATCCTGTGGTTCTCCTTTGGCGCCTCTTGTTTCGTCCTTCTCGCCGTGGTGATGCGATTATTTT TCTTCTGCCTCAAGAATGGAAATATGCGGTGTACGATCTGTATT
3063	ATGTACAAAGTGGTGGTGATAATTGCTCTCCTCGGCGCCGTGAAAGGCCTCGACAAGATATGTCTGGGCCATCATGCAGTCGCCAACGGGA CCATCGTCAAGACCCTCACCAATGAACAGGAGGAGGTTACCAACGCCACTGAAACAGTTGAGTCTACTGGAATCAATCGACTGTGTATGAA GGGGAGAAAACACAAGGACCTCGGTAACTGTCATCCTATTGGGATGCTTATTGGAACACCAGCCTGTGATTTACATCTGACGGGCACCTGG GACACCCTGATCGAAAGAGAGAACGCAATTGCATATTGCTACCCTGGCGCTACAGTTAATGTTGAAGCTCTGAGGCAAAAAATCATGGAGTC CGGGGGAATCGACAAGATTTCAACCGGCTTCACGTACGGCAGTTCTATCAACAGCGCAGGTACCACAAGGGCCTGTATGAGGAACGGCGG AAATAGCTTTTACGCCGAACTGAAATGGCTGGTGTCAAAGAACAAAGGACAGAACTTCCCTCAGACCACAAACACATACCGGAATACAGACA CCGCAGAGCACCTTATAATGTGGGGCATCCATCATCCCTCCTCTATTCAGGAGAAGAATGATCTGTACGGTACACAGTCACTCTCCATCTCT GTAGGCAGCTCCACATATCGTAATAACTTTGTCCCAGTTGTGGGAGCCCGACCACAGGTGAACGGCCAATCAGGACGAATTGATTTTCACT GGACCCTCGTGCAGCCTGGCGACAACATAACATTCAGCCATAATGGAGGACTCATCGCCCCGTCCCGCGTGTCTAAATTAATCGGTCGGG GGCTGGGGATCCAGTCCGATGCGCCCATAGATAACAACTGCGAGTCTAAATGCTTCTGGCGGGGCGGGTCCATTAACACTCGACTGCCCT TCCAAAATCTCAGCCCACGCACAGTAGGCCAGTGTCCAAAATATGTCAATAGACGTTCTCTAATGCTCGCCACCGGAATGAGAAATGTGCCT GAGCTGATTCAGGGAAGGGGATTGTTTGGAGCTATAGCCGGATTCTTGGAGAATGGTTGGGAAGGCATGGTCGACGGATGGTATGGATTC AGACACCAGAACGCTCAAGGCACGGGCCAAGCAGCCGACTACAAATCCACCCAAGCAGCCATTGATCAGATTACAGGGAAGCTGAACAGG CTGGTGGAGAAGACAAACACAGAATTCGAAAGTATTGAGAGCGAATTTTCCGAGATCGAGCACCAGATCGGTAATGTGATCAATTGGACCAT GGACAGCATAACGGACATCTGGACCTATCAGGCCGAACTCCTGGTCGCAATGGAAAATCAGCATACTATCGACATGGCCGATAGTGAGATG TTGAACCTGTATGAGAGGGTTCGTAAGCAGCTCCGTCAGAACGCAGAGGAGGATGGTAAAGGATGTTTTGAGATCTATCACGCATGCGATG ACAGTTGCATGGAATCTATTCGTAACAATACGTACGATCATAGCCAATACCGCGAAGAAGCATTGCTGAACCGGCTGAATATTAACCCCGTG ACATTATCCTCCGGGTACAAGGACATAATTCTGTGGTTCTCCTTCGGGGCCAGCTGCTTCGTGCTGCTCGCTGTGGTCATGGGCCTGGTCT TCTTCTGCCTGAAAAATGGCAACATGCGATGCACAATATGCATC
3064	ATGTACAAAATCGTCGTGATCATAGCACTGCTGGGAGCCGTAAAGGGTCTCGATAAGATCTGCCTAGGGCACCACGCAGTAGCCAATGGAA CCATAGTAAAAACACTCACGAATGAGCAGGAGGAAGTTACCAACGCCACTGAAACCGTCGAATCTACCGGGATCAACAGATTGTGCATGAA AGGGCGCAAACACAAGGATCTGGGGAACTGTCACCCAATCGGAATGCTAATTGGAACACCAGCCTGTGATCTCCATTTGACAGGAATGTGG GATACACTGATAGAGCGTGAAAACGCCATAGCGTACTGTTACCCCGGCGCCACAGTGAACGTGGAAGCGTTACGACAAAAGATAATGGAAT CAGGAGGCATTAATAAGATAAGTACTGGCTTTACCTACGGGTCTTCAATCAACTCTGCCGGGACAACTAGAGCCTGTATGAGAAACGGTGGT AATTCGTTCTACGCTGAGCTGAAGTGGCTGGTGTCCAAGAGCAAAGGGCAGAATTTCCCACAGACTACAAACACCTACAGAAACACCGATA CAGCCGAACACCTCATCATGTGGGGAATTCATCACCCAAGCAGCACGCAGGAGAAGAACGATTTATATGGGACACAAAGCTTGTCAATTTC CGTGGGCAGTAGCACATACAGAAATAACTTCGTGCCAGTGGTAGGGGCGGGTCCCCAGGTGAATGGACAGTCCGGCCGCATTGATTTCCA TTGGACCCTCGTGCAGCCGGGAGACAACATCACATTCAGCCACAATGGTGGACTAATCGCCCCATCCCGCGTCTCCAAACTGATCGGGAG AGGACTGGGTATTCAGTCTGACGCGCCCATTGATAATAACTGCGAGTCAAAATGTTTCTGGCGAGGGGGGAGCATCAATACACGATTGCCT TTTCAGAACTTGTCCCCTCGGACTGTCGGCCAGTGCCCTAAGTATGTGAATAGGAGATCCCTGATGCTTGCCACCGGAATGAGAAACGTTC CTGAGCTGATTCAGGGTCGCGGTCTATTTGGCGCAATCGCCGGCTTTCTCGAAAACGGCTGGGAGGGCATGGTAGATGGTTGGTACGGTT TCCGGCATCAGAACGCCCAGGGAACTGGACAAGCCGCTGATTACAAGTCAACACAGGCAGCAATCGACCAAATTACTGGCAAACTGAACCG GCTTGTCGAGAAAACCAATACTGAATTTGAGTCAATTGAATCCGAATTTAGTGAAATTGAACACCAGATCGGCAATGTGATTAATTGGACTAA GGACTCTATTACCGATATTTGGACTTATCAGGCTGAGCTACTTGTGGCCATGGAGAACCAGCACACAATCGACATGGCTGATAGTGAGATGC TGAACCTTTACGAGCGAGTCAGGAAACAGCTGAGACAGAACGCCGAGGAGGATGGAAAAGGCTGTTTCGAAATCTATCATGCTTGCGACGA TTCATGCATGGAGTCAATCAGAAATAACACCTACGACCACAGTCAGTATCGCGAGGAAGCTTTACTCAACAGACTCAATATCAATCCTGTAAC ATTGTCTTCCGGCTACAAAGACATCATCCTGTGGTTCTCATTTGGAGCCAGCTGCTTTGTCTTGTTGGCTGTGGTGATGGGACTGTTCTTTTT TTGCCTCAAAAATGGCAATATGAGATGTACCATTTGTATC

3065	ATGTATAAGATCGTGGTGATCATTGCTCTCCTGGGCGCAGTAAAGGGTCTTGACAAGATATGTCTTGGACACCATGCAGTGGCAAACGGCA CAATTGTTAAAACACTCACCAACGAGCAGGAGGAAGTGACCAATGCCACTGAAACAGTAGAGTCCACGGGGATCAATAGGCTCTGCATGAA AGGGAGGAAGCACAAAGATCTGGGTAATTGCCATCCAATCGGAATGCTCATAGGAACACCAGCCTGCGACCTTCATCTGACCGGTATGTGG GATACATTGATCGAACGCGAGAACGCGATAGCTTACTGCTATCCCGGCGCCACAGTGAATGTGGAGGCCCTCCGACAGAAGATTATGGAGT CGGGCGGTATTAACAAAATATCCACTGGTTTTACTTACGGATCTTCGATCAATTCCGCAGGAACAACACGAGCCTGTATGAGGAATGGCGG GAACTCTTTCTATGCTGAGCTTAAGTGGCTCGTTTCTAAGTCTAAGGGCCAGAATTTCCCTCAGACGACTAATACCTACAGGAACACGGATA CAGCAGAGCACCTGATTATGTGGGGTATCCATCATCCATCTAGCACCCAAGAGAAAAATGATCTGTATGGGACCCAGTCACTTTCCATTAGC GTGGGCTCTAGTACATACAGGAATAACTTTGTACCTGTGGTGGGGGCTAGACCGCAAGTGAATGGCCAGAGCGGCCGGATTGACTTCCACT GGACTTTAGTCCAGCCTGGAGATAACATTACATTTTCACACAACGGCGGGCTGATTGCGCCTAGTAGGGTGAGCAAACTTATTGGACGAGG ATTAGGAATCCAGAGCGATGCCCCCATCGACAACAACTGCGAAAGCAAATGCTTTTGGAGGGGCGGTTCAATCAACACCCGTCTCCCTTTC CAGAATCTGTCCCCCCGTACCGTGGGGCAGTGTCCAAAGTATGTTAATAGAAGGTCACTGATGCTAGCCACTGGCATGCGGAATGTGCCCG AGCTGATCCAGGGCCGTGGGCTGTTCGGTGCTATTGCAGGATTTCTGGAAAATGGCTGGGAAGGCATGGTAGATGGTTGGTACGGTTTTC GGCACCAGAACGCCCAGGGGACCGGACAGGCGGCCGACTACAAATCCACTCAAGCCGCTATCGATCAGATTACGGGAAAGCTAAATAGAC TGGTCGAAAAAACCAACACAGAGTTCGAATCCATAGAGTCCGAGTTCTCTGAGATCGAGCACCAGATCGGGAATGTGATCAACTGGACTAA GGATAGCATTACTGATATTTGGACCTACCAGGCAGAGCTGCTGGTCGCCATGGAAAACCAACACACAATCGATATGGCGGATAGTGAGATG CTGAACCTGTATGAAAGAGTGAGGAAACAGCTGAGACAGAACGCAGAGGAGGACGGAAAAGGGTGCTTCGAGATTTACCACGCTTGCGAT GACAGTTGCATGGAAAGCATACGCAATAACACTTACGACCATAGTCAATACCGGGAGGAGGCTCTCCTGAATAGGCTCAACATCAATCCTGT GACCCTGAGTTCAGGTTACAAAGATATCATCTTGTGGTTTTCATTTGGGGCGAGCTGCTTTGTGCTGCTGGCCGTCGTGATGGGCCTGTTCT TTTTTTGCCTGAAGAATGGCAATATGCGATGTACCATATGCATC
3066	ATGTATAAAATAGTGGTGATCATAGCACTCTTGGGAGCCGTAAAGGGACTGGATAAGATCTGCCTGGGGCATCATGCCGTTGCCAACGGCA CAATCGTGAAGACCCTTACTAATGAACAAGAAGAAGTGACCAACGCTACTGAGACCGTTGAATCCACCGGCATTAACCGCCTCTGCATGAA GGGGCGGAAGCACAAAGATCTCGGCAACTGCCACCCAATCGGGATGCTGATTGGGACCCCCGCGTGCGATTTACATCTGACAGGTATGTG GGATACATTAATTGAAAGGGAGAACGCGATTGCGTACTGCTACCCTGGGGCTACCGTGAATGTCGAAGCACTCCGACAGAAGATCATGGAA TCTGGGGGGATCAACAAGATCTCGACCGGTTTTACCTATGGGAGTTCTATTAACTCTGCTGGCACAACACGGGCATGTATGCGCAATGGGG GGAATAGTTTCTATGCTGAGCTCAAATGGCTTGTATCCAAATCCAAAGGGCAGAATTTCCCCCAGACTACAAATACCTACAGAAACACGGAT ACCGCTGAGCACCTAATAATGTGGGGAATTCACCATCCTTCCTCAACCCAAGAGAAGAATGATCTTTACGGGACCCAGAGTCTCAGCATTAG CGTCGGATCGTCGACTTACCGCAACAATTTCGTCCCAGTGGTCGGAGCCAGGCCACAGGTCAACGGCCAGTCTGGCAGGATAGACTTCCA CTGGACCCTGGTTCAGCCTGGAGATAACATTACATTTAGTCATAACGGCGGCCTGATTGCTCCCAGTCGCGTGTCTAAACTAATCGGACGT GGTCTGGGCATCCAATCCGATGCGCCGATCGACAATAACTGCGAAAGCAAGTGCTTTTGGAGGGGAGGATCCATCAATACACGACTTCCCT TCCAGAACCTATCACCAAGAACCGTAGGCCAGTGCCCGAAATACGTGAATCGACGGAGTCTGATGCTGGCCACCGGCATGAGAAATGTGC CTGAGCTGATTCAGGGACGAGGACTGTTTGGCGCAATCGCTGGCTTCCTGGAAAACGGGTGGGAAGGAATGGTCGATGGCTGGTACGGAT TTCGGCATCAGAACGCCCAAGGCACTGGCCAAGCCGCCGACTACAAGAGTACACAGGCCGCAATCGACCAGATCACAGGTAAGTTGAATC GATTGGTGGAGAAGACCAACACTGAGTTCGAATCTATTGAGAGCGAATTCAGTGAAATCGAGCATCAGATCGGAAATGTGATAAATTGGACC AAGGACTCAATTACCGACATCTGGACATACCAAGCAGAGCTTCTGGTTGCGATGGAAAACCAACACACAATTGATATGGCAGATTCCGAGAT GCTGAACCTCTATGAGCGCGTCAGGAAGCAATTGAGGCAGAATGCAGAGGAAGATGGAAAAGGGTGTTTCGAAATCTATCACGCCTGCGAC GACAGTTGCATGGAATCGATCCGCAACAATACTTATGACCACTCACAGTATAGAGAGGAGGCTTTGCTGAACCGCTTAAATATCAATCCTCT CACGTTGAGTAGCGGCTACAAGGATATCATCCTATGGTTTTCCTTCGGGGCGTCTTGTTTTGTGCTGTTAGCAGTGGTGATGGGACTTTTCT TCTTCTGTCTTAAAAACGGAAACATGAGATGTACCATCTGCATA
3067	ATGTATAAGATCGTGGTTATCATTGCCCTCCTGGGCGCAGTCAAAGGCTTGGACAAAATCTGTCTCGGACATCACGCCGTTGCTAACGGCA CCATTGTTAAGACCCTTACCAACGAGCAGGAAGAAGTGACAAATGCAACAGAAACGGTCGAGTCCACAGGCATCAACCGGCTCTGTATGAA AGGTCGGAAGCACAAGGATCTTGGGAATTGTCACCCAATAGGTATGCTGATTGGCACTCCTGCCTGCGATCTTCATCTAACTGGGATGTGG GACACCCTTATCGAACGAGAAAATGCCATTGCATACTGTTACCCTGGTGCTACTGTCAACGTGGAAGCCCTCAGGCAGAAAATAATGGAGTC GGGTGGGATAAATAAGATAAGCACCGGGTTCACTTACGGCAGCTCCATTAATTCCGCCGGAACTACGCGCGCCTGTATGAGAAATGGCGGT AACTCCTTCTACGCGGAGCTGAAGTGGCTGGTCTCGAAGTCTAAAGGCCAGAACTTCCCCCAAACAACAAACACCTACAGGAACACTGACA CCGCTGAACACCTCATCATGTGGGGCATTCACCACCCCTCCAGCACCCAGGAAAAAAACGACTTATACGGCACACAGAGCCTTAGTATAAG CGTGGGAAGCTCAACTTATCGCAACAATTTTGTTCCAGTGGTGGGCGCAAGACCGCAGGTGAACGGGCAAAGTGGTCGAATTGACTTTCAC TGGACACTCGTGCAACCAGGCGACAATATCACCTTTTCACATAACGGAGGACTCATTGCCCCGTCCCGCGTCAGTAAGTTAATTGGGCGGG GCCTCGGCATTCAGTCAGACGCCCCTATCGATAACAACTGTGAATCTAAGTGCTTTTGGAGAGGAGGGTCTATAAACACGAGACTCCCCTTC CAGAACTTATCGCCTCGCACTGTGGGTCAGTGTCCTAAGTACGTTAACAGACGTTCCCTGATGCTGGCAACGGGAATGAGAAACGTTCCAG AACTGATCCAGGGGCGCGGGTTATTTGGGGCTATCGCTGGTTTCCTGGAGAATGGCTGGGAGGGAATGGTTGACGGGTGGTATGGGTTTC GTCACCAAAACGCCCAGGGGACGGGGCAGGCCGCAGATTACAAGTCTACTCAGGCGGCCATAGACCAGATTACAGGCAAGCTGAATAGAC TGGTCGAGAAGACGAACACGGAGTTTGAGTCTATTGAGTCCGAGTTCAGCGAGATTGAACATCAGATAGGTAATGTTATCAATTGGACAAAG GATAGCATAACCGATATTTGGACATACCAGGCAGAGTTACTGGTTGCAATGGAAAATCAACACACTATTGACATGGCTGACTCAGAGATGCT CAACTTATACGAACGAGTGAGGAAGCAACTACGACAGAACGCCCGAAGAGGACGGAAAGGGTTGCTTTGAGATCTACCACGCCTGCGACGA TAGCTGCATGGAGAGCATCAGGAATAATACTTACGATCACTCACAGTACAGGGAGGAAGCCCTCTTGAACCGTCTGAACATAAATCCTGTCA CACTTAGCTCCGGCTACAAAGATATCATCTTGTGGTTTTCATTTGGCGCGTCATGTTTTGTGCTCCTAGCTGTGGTCATGGGCTTATTCTTTT TCTGCCTGAAAAATGGCAATATGAGGTGCACAATTTGTATC

3068	ATGTACAAGATAGTGGTAATTATCGCGCTGTTGGGCGCAGTGAAAGGACTGGACAAGATCTGCCTGGGTCACCACGCCGTGGCCAATGGA ACGATTGTGAAAACCTTGACAAACGAACAGGAGGAAGTGACCAACGCAACGGAGACTGTGGAATCAACTGGTATCAATCGGCTATGCATGA AAGGCAGAAAACACAAAGATTTGGGCAATTGCCATCCCATCGGTATGCTCATTGGAACACCCGCTTGCGACCTCCACCTGACTGGGACGTG GGATACTTTGATCGAGAGGGAGAATGCTATTGCCTACTGTTACCCTGGCGCAACAGCTAATGTGGAGGCCCTCCGGCAGAAGATCATGGAG AGCGGTGCAATCGATAACATTTCAACTGGCTTCACATACGGCTCCTCGATCAATTCTGCCGGCACTACTCGGGCTTGCATGAGAAACGGAG GAAATAGTTTCTATGCTGAGCTGAAGTGGCTTGTGAGTAAATCTAAAGGCCAGAACTTTCCCCAGACAACCAACACATACCGGAATACCGAC ACCGCAGAACATCTGATTATGTGGGGGATACATCACCCATCTAGCATTCAGGAGAAAAACGACCTGTATGGCACTCAGTCCCTGAGCATTTC CGTAGGCTCTAGTACCTACCGGAACAATTTCGTACCCGTCGTAGGCGCCCGCCCCCAAGTGAACGGCCAGTCCGGTCGAATTGACTTCCA CTGGACACTAGTGCAGCCTGGGGACAATATCACATTTAGTCATAACGGCGGACTGATCGCCCCATCGAGAGTGTCTAAACTGATCGGACGC GGGCTGGGTATTCAATCAGATGCCCCCATCGACAACAACTGCGAAAGCAAATGCTTCTGGCGCGGCGGGAGTATAAACACCCGGCTTCCAT TTCAGAACCTCTCTCCTCGGACCGTCGGACAGTGCCCGAAATATGTGAACCGCAGGTCACTGATGTTGGCAACTGGAATGAGGAATGTCCC CGAATTAATACAGGGCCGGGGCTTATTCGGAGCCATCGCAGGCTTCCTGGAGAACGGTTGGGAAGGCATGGTGGATGGATGGTATGGATT TAGACATCAAAATGCACAGGGAACAGGCCAGGCTGCCGACTACAAATCTACCCAGGCCGCGATAGATCAAATTACAGGAAAGTTGAACAGG CTGGTGGAAAAGACTAATACCGAATTTGAATCAATCGAGTCCGAATTCTCCGAAATCGAACACCAGATCGGGAATGTCATTAACTGGACTAT GGATTCTATAACCGATATCTGGACATACCAGGCAGAACTGCTAGTAGCCATGGAAAATCAACACACGATCGACATGGCTGACTCAGAGATG CTGAACCTCTACGAGCGGGTCAGAAAACAGCTCCGACAGAATGCAGAGGAAGACGGAAAAGGCTGTTTTGAGATTTACCACGCTTGCGACG ACTCTTGTATGGAGTCCATACGGAATAATACGTACGATCACTCACAGTATAGGGAGGAGGCCCTACTGAATAGACTGAACATCAACCCCGTC ACCCTGTCTAGTGGCTATAAAGACATCATTTTGTGGTTTTCTTTCGGAGCAAGTTGCTTCGTCCTCCTCGCCCGTGGTGATGGGACTCGTGTT CTTCTGCTTGAAAAATGGGAATATGCGATGTACCATTTGCATC
3069	ATGTATAAAATTGTCGTGATAATCGCTCTGCTTGGCGCTGTGAAGGGCTTAGATAAGATATGCCTCGGACACCATGCAGTTGCCAATGGCAC GATAGTGAAAACATTGACCAACGAGAAAGAAGAGGTGACCAACGCTACCGAGACTGTGGAATCCACCGGTCTGAATAGACTCTGCATGAAG GGGCGGAAGCATAAGGACTTGGGCAATTGTCACCCAATTGGGATGCTCATCGGATCGCCCGCATGTGACCTGCATCTCACTGGGACCTGG GACACATTGATTGAACGAGAGAACGCTATTGCGTACTGCTACCCGGGAGCCACTGTTAATGGTGAAGCACTTCGCCAGAAGATCATGGAGT CCGGAGGCATCGACAAGATCAGCACTGGATTCACCTATGAAAGTTCCATAAACTCCGCAGGCACTACAAGAGCTTGCATGAGGAACGGCGG GAATTCATTCTACGCTGAGCTCAAGTGGCTGGTGAGCAAATCTAAGGGACAGAATTTTCCACAGACAACCAACACGTATCGAAACACCGATA CTGCAGAGCACCTTATTATGTGGGGCATTCACCATCCATCCAGCACGCAGGAGAAAAACGATCTGTACGGTACCCAGTCTCTGTCTATCTCC GTGGGATCTTCCACCTATAGAAACAATTTCGTACCCGTAGTGGGGGCCAGGCCGCAAGTCAATGGCCAATCTGGCCGCATCGACTTCCACT GGACCTTGGTACAGCCTGGCGACAATATCACATTCTCACACAACGGCGGCTTGATTGCTCCGAGCCGGGTGTCTAAGTTGATCGGAAGAGG GCTCGGCATCCAGTCTGATGCCCCTATCGATAACAACTGTGAAAGTAAGTGCTTCTGGCGGGGCGGAAGTATTAATACCAGGCTGCCATTC CAGAATCTTTCCCCCCGCACCGTCGGTCAATGTCCCAAATACGTGAACAAGCGAAGCCTGATGTTGGCCACTGGGATGCGAAATGTTCCTG AGCTGATGCAGGGACGAGGACTGTTCGGAGCCATTGCTGGGTTCCTCGAGAACGGCTGGGAGGGCATGGTTGACGGTTGGTACGGATTCA GACACCAAAACGCGCAGGGCACGGGTCAAGCCGCCGATTACAAATCTACACAGGCGGCGATTGACCAGATTACAGGTAAGCTGAATAGAC TCGTGGAGAAAACAAACACCGAATTCGAGTCAATAGAGTCCGAGTTCTCTGAGATAGAACACCAGATCGGCAATGTAATCAACTGGACGAAA GACTCTATTACAGACATTTGGACCTACCAGGCGGAACTCCTGGTGGCTATGGAGAACCAGCATACTATTGATATGGCTGACTCTGAGATGCT GAATCTGTACGAGCGCGTCCGGAAGCAACTGCGTCAGAATGCAGAGGAAGACGGAAAGGGGTGCTTTGAGATTTATCACGCCTGCGATGA TAGCTGTATGGAGTCAATCCGAAACAACACTTATGATCATAGCCAGTACCGTGAGGAGGCACTACTGAACAGGCTGAACATTAATCCCGTCA CCCTGAGCAGTGGCTACAAAGACATCATCTTGTGGTTCTCTTTCGGCGCCTCATGCTTTGTCTTGCTTGCCGTAGTCATGGGTTTAGTGTTTT TTTGTCTCAAAAACGGTAATATGCGGTGTACAATTTGCATC
3070	ATGTACAAAATCGTGGTAATAATCGCGCTTCTGGGCGCAGTGAAGGGCCTGGACAAAATCTGCCTGGGACATCACGCTGTGGCAAATGGGA CCATCGTGAAAACCCTCACCAATGAGAAAGAGGAAGTGACCAACGCCACGGAAACAGTAGAAAGCACAGGATTAAACCGGCTTTGCATGAA AGGCCGCAAGCACAAAGACCTGGGCAATTGCCATCCCATCGGGATGCTGATCGGATCACCCGCCTGTGATCTCCATCTAACCGCACCTG GGACACTCTGATCGAAAGGGAGAACGCGATCGCTTACTGCTACCCAGGCGCTACCGTTAATGGCGAGGCTTTGCGCCAGAAGATCATGGA GTCAGGAGGGATCGACAAGATTAGTACAGGCTTCACCTATGAGAGCTCGATCAATTCCGCAGGAACAACCCGGGCCTGCATGAGAAACGG CGGCAATTCCTTTTACGCCGAACTTAAGTGGCTGGTGAGTAAAAGCAAGGGACAAAACTTCCCACAGACCACCAACACCTACCGTAATACC GATACCGCTGAGCATTTGATCATGTGGGGAATTCATCATCCCAGCTCCACACAGGAGAAGAATGATCTTTACGGGACACAGTCACTCAGCAT TTCCGTCGGTTCGAGCACTTACCGGAATAATTTTGTCCCTGTGGTCGGCGCCCGCCCTCAGGTCAACGGGCAGAGCGGCAGGATCGACTT CCATTGGACACTAGTTCAGCCTGGCGACAATATTACCTTTAGCCACAACGGCGGTCTGATCGCTCCTTCGCGTGTCTCAAAACTGATCGGTC GGGGGTTAGGCATTCAAAGCGATGCCCCCATTGACAACAATTGTGAATCAAAATGTTTCTGGCGCGGTGGCTCCATTAACACCAGACTGCC TTTCCAGAATCTTAGCCCAAGAACAGTCGGCCAGTGCCCCAAGTATGTGAATAAAAGGTCTCTGATGCTGGCTACCGGCATGCGTAATGTC CCAGAGCTTATGCAAGGCAGAGGACTGTTCGGCGCAATCGCGGGTTTTCTTGAGAACGGATGGGAGGGTATGGTTGACGGATGGTACGGT TTCCGGCACCAGAACGCCCAGGGAACTGGCCAGGCCGCAGACTACAAGTCCACACAGGCGGCAATTGACCAGATCACTGGCAAACTTAAC CGTCTAGTGGAAAAAACCAACACTGAATTCGAGAGTATCGAGAGTGAGTTCTCCGAGATTGAGCATCAGATAGGGAACGTGATCAATTGGA CAAAGGACTCTATTACAGACATTTGGACTTATCAGGCCGAGCTGCTGGTGGCCATGGAGAATCAGCACACAATCGACATGGCCGACTCTGA GATGCTCAATCTTTATGAACCGGTGCGCAAACAGCTGCGACAGAATGCAGAGGACGATGGGAAGGGGTGCTTCGAAATCTATCACGCATG CGACGATTCATGTATGGAGTCCATCCGGAATAATACATACGATCATAGTCAGTATCGCGAGGAAGCCTTATTGAACAGGCTGAACATTAATC CAGTCACTCTCAGCTCAGGGTACAAGGACATCATTTTGTGGTTCTCATTTGGAGCATCGTGCTTCGTGCTCCTTGCGGTTGTCATGGGTCTG GTCTTTTTTTGCCTGAAGAATGGAAACATGAGATGCACCATATGTATC

3071	ATGTACAAGATCGTGGTTATTATCGCCCTCCTGGGTGCTGTGAAAGGACTGGATAAGATCTGTCTGGGACACCATGCTGTGGCAAACGGCA CTATTGTTAAAACTCTGACTAACGAGCAAGAAGAAGTTACTAATGCTACTGAGACGGTCGAAAGTACTGGAATCAATCGGCTTTGCATGAAG GGCCGGAAGCACAAAGACCTTGGGAATTGCCATCCTATTGGGATGCTGATCGGTACCCCCGCATGTGACCTCCATCTGACTGGGATGTGG GACACGCTCATCGAGAGGGAGAATGCCATCGCTTACTGTTATCCCGGAGCCACAGTTAATGTCGAAGCCTTACGACAAAAAATCATGGAGT CAGGTGGTATCAACAAGATCAGCACCGGTTTCACATACGGCTCCTCCATCAACTCAGCAGGTACTACACGCGCATGCATGCGGAACGGCG GAAACAGTTTCTACGCCGAGCTGAAGTGGCTTGTTAGCAAATCAAAAGGTCAGAATTTCCCTCAGACTACAAACACATACCGTAACACCGAT ACCGCAGAACATCTCATCATGTGGGGTATTCATCACCCAAGTTCCACACAAGAAAAGAACGATCTCTACGGGACTCAATCACTCTCCATCTC CGTCGGATCCTCTACATACCGTAACAACTTTGTTCCTGTGGTGGGCGCAAGACCTCAGGTTAATGGGCAGTCAGGTAGAATTGACTTCCATT GGACACTGGTGCAACCTGGGGACAACATCACCTTCTCCCACAACGGCGGTCTGATCGCGCCCTCAAGGGTCTCCAAGCTGATAGGTCGGG GCCTCGGCATCCAGAGCGACGCCCCGATCGATAACAATTGTGAAAGCAAATGCTTCTGGAGGGGGGGCTCTATCAACACACGACTGCCTTT CCAGAATCTCTCTCCCCGAACCGTGGGACAGTGTCCTAAATACGTCAACAGGAGGAGCCTGATGCTCGCCACAGGTATGCGTAATGTCCCA GAGCTGATTCAAGGCCGCGGGTTATTTGGCGCGATAGCCGGTTTTCTGGAGAACGGCTGGGAGGGAATGGTGGATGGCTGGTATGGATTC CGGCACCAAAATGCTCAGGGAACTGGTCAAGCAGCCGACTATAAATCCACGCAGGCCGCCATCGACCAGATTACTGGCAAATTAAATCGCC TCGTGGAGAAGACTAACACTGAGTTCGAATCCATAGAGTCAGAGTTTTCCGAGATCGAGCACCAGATCGGTAATGTGATTAATTGGACCAAA GATTCCATCACTGATATTTGGACATACCAAGCCGAGCTCTTGGTGGCTATGGAGAACCAACATACTATCGATATGGCCGATAGCGAAATGCT CAACCTGTACGAGAGGGTCAGGAAACAGCTTCGCCAGAACGCAGAGGAAGACGGCAAAGGATGTTTCGAGATTTACCATGCGTGCGACGA TTCTTGCATGGAGAGCATTCGGAACAACACGTATGATCACAGCCAGTACAGGGAAGAGGCACTTCTGAACCGCCTGAACATAAATCCAGTC ACATTGAGCAGCGGCTACAAAGACATTATACTCTCGTTTTCATTTGGGGCCTCTTGCTTCGTCCTGCTGGCAGTGGTGATCGGATTGTTCTT CTTTTGCCTCAAAAACGGTAATATGCGCTGCACCATTTGCATC
3072	ATGTATAAAATAGTGGTCATTATTGCTCTTCTCGGTGCAGTTAAGGGTCTGGATAAGATCTGCTTGGGCCATCACGCTGTTGCCAATGGGAC CATAGTGAAGACCTTGACAAACGAACAGGAAGAAGTGACAAACGCCACCGAAACCGTCGAATCCACTGGCATTAACCGGCTGTGCATGAAG GGGCGAAAACACAAAGACCTCGGTAATTGTCATCCCATAGGGATGCTGATTGGTACACCTGCCTGTGACCTGCACCTGACCGGAATGTGGG ATACGCTCATTGAACGAGAGAATGCCATCGCCTATTGTTATCCGGGTGCAACTGTGAACGTGGAGGCCCTGCGTCAAAAGATCATGGAAAG CGGGGGGATAAACAAAATCTCGACCGGTTTCACGTATGGTTCAAGCATTAACAGTGCCGGCACAACAAGGGCATGCATGCGCAATGGAGGT AACTCATTCTACGCCGAGCTCAAGTGGCTCGTGAGCAAGTCCAAAGGACAGAACTTCCCTCAGACTACCAATACCTACCGTAATACTGACAC AGCGGAACATCTGATAATGTGGGGCATTCACCATCCTTCCTCGACCCAGGAAAAAAACGACCTGTACGGAACCCAGTCTCTCAGCATCTCC GTCGGGAGTAGCACTTACCGCAATAACTTTGTGCCCGTCGTGGGAGCCCGCCCCCAAGTGAATGGACAGAGCGGGAGGATTGATTTCCAC TGGACCCTCGTGCAACCGGGGGACAATATTACTTTCAGTCATAATGGAGGACTGATCGCCCCCAGTCGCGTGTCCAAACTGATCGGAAGGG GGCTCGGTATCCAATCCGATGCACCGATAGATAACAATTGCGAGTCTAAGTGTTTCTGGCGTGGAGGCAGTATTAACACTCGGTTACCTTTT CAAAATCTTTCGCCGCGGACCGTTGGCCAGTGCCCTAAGTACGTCAATCGTAGGTCCCTCATGCTAGCCACTGGAATGCGCAACGTCCCGG AGCTCATCCAGGGGCGTGGATTGTTTGGAGCTATTGCTGGCTTCCTGGAGAATGGCTGGGAGGGAATGGTCGATGGCTGGTACGGGTTTC GGCATCAGAACGCACAGGGCACAGGCCAGGCCGCCGACTACAAGTCTACACAGGCCGCCATTGATCAGATCACCGGGAAGCTGAACCGG CTGGTTGAGAAGACCAATACCGAATTTGAGAGCATCGAATCGGAATTCTCGGAAATCGAGCACCAGATTGGTAACGTAATAAACTGGACAAA GGATTCCATTACGGACATTTGGACGTATCAAGCAGAACTGCTCGTGGCCATGGAGAACCAACACACGATTGATATGGCAGATTCAGAAATG CTAAACTTGTACGAACGGGTCAGAAAGCAGCTGCGTCAGAATGCTGAGGAGGACGGCAAGGGATGTTTTGAGATCTACCATGCTTGCGATG ATTCTTGTATGGAGTCCATTCGGAATAACACGTACGACCATTCACAATATCGCGAGGAAGCACTTTTAAACCGGCTGAATATCAATCCAGTCA CACTGAGTTCAGGCTACAAGGATATCATATTATGGTTCTCATTCGGTGCCTCATGCTTCGTCCTTTTGGCAGTGGTGATGGGTTTGTTCTTTT TTTGTCTGAAGAATGGCAACATGAGATGTACAATATGCATC
3073	ATGTACAAGATTGTGGTGATCATCGCACTCCTTGGAGCTGTGAAAGGACTGGACAAAATTTGTCTGGGACATCACGCTGTGGCTAACGGGA CCATCGTGAAAACCCTAACGAATGAGCAGGAAGAAGTCACAAATGCGACAGAGACAGTGGAATCCACTGGTATCAATCGACTGTGTATGAA GGGCAGAAAACACAAGGACCTGGGCAATTGTCATCCGATAGGCATGTTAATCGGCACACCAGCCTGTGACCTGCATCTGACCGCATGTG GGATACTCTTATTGAGCGGGAGAACGCAATAGCTTACTGCTACCCAGGCGCTACTGTGAATGTGGAAGCTCTGCGTCAGAAAATTATGGAG AGCGGAGGGATTAACAAAATCTCTACTGGTTTCACATACGGCAGCAGTATTAACTCAGCCGGCACCACCAGGGCGTGTATGCGGAATGGTG GAAACTCATTTTATGCTGAGCTTAAATGGCTTGTCTCAAAGTCGAAAGGCCAGAACTTCCCCCAGACTACAAACACCTACAGGAATACGGAT ACAGCCGAGCATCTCATTATGTGGGGAATACATCATCCCAGCAGTACCCAGGAGAAAAACGATCTGTACGGAACCCAGAGTCTCTCAATCT CAGTGGGATCAAGCACGTACAGAAATAATTTCGTCCCTGTGGTGGGAGCCAGACCCCAGGTAAATGGACAGTCAGGGCGCATTGACTTTCA CTGGACTCTCGTCCAGCCAGGAGACAATATCACATTTAGTCACAATGGCGGCCTGATAGCTCCTTCTCGAGTGAGCAAACTGATCGGACGG GGTCTCGGTATCCAGTCCGACGCCCCCATCGATAACAACTGCGAGTCGAAGTGCTTCTGGCGTGGCGGGTCAATTAACACAAGACTACCAT TTCAGAACCTTTCACCTAGAACCGTGGGACAGTGCCCAAAGTACGTCAATAGACGGAGCCTGATGCTGGCCACAGGGATGAGGAACGTTCC TGAACTGATACAAGGTCGGGGGCTTTTCGGTGCCATCGCAGGTTTTCTCGAGAACGGCTGGGAGGGCATGGTGGATGGGTGGTATGGGTT TAGGCACCAAAATGCACAAGGGACCGGCCAGGCCGCCGATTATAAGTCTACGCAAGCCGCCATTGACCAGATCACTGGGAAGCTTAATAG GTTAGTAGAGAAGACAAATACCGAATTCGAGAGTATTGAGTCAGAATTCTCCGAGATCGAGCACCAGATAGGCAACGTCATTAATTGGACAA AGGACTCTATTACCGACATTTGGACGTATCAAGCCGAACTTCTGGTGGCAATGGAAAATCAGCACACAATCGACATGGCCGACTCGGAGAT GTTGAACTTATATGAAAGGGTGCGCAAACAGCTGCGCCAGAACGCTGAAGAAGACGGTAAGGGATGCTTTGAGATCTATCATGCCTGCGAT GATTCCTGTATGGAGTCGATAAGAAACAATACCTATGACCATAGCCAGTACCGGGAAGAAGCCCTCCTGAACAGACTGAATATTAACCCCGT TACCTTGTCCTCAGGATACAAGGACATTATCCTGTGGTTTTCTTTCGGTGCTAGCTGTTTCGTGCTGCTCGCCGTCGTAATGGGGTTGTTTTT CTTCTGCCTCAAAAACGGTAACATGCGATGCACCATTTGCATT

	ATGTATAAGATCGTCGTCATTATCGCTCTCCTCGGAGCCGTGAAAGGTCTTGACAAAATTTGTCTGGGCCATCACGCAGTGGCCAATGGGA CCATCGTGAAGACGCTGACGAACGAGAAAGAAGAGGTGACAAACGCTACTGAGACGGTCGAGAGCACCGGACTCAATAGGTTGTGCATGA AAGGACGAAAGCATAAAGATCTGGGCAACTGTCACCCCATCGGGATGCTGATTGGCTCCCCAGCCTGCGACTTACACCTGACTGGCACCTG GGATACACTGATCGAGCGCGAGAACGCCATTGCCTACTGTTACCCCGGCGCTACAGTGAATGGTGAGGCCCTTCGCCAAAAGATTATGGA GTCAGGCGGAATAGATAAGATCTCTACAGGCTTCACTTATGAGAGTTCGATTAACTCGGCCGGGACAACTAGGGCGTGTATGCGGAATGGC GGGAATTCCTTCTATGCTGAGCTTAAGTGGTTGGTTTCCAAGAGCAAGGGGCAGAATTTTCCTCAGACCACCAACACCTATCGAAACACAGA CACCGCTGAACACCTCATCATGTGGGGCATTCACCACCCATCTTCAACCCAGGAGAAGAACGACCTCTACGGTACTCAAAGCCTTAGTATCA GCGTGGGGTCATCTACCTACCGGAACAATTTTGTGCCCGTGGTGGGAGCGCGCCCTCAAGTGAACGGACAAAGTGGCAGGATCGATTTCC ATTGGACTTTGGTCCAGCCTGGTGACAACATCACATTCAGTCACAACGGTGGCCTGATCGCACCGTCCAGAGTTTCCAAACTGATCGGCCG GGGTCTAGGGATTCAGAGCGACGCCCCTATTGATAACAACTGCGAGTCAAAATGCTTCTGGAGGGGGGGCTCCATTAACACTCGACTGCCC TTTCAGAATCTGTCACCCCGAACCGTGGGTCAGTGTCCTAAGTACGTGAACAAGAGATCACTCATGCTTGCCACTGGGATGCGCAATGTGC CCGAGCTCATGCAGGGAAGAGGACTGTTCGGAGCGATTGCTGGGTTTCTGGAAAATGGGTGGGAAGGGATGGTAGATGGATGGTACGGAT TTCGTCATCAAAACGCACAGGGCACTGGACAAGCGGCGGACTATAAGTCAACTCAGGCTGCCATAGACCAAATAACCGGGAAACTAAACCG GCTCGTCGAGAAGACCAACACCGAATTTGAATCTATCGAGTCGGAATTTAGCGAGATTGAGCACCAGATAGGAAATGTTATCAATTGGACCA AGGACTCCATTACGGATATTTGGACTTACCAGGCGGAGCTGCTGGTTGCCATGGAGAATCAGCACACTATTGACATGGCCGATAGTGAGAT GTTGAATTTGTACGAAAGAGTGCGCAAGCAGCTGAGGCAGAACGCTGAAGAGGACGGTAAGGGATGCTTCGAAATTTATCACGCATGCGAT GATTCCTGCATGGAATCGATCAGAAACAATACATATGATCATTCTCAATACCGGGAGGAGGCTTTGCTCAACCGCCTGAATATAAATCCTGT GACTCTCAGTTCGGGCTATAAAGACATCATCCTGTGGTTCTCTTTTGGCGCCTCCTGCTTTGTCCTGCTGGCGGTGGTGATGGGACTCGTCT
3074	TCTTCTGCCTAAAGAATGGCAACATGCGGTGTACAATCTGTATT

[0125] An alignment of amino acid sequences of hemagglutinin proteins from influenza A H7N9 strains are shown relative to a consensus sequence in Figure 22. An alignment of amino acid sequences of hemagglutinin proteins from influenza A H10N8 strains are shown relative to a consensus sequence in Figure 23.

[0126] Aspects of the present disclosure provide RNA polynucleotides encoding an influenza hemagglutinin protein or a fragment thereof. The terms "hemagglutinin," "hemagglutinin protein," and "HA" may be used interchangeably throughout and refer to a hemagglutinin protein that may be present on the surface of an influenza virus. On the viral surface, the hemagglutinin protein is present in homotrimers, each monomer of which is comprised of two subunits, HA1 and HA2, linked by a disulfide bond. Structurally, hemagglutinin proteins are comprised of several domains: a globular head domain, a stalk domain (also referred to as a stem domain), a transmembrane domain, and a cytoplasmic domain. It is generally though that during infection of a host cell (*e.g.*, a eukaryotic cell such as a human cell) with an influenza virus, the hemagglutinin protein recognizes and binds to sialic acid of a receptor on the surface of a host cell facilitating attachment of the virus to the host cell (Gamblin et al., 2004; Ha et al., 2000). Following endocytosis of the virus and acidification of the endosome, the hemagglutinin protein undergoes a pH-dependent conformational change that allows for the hemagglutinin protein to facilitate fusion of the viral envelope with the endosome membrane of host cell and entry of the viral nucleic acid into the host cell.

[0127] In general, influenza viruses are classified based on the amino acid sequence of the viral hemagglutinin protein and/orthe amino acid sequence of the viral neuraminidase (NA). In some embodiments, the hemagglutinin is of the subtype H1, H2, H3, H4, H5, H6, H7, H8, H9, H10, H11, H12, H13, H14, H15, H16, H17, or H18. The differences in amino acid sequences between hemagglutinin proteins of different subtypes are largely found within the sequence of the head domain of the protein. The amino acid sequence of the stalk domain is considered to be more conserved between hemagglutinin subtypes compared to sequence of the head domain. Domains of the hemagglutinin protein may be predicted using conventional methods known in the art.

[0128] Many naturally occurring and experimentally derived antibodies that bind and neutralize the hemagglutinin protein are thought to bind epitopes within the head domain of hemagglutinin and prevent or reduce interaction of hemagglutinin with sialic acid on receptors of host cells, thereby preventing or reducing infection of the cell. Alternatively or in addition, neutralizing antibodies may prevent or reduce fusion of the virus membrane with the membrane of the endosome. Such antibodies may bind epitopes within the stalk domain, thereby inhibiting the conformations change of the protein.

[0129] In some embodiments, the RNA polynucleotides described herein encode a fragment of a hemagglutinin protein, such as a truncated hemagglutinin protein. In some embodiments, the fragment is a headless hemagglutinin, meaning the fragment does not comprise the head domain In some embodiments, the fragment comprises a portion of the head domain. In some embodiments, the fragment is a stalk fragment (see, *e.g.*, Mallajosyula et al. PNAS (2014) E2514-E2523). In some embodiments, fragment does not comprise the cytoplasmic domain. In some embodiments, the fragment does not comprise the transmembrane domain. In such embodiments, the fragment may be referred to as a soluble or secreted hemagglutinin protein or fragment.

[0130] In some embodiments, the RNA polypeptide encodes a hemagglutinin protein or fragment thereof that is at least 90% (*e.g.*, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99%) identical to a hemagglutinin protein provided by

an amino acid sequence in Table 14-16. The terms "identical" or percent "identity" in the context of two or more polypeptide sequences refer to two or more sequences that are the same. The percent identity between polypeptide sequences may be performed using algorithms known in the art, such as BLAST and CLUSTAL.

[0131] The sequence of the hemagglutinin protein or fragment thereof may be obtained from any source. In some embodiments, the sequence of the hemagglutinin protein or fragment thereof is from an avian influenza strain. In some embodiments, the sequence of the hemagglutinin protein or fragment thereof is from an avian influenza strain that is capable of or at risk of infecting human subjects. In some embodiments, the sequence of the hemagglutinin protein or fragment thereof is from a porcine influenza strain. In some embodiments, the sequence of the hemagglutinin protein or fragment thereof is from a porcine influenza strain that is capable of or at risk of infecting human subjects.

[0132] In any of the embodiments described herein, the sequence of the hemagglutinin or fragment thereof may be modified or optimized (such as codon optimized) for expression in a particular cell or host organism.

Dengue

[0133] Dengue fever is a mosquito-borne virus and is epidemic (SE Asia) and endemic (Sub-Saharan Africa, India). As much as 40% of world's population is at risk with over 100 million infections per year according to the World Health Organiztion. There have been about 1-2 million clinically documented cases. The mortality rate depends on access to healthcare and can reach as high as 20%.

[0134] Symptoms include acute onset fevers with terrible joint and muscle pain for 5-7 days, followed by weeks of lethargy and fatigue. Dehydration and hemorrhage are the main drivers of mortality, hence need for access to IV fluids to avoid shock. Supportive care includes fluid resuscitation but prevention is the primary means of limiting the impact of virus (e.g., mosquito control, personal protection).

[0135] For Dengue fever, disease characteristics requires a neutralizing - but not enhancing - antibody response to the four most critical Dengue serotypes (DENV1-4). Therefore a multivalent antigen targeting key proteins/protein domains of four serotypes of Dengue virus (DENV 1-4) would have value as a vaccine.

[0136] In one embodiment, NAVs of the invention comprise one or more polynucleotides which encode the E protein domain III (DENV1-4 tandem mRNA), the E protein domain I/II hinge region (DENV1-4 individual mRNAs), the prM protein (DENV1-4 tandem or individual mRNAs) and the C protein (DENV1-4 tandem or single mRNAs).

[0137] In one embodiment, the most potent NAV vaccine is selected by measuring the antibody titer in Balb/c mice followed by tests of selected vaccines in Dengue disease models. Upon rescue in disease model, crossreactivity analysis in in-vitro viral assays to ensure activity of multi-valent vaccine against all serotypes is performed.

[0138] In some embodiments cross-neutralizing Ab-titers agains each of the four Dengue strains (DENV1-4) in a virus neutralization assay are tested either in vitro or in vivo as in (BMC Microbiol. 2014; 14: 44; and Immunology. 2012 Jul; 136 (3):334-43).

[0139] In some embodiments, neutralizing vs enhancing Ab-titers is evaluated and in another embodiment the RNAVare tested in humanized mouse model for Dengue disease (J Virol. 2014 Feb;88(4):2205-18).

[0140] The NAV targeting Dengue is referred to as a multi-genotype antigen NAV.

Middle-east respiratory syndrome Coronavirus (MERS-CoV)

[0141] MERS-CoV, previously known as the Novel Coronavirus or SARS-like virus, is a member of the coronavirus family. Symptoms are similar to SARS infections and include coughing, production of mucous, shortness of breath, malaise - a general feeling of being unwell, chest pain, fever, diarrhea (in some cases) and renal (kidney) failure.*Human enterovirus 71 and Human enterovirus 68*

[0142] Enterovirus 71 (EV-71) is one of the major causative agents for hand, foot and mouth disease (HFMD), and is sometimes associated with severe central nervous system diseases. The Enterovirus 71 (EV71) infection may be asymptomatic.

[0143] Enterovirus 68 (EV68, EV-D68) is a member of the Picornaviridae family, an enterovirus (a group of ssRNA viruses containing the polioviruses, coxsackieviruses, and echoviruses). First isolated in 1962, it has been been on a worldwide upswing in the last few years. It may be involved in cases of a recent outbreak of polio-like disease in California.

Antigens

[0144] Antigens encoded by the NAVs of the present invention include polypeptides, peptides and/or polypeptides of interest and are encoded by the polynucleotides of the invention. Polynucleotides encoding such antigens are described in more detail below under "Design, Synthesis and Quantitation of NAV Polynucleotides". References herein to "NAVs of the invention" relate to those as set out in the appended claims. References to polynucleotides, polypeptides or antigens "of the invention" relate to those in the context of the NAVs of the invention.

III. Design, Synthesis and Quantification of NAV Polynucleotides

[0145] According to the present disclosure, the polynucleotides encode at least one polypeptide of interest, e.g., an antigen. Antigens of the present disclosuremay be wild type (i.e., derived from the infectious agent) or modified (e.g., engineered, designed or artificial). They may have any combination of the features described herein.

[0146] NAVs of the present invention comprise nucleic acid molecules, specifically mRNA polynucleotides which encode one or more peptides or polypeptides of interest. Such peptides or polypeptides, according to the invention serve as an antigen or antigenic molecule.

[0147] In one embodiment, linear polynucleotides encoding one or more antigens of the NAVs of the present invention which are made using only *in vitro* transcription (IVT) enzymatic synthesis methods are referred to as "IVT polynucleotides."

[0148] In another embodiment, the polynucleotides of the present invention which have portions or regions which differ in size and/or chemical modification pattern, chemical modification position, chemical modification percent or chemical modification population and combinations of the foregoing are known as "chimeric polynucleotides" A "chimera" according to the present invention is an entity having two or more incongruous or heterogeneous parts or regions As used herein a "part" or "region" of a polynucleotide is defined as any portion of the polynucleotide which is less than the entire length of the polynucleotide.

[0149] In yet another embodiment, the polynucleotides of the present invention that are circular are known as "circular polynucleotides" or "circP." As used herein, "circular polynucleotides" or "circP" means a single stranded circular polynucleotide which acts substantially like, and has the properties of, an RNA. The term "circular" is also meant to encompass any secondary or tertiary configuration of the circP.

[0150] In some embodiments, the polynucleotide includes from about 30 to about 100,000 nucleotides (e.g., from 30 to 50, from 30 to 100, from 30 to 250, from 30 to 500, from 30 to 1,000, from 30 to 1,500, from 30 to 3,000, from 30 to 5,000, from 30 to 7,000, from 30 to 10,000, from 30 to 25,000, from 30 to 50,000, from 30 to 70,000, from 100 to 250, from 100 to 500, from 100 to 1,000, from 100 to 1,500, from 100 to 3,000, from 100 to 5,000, from 100 to 7,000, from 100 to 10,000, from 100 to 25,000, from 100 to 50,000, from 100 to 70,000, from 100 to 100,000, from 500 to 1,000, from 500 to 1,500, from 500 to 2,000, from 500 to 3,000, from 500 to 5,000, from 500 to 7,000, from 500 to 10,000, from 500 to 25,000, from 500 to 50,000, from 500 to 70,000, from 500 to 100,000, from 1,000 to 1,500, from 1,000 to 2,000, from 1,000 to 3,000, from 1,000 to 5,000, from 1,000 to 7,000, from 1,000 to 10,000, from 1,000 to 25,000, from 1,000 to 50,000, from 1,000 to 70,000, from 1,000 to 100,000, from 1,500 to 3,000, from 1,500 to 5,000, from 1,500 to 7,000, from 1,500 to 10,000, from 1,500 to 25,000, from 1,500 to 50,000, from 1,500 to 70,000, from 1,500 to 100,000, from 2,000 to 3,000, from 2,000 to 5,000, from 2,000 to 7,000, from 2,000 to 10,000, from 2,000 to 25,000, from 2,000 to 50,000, from 2,000 to 70,000, and from 2,000 to 100,000).

[0151] In one embodiment, the length of a region encoding at least one peptide polypeptide of interest of the polynucleotides present invention is greater than about 30 nucleotides in length (e.g., at least or greater than about 35, 40, 45, 50, 55, 60, 70, 80, 90, 100, 120, 140, 160, 180, 200, 250, 300,350,400,450,500,600,700,800,900,1,000,1,100,1,200,1,300,1,400,1,500,1,600,1,700,1,800,1,900,2,000,2,500, and 3,000, 4,000, 5,000, 6,000, 7,000, 8,000, 9,000, 10,000, 20,000, 30,000, 40,000, 50,000, 60,000, 70,000, 80,000, 90,000 or up to and including 100,000 nucleotides). As used herein, such a region may be referred to as a "coding region" or "region encoding."

[0152] The polynucleotides of the present invention is or functions as a messenger RNA (mRNA). As used herein, the term "messenger RNA" (mRNA) refers to any polynucleotide which encodes at least one peptide or polypeptide of interest and which is capable of being translated to produce the encoded peptide polypeptide of interest *in vitro, in vivo, in situ* or *ex vivo.*

[0153] The polynucleotides of the present invention are chemically modified. As used herein, a "structural" modification is one in which two or more linked nucleosides are inserted, deleted, duplicated, inverted or randomized in a polynucleotide without significant chemical modification to the nucleotides themselves. Because chemical bonds will necessarily be broken and reformed to effect a structural modification, structural modifications are of a chemical nature and hence are chemical modifications. However, structural modifications will result in a different sequence of nucleotides. For example, the polynucleotide "ATCG" may be chemically modified to "AT-5meC-G". The same polynucleotide may be structurally modified from "ATCG" to "ATCCCG". Here, the dinucleotide "CC" has been inserted, resulting in a structural modification to the polynucleotide.

[0154] In one embodiment, the polynucleotides of the present invention, such as IVT polynucleotides or circular polynucleotides, may have a uniform chemical modification of all or any of the same nucleoside type or a population of modifications produced by mere downward titration of the same starting modification in all or any of the same nucleoside type, or a measured percent of a chemical modification of all any of the same nucleoside type but with random incorporation, such as where all uridines are replaced by a uridine analog, e.g., pseudouridine. In another embodiment, the polynucleotides may have a uniform chemical modification of two, three, or four of the same nucleoside type throughout

the entire polynucleotide (such as all uridines and all cytosines, etc. are modified in the same way).

[0155] When the polynucleotides of the present invention are chemically and/or structurally modified the polynucleotides may be referred to as "modified polynucleotides."

[0156] In one embodiment, the polynucleotides of the present invention may include a sequence encoding a self-cleaving peptide. The self-cleaving peptide may be, but is not limited to, a 2A peptide. As a non-limiting example, the 2A peptide may have the protein sequence: GSGATNFSLLKQAGDVEENPGP (SEQ ID NO: 2005), fragments or variants thereof. In one embodiment, the 2A peptide cleaves between the last glycine and last proline. As another non-limiting example, the polynucleotides of the present invention may include a polynucleotide sequence encoding the 2A peptide having the protein sequence GSGATNFSLLKQAGDVEENPGP (SEQ ID NO: 2005) fragments or variants thereof.

[0157] One such polynucleotide sequence encoding the 2A peptide is GGAAGCGGAGCTACTAACTTCAGCCTGCTGAAGCAGGCTGGAGACGTGGAGGAGAACCCTGGACCT (SEQ ID NO: 2006). The polynucleotide sequence of the 2A peptide may be modified or codon optimized by the methods described herein and/or are known in the art.

[0158] In one embodiment, this sequence may be used to separate the coding region of two or more polypeptides of interest. As a non-limiting example, the sequence encoding the 2A peptide may be between a first coding region A and a second coding region B (A-2Apep-B). The presence of the 2A peptide would result in the cleavage of one long protein into protein A, protein B and the 2A peptide. Protein A and protein B may be the same or different peptides or polypeptides of interest. In another embodiment, the 2A peptide may be used in the polynucleotides of the present invention to produce two, three, four, five, six, seven, eight, nine, ten or more proteins.

Polynucleotide Architecture

[0159] Traditionally, the basic components of an mRNA molecule include at least a coding region, a 5'UTR, a 3'UTR, a 5' cap and a poly-A tail. The IVT polynucleotides of the present invention may function as mRNA but are distinguished from wild-type mRNA in their functional and/or structural design features which serve to overcome existing problems of effective polypeptide production using nucleic-acid based therapeutics. It is to be understood that the antigens of the NAVs of the present invention may be encoded by IVT polynucleotides, as described herein.

[0160] Figure 1 shows a primary construct **100** of an IVT polynucleotide of the present invention. Such polynucleotides are useful in NAV compositions, RNAV compositions or mRNA vaccines. As used herein, "primary construct" refers to a polynucleotide of the present invention which encodes one or more polypeptides of interest and which retains sufficient structural and/or chemical features to allow the polypeptide of interest encoded therein to be translated.

[0161] According to FIG. **1A** and **1B,** the polynucleotide **100** here contains a first region of linked nucleotides **102** that is flanked by a first flanking region **104** and a second flaking region **106.** The polypeptide of interest may comprise at its 5' terminus one or more signal sequences encoded by a signal sequence region **103.** The flanking region **104** may comprise a region of linked nucleotides comprising one or more complete or incomplete 5' UTRs sequences which may be completely codon optimized or partially codon optimized. The flanking region **104** may include at least one nucleic acid sequence including, but not limited to, miR sequences, TERZAK™ sequences and translation control sequences. The flanking region **104** may also comprise a 5' terminal cap **108.** The 5' terminal capping region **108** may include a naturally occurring cap, a synthetic cap or an optimized cap. Non-limiting examples of optimized caps include the caps taught by Rhoads in US Patent No. US7074596 and International Patent Publication No. WO2008157668, WO2009149253 and WO2013103659. The second flanking region **106** may comprise a region of linked nucleotides comprising one or more complete or incomplete 3' UTRs. The second flanking region **106** may be completely codon optimized or partially codon optimized. The flanking region **106** may include at least one nucleic acid sequence including, but not limited to, miR sequences and translation control sequences. The flanking region **106** may also comprise a 3' tailing sequence **110.** The 3' tailing sequence **110** may include a synthetic tailing region **112** and/or a chain terminating nucleoside **114.** Non-liming examples of a synthetic tailing region include a polyA sequence, a polyC sequence, and/or a polyA-G quartet. Non-limiting examples of chain terminating nucleosides include 2'-0 methyl, F and locked nucleic acids (LNA).

[0162] Bridging the 5' terminus of the first region **102** and the first flanking region **104** is a first operational region **106.** Traditionally this operational region comprises a Start codon. The operational region may alternatively comprise any translation initiation sequence or signal including a Start codon.

[0163] Bridging the 3' terminus of the first region **102** and the second flanking region **106** is a second operational region **107.** Traditionally this operational region comprises a Stop codon. The operational region may alternatively comprise any translation initiation sequence or signal including a Stop codon. Multiple serial stop codons may also be used in the IVT polynucleotide. In one embodiment, the operation region of the present invention may comprise two stop codons. The first stop codon may be "TGA" or "UGA" and the second stop codon may be selected from the group consisting of "TAA," "TGA," "TAG," "UAA," "UGA" or "UAG."

[0164] The shortest length of the first region of the primary construct of the IVT polynucleotide of the present invention

can be the length of a nucleic acid sequence that is sufficient to encode for a dipeptide, a tripeptide, a tetrapeptide, a pentapeptide, a hexapeptide, a heptapeptide, an octapeptide, a nonapeptide, or a decapeptide. In another embodiment, the length may be sufficient to encode a peptide of 2-30 amino acids, e.g. 5-30, 10-30, 2-25, 5-25, 10-25, or 10-20 amino acids. The length may be sufficient to encode for a peptide of at least 11, 12, 13, 14, 15, 17, 20, 25 or 30 amino acids, or a peptide that is no longer than 40 amino acids, e.g. no longer than 35, 30, 25, 20, 17, 15, 14, 13, 12, 11 or 10 amino acids. Examples of dipeptides that the polynucleotide sequences can encode or include, but are not limited to, carnosine and anserine. It is understood that the NAV, RNAV or mRNA vaccines of the invention may be translatable and include such first region of a primary construct.

[0165] The length of the first region of the primary construct of the IVT polynucleotide encoding the polypeptide of interest of the present invention is greater than about 30 nucleotides in length (e.g., at least or greater than about 35, 40, 45, 50, 55, 60, 70, 80, 90, 100, 120, 140, 160, 180, 200, 250, 300, 350, 400, 450, 500, 600, 700, 800, 900, 1,000, 1,100, 1,200, 1,300, 1,400, 1,500, 1,600, 1,700, 1,800, 1,900, 2,000, 2,500, and 3,000, 4,000, 5,000, 6,000, 7,000, 8,000, 9,000, 10,000, 20,000, 30,000, 40,000, 50,000, 60,000, 70,000, 80,000, 90,000 or up to and including 100,000 nucleotides).

[0166] In some embodiments, the IVT polynucleotide includes from about 30 to about 100,000 nucleotides (e.g., from 30 to 50, from 30 to 100, from 30 to 250, from 30 to 500, from 30 to 1,000, from 30 to 1,500, from 30 to 3,000, from 30 to 5,000, from 30 to 7,000, from 30 to 10,000, from 30 to 25,000, from 30 to 50,000, from 30 to 70,000, from 100 to 250, from 100 to 500, from 100 to 1,000, from 100 to 1,500, from 100 to 3,000, from 100 to 5,000, from 100 to 7,000, from 100 to 10,000, from 100 to 25,000, from 100 to 50,000, from 100 to 70,000, from 100 to 100,000, from 500 to 1,000, from 500 to 1,500, from 500 to 2,000, from 500 to 3,000, from 500 to 5,000, from 500 to 7,000, from 500 to 10,000, from 500 to 25,000, from 500 to 50,000, from 500 to 70,000, from 500 to 100,000, from 1,000 to 1,500, from 1,000 to 2,000, from 1,000 to 3,000, from 1,000 to 5,000, from 1,000 to 7,000, from 1,000 to 10,000, from 1,000 to 25,000, from 1,000 to 50,000, from 1,000 to 70,000, from 1,000 to 100,000, from 1,500 to 3,000, from 1,500 to 5,000, from 1,500 to 7,000, from 1,500 to 10,000, from 1,500 to 25,000, from 1,500 to 50,000, from 1,500 to 70,000, from 1,500 to 100,000, from 2,000 to 3,000, from 2,000 to 5,000, from 2,000 to 7,000, from 2,000 to 10,000, from 2,000 to 25,000, from 2,000 to 50,000, from 2,000 to 70,000, and from 2,000 to 100,000).

[0167] According to the present invention, the first and second flanking regions of the IVT polynucleotide may range independently from 15-1,000 nucleotides in length (e.g., greater than 30, 40, 45, 50, 55, 60, 70, 80, 90, 100, 120, 140, 160, 180, 200, 250, 300, 350, 400, 450, 500, 600, 700, 800, and 900 nucleotides or at least 30, 40, 45, 50, 55, 60, 70, 80, 90, 100, 120, 140, 160, 180, 200, 250, 300, 350, 400, 450, 500, 600, 700, 800, 900, and 1,000 nucleotides).

[0168] According to the present invention, the tailing sequence of the IVT polynucleotide may range from absent to 500 nucleotides in length (e.g., at least 60, 70, 80, 90, 120, 140, 160, 180, 200, 250, 300, 350, 400, 450, or 500 nucleotides). Where the tailing region is a polyA tail, the length may be determined in units of or as a function of polyA Binding Protein binding. In this embodiment, the polyA tail is long enough to bind at least 4 monomers of PolyA Binding Protein. PolyA Binding Protein monomers bind to stretches of approximately 38 nucleotides. As such, it has been observed that polyA tails of about 80 nucleotides (SEQ ID NO: 2276) and 160 nucleotides (SEQ ID NO: 2277) are functional.

[0169] According to the present invention, the capping region of the IVT polynucleotide may comprise a single cap or a series of nucleotides forming the cap. In this embodiment the capping region may be from 1 to 10, e.g. 2-9, 3-8, 4-7, 1-5, 5-10, or at least 2, or 10 or fewer nucleotides in length In some embodiments, the cap is absent.

[0170] According to the present invention, the first and second operational regions of the IVT polynucleotide may range from 3 to 40, e.g., 5-30, 10-20,15, or at least 4, or 30 or fewer nucleotides in length and may comprise, in addition to a Start and/or Stop codon, one or more signal and/or restriction sequences

[0171] The IVT polynucleotides of the present invention are chemically modified. When the IVT polynucleotides of the present invention are chemically and/or structurally modified the polynucleotides may be referred to as "modified IVT polynucleotides."

[0172] In one embodiment, the IVT polynucleotides of the present invention may have a uniform chemical modification of all or any of the same nucleoside type or a population of modifications produced by mere downward titration of the same starting modification in all or any of the same nucleoside type, or a measured percent of a chemical modification of all any of the same nucleoside type but with random incorporation, such as where all uridines are replaced by a uridine analog, e.g., pseudouridine. In another embodiment, the IVT polynucleotides may have a uniform chemical modification of two, three, or four of the same nucleoside type throughout the entire polynucleotide (such as all uridines and all cytosines, etc. are modified in the same way).

[0173] In one embodiment, the IVT polynucleotides of the present invention may include a sequence encoding a self-cleaving peptide, described herein, such as but not limited to the 2A peptide. The polynucleotide sequence of the 2A peptide in the IVT polynucleotide may be modified or codon optimized by the methods described herein and/or are known in the art.

[0174] In one embodiment, this sequence may be used to separate the coding region of two or more polypeptides of

interest in the IVT polynucleotide.

[0175] The IVT polynucleotide of the present invention are chemically modified. When chemically modified and/or structurally modified the IVT polynucleotide may be referred to as a "modified IVT polynucleotide."

[0176] The IVT polynucleotide encodes at least one peptide or polypeptide of interest. In another embodiment, the IVT polynucleotide may encode two or more peptides or polypeptides of interest. Non-limiting examples of peptides or polypeptides of intest include heavy and light chains of antibodies, an enzyme and its substrate, a label and its binding molecule, a second messenger and its enzyme or the components of multimeric proteins or complexes.

Chimeric Polynucleotide Architecture

[0177] The chimeric polynucleotides or RNA constructs of the present invention maintain a modular organization similar to IVT polynucleotides, but the chimeric polynucleotides comprise one or more structural and/or chemical modifications or alterations which impart useful properties to the polynucleotide. As such, the chimeric polynucleotides which are modified mRNA molecules of the present invention are termed "chimeric modified mRNA" or "chimeric mRNA."

[0178] It is to be understood that the antigens of the NAVs of the present invention may be encoded by chimeric modified mRNA or chimeric mRNA.

[0179] Chimeric polynucleotides have portions or regions which differ in size and/or chemical modification pattern, chemical modification position, chemical modification percent or chemical modification population and combinations of the foregoing.

[0180] Examples of parts or regions, where the chimeric polynucleotide functions as an mRNA and encodes a polypeptide of interest include, but are not limited to, untranslated regions (UTRs, such as the 5' UTR or 3' UTR), coding regions, cap regions, polyA tail regions, start regions, stop regions, signal sequence regions, and combinations thereof. Figure 2 illustrates certain embodiments of the chimeric polynucleotides of the invention which may be used as mRNA. Figure 3 illustrates a schematic of a series of chimeric polynucleotides identifying various patterns of positional modifications and showing regions analogous to those regions of an mRNA polynucleotide. Regions or parts that join or lie between other regions may also be designed to have subregions. These are shown in the figure.

[0181] In some instances, the chimeric polynucleotides of the disclosure have a structure comprising Formula I.

$$5' [A_n]_x\text{-}L1\text{-}[B_o]_y\text{-}L2\text{-}[C_p]_z\text{-}L3' \qquad \textbf{Formula I}$$

wherein:

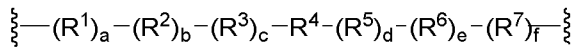
each of A and B independently comprise a region of linked nucleosides;
C is an optional region of linked nucleosides;
at least one of regions A, B, or C is positionally modified, wherein said positionally modified region comprises at least two chemically modified nucleosides of one or more of the same nucleoside type of adenosine, thymidine, guanosine, cytidine, or uridine, and wherein at least two of the chemical modifications of nucleosides of the same type are different chemical modifications;
n, o and p are independenty an integer between 15-1000;
x and y are independently 1-20;
z is 0-5,
L1 and L2 are independently optional linker moieties, said linker moieties being either nucleic acid based or non-nucleic acid based; and
L3 is an optional conjugate or an optional linker moiety, said linker moiety being either nucleic acid based or non-nucleic acid based.

[0182] In some instances the chimeric polynucleotide of Formula I encodes one or more peptides or polypeptides of interest. Such encoded molecules may be encoded across two or more regions

[0183] In another aspect, the disclosure features a chimeric polynucleotide encoding a polypeptide, wherein the polynucleotide has a sequence including Formula II:

$$[A_n]\text{-}L^1[B_o] \qquad \textbf{Formula II}$$

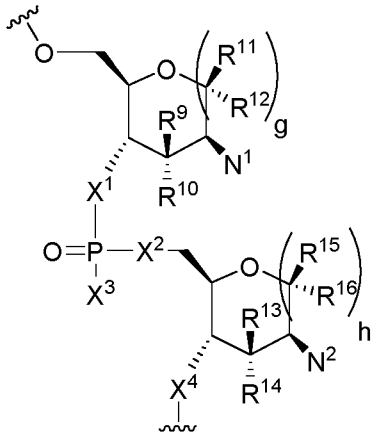
wherein each A and B is independently any nucleoside;
n and o are, independently 15 to 1000; and
L^1 has the structure of Formula III:



Formula III

wherein a, b, c, d, e, and f are each, independently, 0 or 1;
each of R^1, R^3, R^5, and R^7, is, independently, selected from optionally substituted C_1-C_6 alkylene, optionally substituted C_1-C_6 heteroalkylene, O, S, and NR^8;
R^2 and R^6 are each, independently, selected from carbonyl, thiocarbonyl, sulfonyl, or phosphoryl;
R^4 is optionally substituted C_1-C_{10} alkylene, optionally substituted C_2-C_{10} alkenylene, optionally substituted C_2-C_{10} alkynylene, optionally substituted C_2-C_9 heterocyclylene, optionally substituted C_6-C_{12} arylene, optionally substituted C_2-C_{100} polyethylene glycolene, or optionally substituted C_1-C_{10} heteroalkylene, or a bond linking $(R^1)_a$-$(R^2)_b$-$(R^3)_c$ to $(R^5)_d$-$(R^6)_e$-$(R^7)_f$, wherein if c, d, e, f, g, and h are 0, R^4 is not a bond; and
R^8 is hydrogen, optionally substituted C_1-C_4 alkyl, optionally substituted C_2-C_4 alkenyl, optionally substituted C_2-C_4 alkynyl, optionally substituted C_2-C_6 heterocyclyl, optionally substituted C_6-C_{12} aryl, or optionally substituted C_1-C_7 heteroalkyl;
wherein L^1 is attached to $[A_n]$ and $[B_o]$ at the sugar of one of the nucleosides (e.g., at the 3' position of a five-membered sugar ring or 4' position of a six membered sugar ring of a nucleoside of $[A_n]$ and the 5' position of a five-membered sugar ring or 6' position of of a six membered sugar ring of a nucleoside of $[B_o]$ or at the 5' position of a five-membered sugar ring or 6' position of of a six membered sugar ring of a nucleoside of $[A_n]$ and the 3' position of a five-membered sugar ring or 4' position of a six membered sugar ring of a nucleoside of $[B_o]$).

[0184] In some instances, at least one of $[A_n]$ and $[B_o]$ includes the structure of Formula IV:



Formula IV

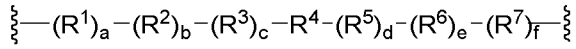
wherein each of N^1 and N^2 is independently a nucleobase;
each of R^9, R^{10}, R^{11}, R^{12}, R^{13}, R^{14}, R^{15}, and R^{16} is, independently, H, halo, hydroxy, thiol, optionally substituted C_1-C_6 alkyl, optionally substituted C_1-C_6 heteroalkyl, optionally substituted C_2-C_6 heteroalkenyl, optionally substituted C_2-C_6 heteroalkynyl, optionally substituted amino, azido, or optionally substituted C_6-C_{10} aryl;
each of g and h is, independently, 0 or 1;
each X^1 and X^4 is, independently, 0, NH, or S;
each X^2 is independently 0 or S; and
each X^3 is OH or SH, or a salt thereof.

[0185] In another aspect, the disclosure features a chimeric polynucleotide encoding a polypeptide, wherein the polynucleotide has a sequence including Formula II:

$[A_n]$-$L^1$$[B_o]$ **Formula II**

wherein each A and B is independently any nucleoside;
n and o are, independently 15 to 1000; and

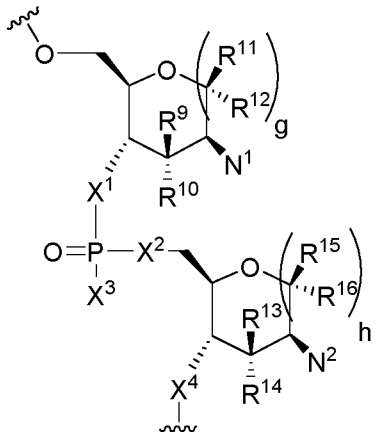
L^1 is a bond or has the structure of Formula III:



Formula III

wherein a, b, c, d, e, and f are each, independently, 0 or 1;
each of R^1, R^3, R^5, and R^7, is, independently, selected from optionally substituted C_1-C_6 alkylene, optionally substituted C_1-C_6 heteroalkylene, 0, S, and NR^8;

R^2 and R^6 are each, independently, selected from carbonyl, thiocarbonyl, sulfonyl, or phosphoryl;
R^4 is optionally substituted C_1-C_{10} alkylene, optionally substituted C_2-C_{10} alkenylene, optionally substituted C_2-C_{10} alkynylene, optionally substituted C_2-C_9 heterocyclylene, optionally substituted C_6-C_{12} arylene, optionally substituted C_2-C_{100} polyethylene glycolene, or optionally substituted C_1-C_{10} heteroalkylene, or a bond linking $(R^1)_a$-$(R^2)_b$-$(R^3)_c$ to $(R^5)_d$-$(R^6)_e$-$(R^7)_f$; and
R^8 is hydrogen, optionally substituted C_1-C_4 alkyl, optionally substituted C_2-C_4 alkenyl, optionally substituted C_2-C_4 alkynyl, optionally substituted C_2-C_6 heterocyclyl, optionally substituted C_6-C_{12} aryl, or optionally substituted C_1-C_7 heteroalkyl;
wherein L^1 is attached to $[A_n]$ and $[B_o]$ at the sugar of one of the nucleosides (e.g., at the 3' position of a five-membered sugar ring or 4' position of a six membered sugar ring of a nucleoside of $[A_n]$ and the 5' position of a five-membered sugar ring or 6' position of of a six membered sugar ring of a nucleoside of $[B_o]$ or at the 5' position of a five-membered sugar ring or 6' position of of a six membered sugar ring of a nucleoside of $[A_n]$ and the 3' position of a five-membered sugar ring or 4' position of a six membered sugar ring of a nucleoside of $[B_o]$).
wherein at least one of $[A_n]$ or $[B_o]$ includes the structure of Formula IV:



Formula IV

wherein each of N^1 and N^2 is independently a nucleobase;
each of R^9, R^{10}, R^{11}, R^{12}, R^{13}, R^{14}, R^{15}, and R^{16} is, independently, H, halo, hydroxy, thiol, optionally substituted C_1-C_6 alkyl, optionally substituted C_1-C_6 heteroalkyl, optionally substituted C_2-C_6 heteroalkenyl, optionally substituted C_2-C_6 heteroalkynyl, optionally substituted amino, azido, or optionally substituted C_6-C_{10} aryl;
each of g and h is, independently, 0 or 1;
each X^1 and X^4 is, independently, 0, NH, or S; and
each X^2 is independently 0 or S; and
each X^3 is OH or SH, or a salt thereof,
wherein at least one of X^1, X^2, or X^4 is NH or S.

[0186] In some instances, X^1 is NH. In other instances, X^4 is NH. In certain instances, X^2 is S.

[0187] In some instances, the polynucleotide includes: (a) a coding region; (b) a 5' UTR including at least one Kozak sequence, (c) a 3' UTR; and (d) at least one 5' cap structure. In other instances, the polynucleotide further includes (e) a poly-A tail.

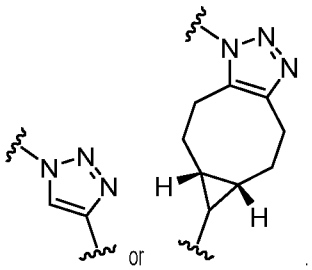
[0188] In some instances, one of the coding region, the 5' UTR including at least one Kozak sequence, the 3' UTR,

the 5' cap structure, or the poly-A tail includes $[A_n]$-L^1-$[B_o]$.

[0189] In other instances, one of the coding region, the 5' UTR including at least one Kozak sequence, the 3' UTR, the 5' cap structure, or the poly-A tail includes $[A_n]$ and another of the coding region, the 5' UTR including at least one Kozak sequence, the 3' UTR, the 5' cap structure, or the poly-A tail includes $[B_o]$.

[0190] In certain instances, the polynucleotide includes at least one modified nucleoside (e.g., a nucleoside of Table 2).

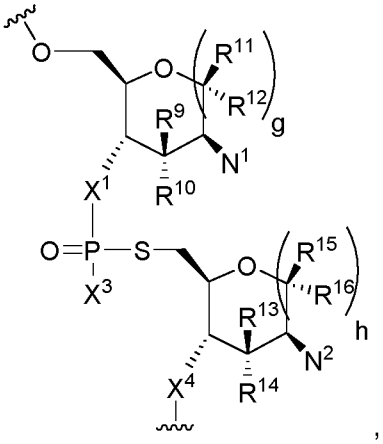
[0191] In some instances, R^4 is optionally substituted $C_{2\text{-}9}$ heterocyclylene, for example, the heterocycle may have the structure:



[0192] In certain instances, L^1 is attached to $[A_n]$ at the 3' position of a five-membered sugar ring or 4' position of a six membered sugar ring of one of the nucleosides and to $[B_o]$ at the 5' position of a five-membered sugar ring or 6' position of of a six membered sugar ring of one of the nucleosides.

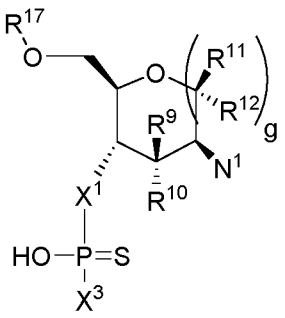
[0193] In some instances, the polynucleotide is circular.

[0194] In another aspect, the disclosure features a method of producing a composition including a chimeric polynucleotide encoding a polypeptide, wherein the polynucleotide includes the structure of Formula V:



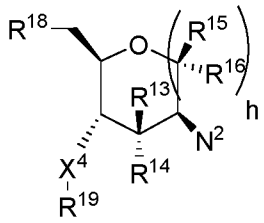
Formula V

[0195] This method includes reacting a compound having the structure of Formula VI:



Formula VI

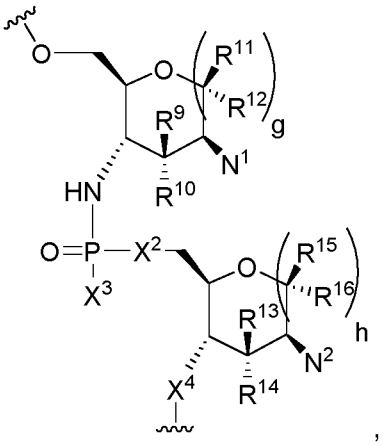
with a compound having the structure of Formula VII:



Formula VII

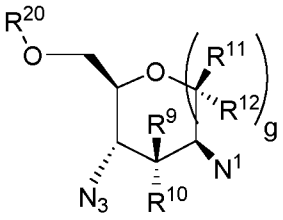
wherein each of N^1 and N^2 is, independently, a nucleobase;
each of R^9, R^{10}, R^{11}, R^{12}, R^{13}, R^{14}, R^{15}, and R^{16} is, independently, H, halo, hydroxy, thiol, optionally substituted C_1-C_6 alkyl, optionally substituted C_1-C_6 heteroalkyl, optionally substituted C_2-C_6 heteroalkenyl, optionally substituted C_2-C_6 heteroalkynyl, optionally substituted amino, azido, or optionally substituted C_6-C_{10} aryl;
each of g and h is, independently, 0 or 1;
each X^1 and X^4 is, independently, 0, NH, or S; and
each X^3 is independently OH or SH, or a salt thereof,
each of R^{17} and R^{19} is, independently, a region of linked nucleosides; and
R^{18} is a halogen.

[0196] In another aspect, the disclosure features a method of producing a composition including a chimeric polynucleotide encoding a polypeptide, wherein the polynucleotide includes the structure of Formula VIII:



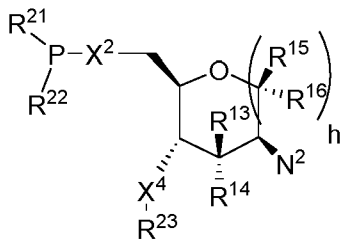
Formula VIII

[0197] This method includes reacting a compound having the structure of Formula IX:



Formula IX

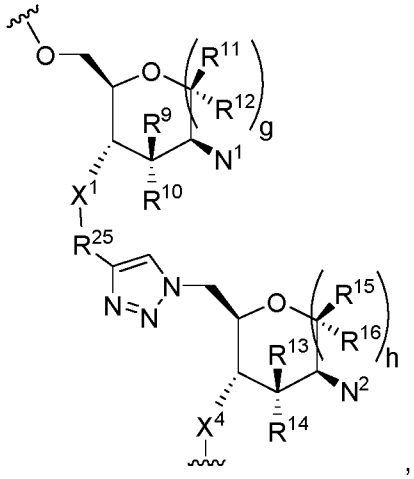
with a compound having the structure of Formula X:



Formula X

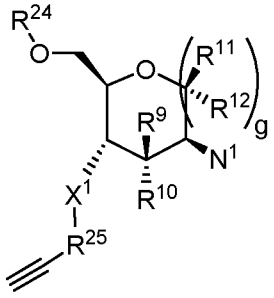
wherein each of N^1 and N^2 is, independently, a nucleobase;
each of R^9, R^{10}, R^{11}, R^{12}, R^{13}, R^{14}, R^{15}, and R^{16} is, independently, H, halo, hydroxy, thiol, optionally substituted C_1-C_6 alkyl, optionally substituted C_1-C_6 heteroalkyl, optionally substituted C_2-C_6 heteroalkenyl, optionally substituted C_2-C_6 heteroalkynyl, optionally substituted amino, azido, or optionally substituted C_6-C_{10} aryl;
each of g and h is, independently, 0 or 1;
each X^4 is, independently, O, NH, or S; and
each X^2 is independently O or S;
each X^3 is independently OH, SH, or a salt thereof,
each of R^{20} and R^{23} is, independently, a region of linked nucleosides; and
each of R^{21} and R^{22} is, independently, optionally substituted C_1-C_6 alkoxy.

[0198] In another aspect, the disclosure features a method of producing a composition including a chimeric polynucleotide encoding a polypeptide, wherei the polynucleotide includes the structure of Formula XI:



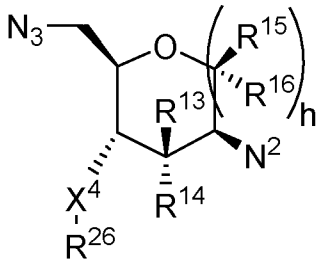
Formula XI

[0199] This method includes reacting a compound having the structure of Formula XII:



Formula XII

with a compound having the structure of Formula XIII:



Formula XIII

wherein each of N^1 and N^2 is, independently, a nucleobase;
each of R^9, R^{10}, R^{11}, R^{12}, R^{13}, R^{14}, R^{15}, and R^{16} is, independently, H, halo, hydroxy, thiol, optionally substituted C_1-C_6 alkyl, optionally substituted C_1-C_6 heteroalkyl, optionally substituted C_2-C_6 heteroalkenyl, optionally substituted C_2-C_6 heteroalkynyl, optionally substituted amino, azido, or optionally substituted C_6-C_{10} aryl;
each of g and h is, independently, 0 or 1;
each X^4 is, independently, 0, NH, or S; and
each X^2 is independently 0 or S;
each X^3 is independently OH, SH, or a salt thereof,
each of R^{24} and R^{26} is, independently, a region of linked nucleosides; and
R^{25} is optionally substituted C_1-C_6 alkylene or optionally substituted C_1-C_6 heteroalkylene or R^{25} and the alkynyl group together form optionally substituted cycloalkynyl.

[0200] In another aspect, the disclosure features a method of producing a composition including a chimeric polynucleotide encoding a polypeptide, wherein the polynucleotide has a sequence including Formula II:

$$[A_n]\text{-}L^1\text{-}[B_o], \qquad \textbf{Formula II}$$

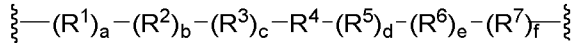
[0201] This method includes reacting a compound having the structure of Formula XIV

$$[A_n]\text{-}(R^1)_a\text{-}(R^2)_b\text{-}(R^3)_c\text{-}N_3 \qquad \textbf{Formula XIII}$$

with a compound having the structure of Formula XV:

$$R^{27}\text{-}(R^5)_d\text{-}(R^6)_e\text{-}(R^7)_f\text{-}[B_o] \qquad \textbf{Formula XII}$$

wherein each A and B is independently any nucleoside;
n and o are, independently 15 to 1000; and
L^1 has the structure of Formula III:

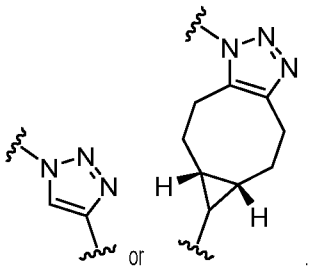


Formula III

wherein a, b, c, d, e, and f are each, independently, 0 or 1;
wherein each A and B is independently any nucleoside;
n and o are, independently 15 to 1000;
R^1, R^3, R^5, and R^7 each, independently, is selected from optionally substituted C_1-C_6 alkylene, optionally substituted C_1-C_6 heteroalkylene, 0, S, and NR^8;
R^2 and R^6 are each, independently, selected from carbonyl, thiocarbonyl, sulfonyl, or phosphoryl;
R^4 is an optionally substituted triazolene; and
R^8 is hydrogen, optionally substituted C_1-C_4 alkyl, optionally substituted C_3-C_4 alkenyl, optionally substituted C_2-C_4 alkynyl, optionally substituted C_2-C_6 heterocyclyl, optionally substituted C_6-C_{12} aryl, or optionally substituted C_1-C_7 heteroalkyl; and

R^{27} is an optionally substituted C_2-C_3 alkynyl or an optionally substituted C_8-C_{12} cycloalkynyl, wherein L^1 is attached to $[A_n]$ and $[B_o]$ at the sugar of one of the nucleosides.

[0202] In some instances, the optionally substituted triazolene has the structure:



[0203] In one instance, at least one of the regions of linked nucleosides of A may comprise a sequence of linked nucleosides which can function as a 5' untranslated region (UTR). The sequence of linked nucleosides may be a natural or synthetic 5' UTR. As a non-limiting example, the chimeric polynucleotide may encode a polypeptide of interest and the sequence of linked nucleosides of A may encode the native 5' UTR of a polypeptide encoded by the chimeric polynucleotide or the sequence of linked nucleosides may be a non-heterologous 5' UTR such as, but not limited to a synthetic UTR.

[0204] In another instance, at least one of the regions of linked nucleosides of A may be a cap region. The cap region may be located 5' to a region of linked nucleosides of A functioning as a 5'UTR. The cap region may comprise at least one cap such as, but not limited to, CapO, Cap1, ARCA, inosine, N1-methyl-guanosine, 2'fluoro-guanosine, 7-deaza-guanosine, 8-oxo-guanosine, 2-amino-guanosine, LNA-guanosine, 2-azido-guanosine, Cap2 and Cap4.

[0205] In one instance, at least one of the regions of linked nucleosdies of B may comprise at least one open reading frame of a nucleic acid sequence. The nucleic acid sequence may be codon optimized and/or comprise at least one modification.

[0206] In one instance, at least one of the regions of linked nucleosides of C may comprise a sequence of linked nucleosides which can function as a 3' UTR. The sequence of linked nucleosides may be a natural or synthetic 3' UTR. As a non-limiting example, the chimeric polynucleotide may encode a polypeptide of interest and the sequence of linked nucleosides of C may encode the native 3' UTR of a polypeptide encoded by the chimeric polynucleotide or the sequence of linked nucleosides may be a non-heterologous 3' UTR such as, but not limited to a synthetic UTR.

[0207] In one instance, at least one of the regions of linked nucleosides of A comprises a sequence of linked nucleosides which functions as a 5' UTR and at least one of the regions of linked nucleosides of C comprises a sequence of linked nucleosides which functions as a 3' UTR. In one instance, the 5' UTR and the 3' UTR may be from the same or different species. In another instance, the 5' UTR and the 3' UTR may encode the native untranslated regions from different proteins from the same or different species

[0208] Figures 4 and 5 provide schematics of a series of chimeric polynucleotides illustrating various patterns of positional modifications based on Formula I as well as those having a blocked or structured 3' terminus.

[0209] Chimeric polynucleotides, including the parts or regions thereof, of the present invention may be classified as hemimers, gapmers, wingmers, or blockmers.

[0210] As used herein, a "hemimer" is chimeric polynucleotide comprising a region or part which comprises half of one pattern, percent, position or population of a chemical modification(s) and half of a second pattern, percent, position or population of a chemical modification(s). Chimeric polynucleotides of the present invention may also comprise hemimer subregions. In one embodiment, a part or region is 50% of one and 50% of another

[0211] In one embodiment the entire chimeric polynucleotide can be 50% of one and 50% of the other Any region or part of any chimeric polynucleotide of the invention may be a hemimer. Types of hemimers include pattern hemimers, population hemimers or position hemimers. By definition, hemimers are 50:50 percent hemimers.

[0212] As used herein, a "gapmer" is a chimeric polynucleotide having at least three parts or regions with a gap between the parts or regions The "gap" can comprise a region of linked nucleosides or a single nucleoside which differs from the chimeric nature of the two parts or regions flanking it. The two parts or regions of a gapmer may be the same or different from each other.

[0213] As used herein, a "wingmer" is a chimeric polynucleotide having at least three parts or regions with a gap between the parts or regions Unlike a gapmer, the two flanking parts or regions surrounding the gap in a wingmer are the same in degree or kind. Such similarity may be in the length of number of units of different modifications or in the number of modifications. The wings of a wingmer may be longer or shorter than the gap. The wing parts or regions may be 20, 30, 40, 50, 60 70, 80, 90 or 95% greater or shorter in length than the region which comprises the gap.

[0214] As used herein, a "blockmer" is a patterned polynucleotide where parts or regions are of equivalent size or number and type of modifications. Regions or subregions in a blockmer may be 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61 62, 63, 64, 65, 66, 67, 68, 69, 70, 71, 72, 73, 74, 75,76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98,99,100,101,102,103,104,105,106,107,108,109,110,
111,112,113,114,115,116,117,118,119,120,121,122,123,124,125,126,127,128,129,130,131,132,133,134,135,136,137,138,139,140,
141,142,143,144,145,146,147,148,149,150,151,152,153,154,155,156,157,158,159,160,161,162,163,164,165,166,167,168,169,170,
171,172,173,174,175,176,177,178,179,180,181,182,183,184,185,186,187,188,189,190,191,192,193,194,195,196,197,198,199,200, 201, 202, 203, 204, 205, 206, 207, 208, 209, 210, 211, 212, 213, 214, 215, 216, 217, 218, 219, 220, 221, 222, 223, 224, 225, 226, 227, 228, 229, 230, 231,232,233,234,235,236,237,238,239,240,241,242,243,244,245,246,247,248,249,250,251,252,253,254,255,256,257,258,259,260, 261, 262, 263, 264, 265, 266, 267, 268, 269, 270, 271, 272, 273, 274, 275, 276, 277, 278, 279, 280, 281, 282, 283, 284, 285, 286, 287, 288, 289, 290, 291, 292, 293, 294, 295, 296, 297, 298, 299, 300, 310, 320, 330, 340, 350, 360, 370, 380, 390, 400, 410, 420, 430, 440, 450, 460, 470, 480, 490 or 500, nuclesides long.

[0215] Chimeric polynucleotides, including the parts or regions thereof, of the present invention having a chemical modification pattern are referred to as "pattern chimeras." Pattern chimeras may also be referred to as blockmers. Pattern chimeras are those polynucleotides having a pattern of modifications within, across or among regions or parts.

[0216] Patterns of modifications within a part or region are those which start and stop within a defined region. Patterns of modifcations across a part or region are those patterns which start in on part or region and end in another adjacent part or region. Patterns of modifications among parts or regions are those which begin and end in one part or region and are repeated in a different part or region, which is not necessarily adjacent to the first region or part.

[0217] The regions or subregions of pattern chimeras or blockmers may have simple alternating patterns such as ABAB[AB]n where each "A" and each "B" represent different chemical modifications (at at least one of the base, sugar or backbone linker), different types of chemical modifications (e.g., naturally occurring and non-naturally occurring), different percentages of modifications or different populations of modifications. The pattern may repeat n number of times where n=3-300. Further, each A or B can represent from 1-2500 units (e.g., nucleosides) in the pattern. Patterns may also be alternating multiples such as AABBAABB[AABB]n (an alternating double multiple) or AAABB-BAAABBB[AAABBB]n (an alternating triple multiple) pattern. The pattern may repeat n number of times where n=3-300.

[0218] Different patterns may also be mixed together to form a second order pattern. For example, a single alternating pattern may be combined with a triple alternating pattern to form a second order alternating pattern A'B'. One example would be [ABABAB][AAABBBAAABBB] [ABABAB][AAABBBAAABBB] [ABABAB][AAABBBAAABBB], where [ABABAB] is A' and [AAABBBAAABBB] is B'. In like fashion, these patterns may be repeated n number of times, where n=3-300.

[0219] Patterns may include three or more different modifications to form an ABCABC[ABC]n pattern. These three component patterns may also be multiples, such as AABBCCAABBCC[AABBCC]n and may be designed as combinations with other patterns such as ABCABCAABBCCABCABCAABBCC, and may be higher order patterns.

[0220] Regions or subregions of position, percent, and population modifications need not reflect an equal contribution from each modification type. They may form series such as "1-2-3-4", "1-2-4-8", where each integer represents the number of units of a particular modification type. Alternatively, they may be odd only, such as '1-3-3-1-3-1-5" or even only "2-4-2-4-6-4-8" or a mixuture of both odd and even number of units such as "1-3-4-2-5-7-3-3-4".

[0221] Pattern chimeras may vary in their chemical modification by degree (such as those described above) or by kind (e.g., different modifications).

[0222] Chimeric polynucleotides, including the parts or regions thereof, of the present disclosure having at least one region with two or more different chemical modifications of two or more nucleoside members of the same nucleoside type (A, C, G, T, or U) are referred to as "positionally modified" chimeras. Positionally modified chimeras are also referred to herein as "selective placement" chimeras or "selective placement polynucleotides". As the name implies, selective placement refers to the design of polynucleotides which, unlike polynucleotides in the art where the modification to any A, C, G, T or U is the same by virtue of the method of synthesis, can have different modifications to the individual As, Cs, Gs, Ts or Us in a polynucleotide or region thereof. For example, in a positionally modified chimeric polynucleotide, there may be two or more different chemical modifications to any of the nucleoside types of As, Cs, Gs, Ts, or Us. There may also be combinations of two or more to any two or more of the same nucleoside type. For example, a positionally modified or selective placement chimeric polynucleotide may comprise 3 different modifications to the population of adenines in the moleucle and also have 3 different modifications to the population of cytosines in the construct-all of which may have a unique, non-random, placement.

[0223] Chimeric polynucleotides, including the parts or regions thereof, of the present invention having a chemical modification percent are referred to as "percent chimeras." Percent chimeras may have regions or parts which comprise at least 1%, at least 2%, at least 5%, at least 8%, at least 10%, at least 20%, at least 30%, at least 40%, at least 50%,

at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, or at least 99% positional, pattern or population of modifications. Alternatively, the percent chimera may be completely modified as to modification position, pattern, or population. The percent of modification of a percent chimera may be split between naturally occurring and non-naturally occurring modifications.

[0224] Chimeric polynucleotides, including the parts or regions thereof, of the present invention having a chemical modification population are referred to as "population chimeras." A population chimera may comprise a region or part where nucleosides (their base, sugar or backbone linkage, or combination thereof) have a select population of modifications. Such modifications may be selected from functional populations such as modifications which induce, alter or modulate a phenotypic outcome. For example, a functional population may be a population or selection of chemical modifications which increase the level of a cytokine. Other functional populations may individually or collectively function to decrease the level of one or more cytokines. Use of a selection of these like-function modifications in a chimeric polynucleotide would therefore constitute a "functional population chimera." As used herein, a "functional population chimera" may be one whose unique functional feature is defined by the population of modifications as described above or the term may apply to the overall function of the chimeric polynucleotide itself. For example, as a whole the chimeric polynucleotide may function in a different or superior way as compared to an unmodified or non-chimeric polynucleotide.

[0225] It should be noted that polynucleotides which have a uniform chemical modification of all of any of the same nucleoside type or a population of modifications produced by mere downward titration of the same starting modification in all of any of the same nucleoside type, or a measured percent of a chemical modification of all any of the same nucleoside type but with random incorporation, such as where all uridines are replaced by a uridine analog, e.g., pseudouridine, are not considred chimeric. Likewise, polynucleotides having a uniform chemical modification of two, three, or four of the same nucleoside type throughout the entire polynucleotide (such as all uridines and all cytosines, etc. are modified in the same way) are not considered chimeric polynucleotides. One example of a polynucleotide which is not chimeric is the canonical pseudouridine/5-methyl cytosine modified polynucleotide of the prior art. These uniform polynucleotides are arrived at entirely via *in vitro* transcription (IVT) enzymatic synthesis; and due to the limitations of the synthesizing enzymes, they contain only one kind of modification at the occurrence of each of the same nucleoside type, i.e., adenosine (A), thymidine (T), guanosine (G), cytidine (C) or uradine (U), found in the polynucleotide. Such polynucleotides may be characterized as IVT polynucleotides.

[0226] The chimeric polynucleotides of the present invention are chemically modified. When the chimeric polynucleotides of the present invention are chemically and/or structurally modified the polynucleotides may be referred to as "modified chimeric polynucleotides."

[0227] In some embodiments of the invention, the chimeric polynucleotides may encode two or more peptides or polypeptides of interest. Such peptides or polypeptides of interest include the heavy and light chains of antibodies, an enzyme and its substrate, a label and its binding molecule, a second messenger and its enzyme or the components of multimeric proteins or complexes.

[0228] The regions or parts of the chimeric polynucleotides of the present invention may be separated by a linker or spacer moiety. Such linkers or spaces may be nucleic acid based or non-nucleosidic.

[0229] In one embodiment, the chimeric polynucleotides of the present invention may include a sequence encoding a self-cleaving peptide described herein, such as, but not limited to, a 2A peptide. The polynucleotide sequence of the 2A peptide in the chimeric polynucleotide may be modified or codon optimized by the methods described herein and/or are known in the art.

[0230] Notwithstanding the foregoing, the chimeric polynucleotides of the present invention may comprise a region or part which is not positionally modified or not chimeric as defined herein.

[0231] For example, a region or part of a chimeric polynucleotide may be uniformly modified at one ore more A, T, C, G, or U but according to the disclosure, the polynucleotides will not be uniformly modified throughout the entire region or part.

[0232] Regions or parts of chimeric polynucleotides may be from 15-1000 nucleosides in length and a polynucleotide may have from 2-100 different regions or patterns of regions as described herein.

[0233] In another embodiment, the chimeric polynucleotides have both coding and non-coding regions and parts.

[0234] Figure 2 illustrates the design of certain chimeric polynucleotides of the present invention when based on the scaffold of the polynucleotide of Figure 1. Shown in the figure are the regions or parts of the chimeric polynucleotides where patterned regions represent those regions which are positionally modified and open regions illustrate regions which may or may not be modified but which are, when modified, uniformly modified. Chimeric polynucleotides of the present invention may be completely positionally modified or partially positionally modified. They may also have subregions which may be of any pattern or design. Shown in Figure 2 are a chimeric subregion and a hemimer subregion.

[0235] In one embodiment, the shortest length of a region of the chimeric polynucleotide of the present invention encoding a peptide can be the length that is sufficient to encode for a dipeptide, a tripeptide, a tetrapeptide, a pentapeptide, a hexapeptide, a heptapeptide, an octapeptide, a nonapeptide, or a decapeptide. In another embodiment, the length may be sufficient to encode a peptide of 2-30 amino acids, e.g. 5-30, 10-30, 2-25, 5-25, 10-25, or 10-20 amino acids.

The length may be sufficient to encode for a peptide of at least 11, 12, 13, 14, 15, 17, 20, 25 or 30 amino acids, or a peptide that is no longer than 40 amino acids, e.g. no longer than 35, 30, 25, 20, 17, 15, 14, 13, 12, 11 or 10 amino acids. Examples of dipeptides that the polynucleotide sequences can encode or include, but are not limited to, carnosine and anserine.

[0236] In one embodiment, the length of a region of the chimeric polynucleotide of the present invention encoding the peptide or polypeptide of interest is greater than about 30 nucleotides in length (e.g., at least or greater than about 35, 40, 45, 50, 55, 60, 70, 80, 90, 100, 120, 140, 160, 180, 200, 250,300,350,400,450,500,600,700,800,900,1,000, 1,100, 1,200, 1,300, 1,400, 1,500, 1,600, 1,700, 1,800, 1,900, 2,000, 2,500, and 3,000, 4,000, 5,000, 6,000, 7,000, 8,000, 9,000, 10,000, 20,000, 30,000, 40,000, 50,000, 60,000, 70,000, 80,000, 90,000 or up to and including 100,000 nucleotides). As used herein, such a region may be referred to as a "coding region" or "region encoding."

[0237] In some embodiments, the chimeric polynucleotide includes from about 30 to about 100,000 nucleotides (e.g., from 30 to 50, from 30 to 100, from 30 to 250, from 30 to 500, from 30 to 1,000, from 30 to 1,500, from 30 to 3,000, from 30 to 5,000, from 30 to 7,000, from 30 to 10,000, from 30 to 25,000, from 30 to 50,000, from 30 to 70,000, from 100 to 250, from 100 to 500, from 100 to 1,000, from 100 to 1,500, from 100 to 3,000, from 100 to 5,000, from 100 to 7,000, from 100 to 10,000, from 100 to 25,000, from 100 to 50,000, from 100 to 70,000, from 100 to 100,000, from 500 to 1,000, from 500 to 1,500, from 500 to 2,000, from 500 to 3,000, from 500 to 5,000, from 500 to 7,000, from 500 to 10,000, from 500 to 25,000, from 500 to 50,000, from 500 to 70,000, from 500 to 100,000, from 1,000 to 1,500, from 1,000 to 2,000, from 1,000 to 3,000, from 1,000 to 5,000, from 1,000 to 7,000, from 1,000 to 10,000, from 1,000 to 25,000, from 1,000 to 50,000, from 1,000 to 70,000, from 1,000 to 100,000, from 1,500 to 3,000, from 1,500 to 5,000, from 1,500 to 7,000, from 1,500 to 10,000, from 1,500 to 25,000, from 1,500 to 50,000, from 1,500 to 70,000, from 1,500 to 100,000, from 2,000 to 3,000, from 2,000 to 5,000, from 2,000 to 7,000, from 2,000 to 10,000, from 2,000 to 25,000, from 2,000 to 50,000, from 2,000 to 70,000, and from 2,000 to 100,000).

[0238] According to the present invention, regions or subregions of the chimeric polynucleotides may also range independently from 15-1,000 nucleotides in length (e.g., greater than 30, 40, 45, 50, 55, 60, 70, 80, 90, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 225, 250, 275, 300, 325, 350, 375, 400, 425, 450, 475, 500, 550, 600, 650, 700, 750, 800, 850, 900 and 950 nucleotides or at least 30, 40, 45, 50, 55, 60, 70, 80, 90, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 225, 250, 275, 300, 325, 350, 375, 400, 425, 450, 475, 500, 550, 600, 650, 700, 750, 800, 850, 900, 950 and 1,000 nucleotides).

[0239] According to the present invention, regions or subregions of chimeric polynucleotides may range from absent to 500 nucleotides in length (e.g., at least 60, 70, 80, 90, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 250, 300, 350, 400, 450, or 500 nucleotides). Where the region is a polyA tail, the length may be determined in units of or as a function of polyA Binding Protein binding. In this embodiment, the polyA tail is long enough to bind at least 4 monomers of PolyA Binding Protein. PolyA Binding Protein monomers bind to stretches of approximately 38 nucleotides As such, it has been observed that polyA tails of about 80 nucleotides to about 160 nucleotides (SEQ ID NO: 2278) are functional. The chimeric polynucleotides of the present invention which function as an mRNA need not comprise a polyA tail.

[0240] According to the present invention, chimeric polynucleotides which function as an mRNA may have a capping region. The capping region may comprise a single cap or a series of nucleotides forming the cap. In this embodiment the capping region may be from 1 to 10, e.g. 2-9, 3-8, 4-7, 1-5, 5-10, or at least 2, or 10 or fewer nucleotides in length In some embodiments, the cap is absent.

[0241] The present invention contemplates chimeric polynucleotides which are circular or cyclic. As the name implies circular polynucleotides are circular in nature meaning that the termini are joined in some fashion, whether by ligation, covalent bond, common association with the same protein or other molecule or complex or by hybridization. Any of the cicular polynucleotides as taught in for example U.S. Provisional Application Number 61/873,010 filed September 3,2013 may be made chimeric according to the present invention.

[0242] Chimeric polynucleotides, formulations and compositions comprising chimeric polynucleotides, and methods of making, using and administering chimeric polynucleotides are also described in co-pending US Provisional Application No 61/873,034, filed September 3, 2013, entitled Chimeric Polynucleotides, and US Provisional Application No. 61/877,582, filed September 13, 2013, entitled Chimeric Polynucleotides.

Circular Polynculeotide Architecture

[0243] The present invention contemplates polynucleotides which are circular or cyclic. As the name implies circular polynucleotides are circular in nature meaning that the termini are joined in some fashion, whether by ligation, covalent bond, common association with the same protein or other molecule or complex or by hybridization. Any of the cicular polynucleotides as taught in for example U.S. Provisional Application Number 61/873,010 filed September 3, 2013.

[0244] Circular polynucleotides of the present invention may be designed according to the circular RNA construct scaffolds shown in Figures 6-12. Such polynucleotides are cicular polynucleotides or circular constructs.

[0245] The circular polynucleotides or circPs of the present invention which encode at least one peptide or polypeptide

of interest are known as circular RNAs or circRNA. The antigens of the NAVs of the present invention may be encoded by one or more circular RNAs or circRNAs.

[0246] As used herein, "circular RNA" or "circRNA" means a circular polynucleotide that can encode at least one peptide or polypeptide of interest

[0247] As used herein, "sensor sequence" means a receptor or pseudo-receptor for endogenous nucleic acid binding molecules. Non-limiting examples of sensor sequences include, microRNA binding sites, microRNA seed sequences, microRNA binding sites without the seed sequence, transcription factor binding sites and artificial binding sites engineered to act as pseudo-receptors and portions and fragments thereof.

[0248] The circPs of the present invention which comprise at least one sensor sequence and encode at least one peptide or polypeptide of interest are known as circular RNA sponges or circRNA-SP. As used herein, "circular RNA sponges" or "circRNA-SP" means a circular polynucleotide which comprises at least one sensor sequence and at least one region encoding at least one peptide or polypeptide of interest.

[0249] Figure **6A** shows a representative circular construct **200** of the circular polynucleotides of the present invention. As used herein, the term "circular construct" refers to a circular polynucleotide transcript which may act substantiatlly similar to and have properties of a RNA molecule. In one embodiment the circular construct acts as an mRNA. If the circular construct encodes one or more peptides or polypeptides of interest (e.g., a circRNA or circRNA-SP) then the polynucleotide transcript retains sufficient structural and/or chemical features to allow the polypeptide of interest encoded therein to be translated. Circular constructs may be polynucleotides of the invention When structurally or chemically modified, the construct may be referred to as a modified circP, modified circSP, modified circRNA or modified circRNA-SP.

[0250] Returning to FIG. **6A,** the circular construct **200** here contains a first region of linked nucleotides **202** that is flanked by a first flanking region **204** and a second flanking region **206.** As used herein, the "first region" may be referred to as a "coding region," a "non-coding region" or "region encoding" or simply the "first region." In one embodiment, this first region may comprise nucleotides such as, but is not limited to, encoding at leaset one peptide or polypeptide of interest and/or nucleotides encoding a sensor region. The peptide or polypeptide of interest may comprise at its 5' terminus one or more signal peptide sequences encoded by a signal peptide sequence region **203.** The first flanking region **204** may comprise a region of linked nucleosides or portion thereof which may act similiarly to an untranslated region (UTR) in an mRNA and/or DNA sequence. The first flanking region may also comprise a region of polarity **208.** The region of polarity **208** may include an IRES sequence or portion thereof. As a non-limiting example, when linearlized this region may be split to have a first portion be on the 5' terminus of the first region **202** and second portion be on the 3' terminus of the first region **202.** The second flanking region **206** may comprise a tailing sequence region **210** and may comprise a region of linked nucleotides or portion thereof **212** which may act similiarly to a UTR in an mRNA and/or DNA.

[0251] Bridging the 5' terminus of the first region **202** and the first flanking region **104** is a first operational region **206.** In one embodiment, this operational region may comprise a start codon. The operational region may alternatively comprise any translation initiation sequence or signal including a start codon.

[0252] Bridging the 3' terminus of the first region **202** and the second flanking region **106** is a second operational region **207.** Traditionally this operational region comprises a stop codon. The operational region may alternatively comprise any translation initiation sequence or signal including a stop codon. According to the present invention, multiple serial stop codons may also be used. In one embodiment, the operation region of the present invention may comprise two stop codons. The first stop codon may be "TGA" or "UGA" and the second stop codon may be selected from the group consisting of "TAA," "TGA," "TAG," "UAA," "UGA" or"UAG."

[0253] Turning to Figure **6B,** at least one non-nucleic acid moiety **201** may be used to prepare a circular construct **200** where the non-nucleic acid moiety **201** is used to bring the first flanking region **204** near the second flanking region **206.** Non-limiting examples of non-nucleic acid moieties which may be used in the present invention are described herein. The circular construct **200** may comprise more than one non-nucleic acid moiety wherein the additional non-nucleic acid moeities may be heterologous or homologous to the first non-nucleic acid moiety.

[0254] Turning to Figure **7A,** the first region of linked nucleosides **202** may comprise a spacer region **214.** This spacer region **214** may be used to separate the first region of linked nucleosides **202** so that the circular construct can include more than one open reading frame, non-coding region or an open reading frame and a non-coding region.

[0255] Turning to Figure **7B,** the second flanking region **206** may comprise one or more sensor regions **216** in the the 3'UTR **212.** These sensor sequences as discussed herein operate as pseudo-receptors (or binding sites) for ligands of the local microenvironment of the circular construct. For example, microRNA bindng sites or miRNA seeds may be used as sensors such that they function as pseudoreceptors for any microRNAs present in the environment of the circular polynucleotide. As shown in Figure **9,** the one or more sensor regions **216** may be separated by a spacer region **214.**

[0256] As shown in Figure **8A,** a circular construct **200,** which includes one or more sensor regions **216,** may also include a spacer region **214** in the first region of linked nucleosides **202.** As discussed above for Figure **7,** this spacer region **214** may be used to separate the first region of linked nucleosides **202** so that the circular construct can include more than one open reading frame and/or more than one non-coding region.

[0257] Turning to Figure **8B,** a circular construct **200** may be a non-coding construct known as a circSP comprising

at least one non-coding region such as, but not limited to, a sensor region **216.** Each of the sensor regions **216** may include, but are not limited to, a miR sequence, a miR seed, a miR binding site and/or a miR sequence without the seed.

[0258] Turning to Figure **9,** at least one non-nucleic acid moiety **201** may be used to prepare a circular construct **200** which is a non-coding construct. The circular construct **200** which is a non-coding construct may comprise more than one non-nucleic acid moiety wherein the additional non-nucleic acid moeities may be heterologous or homologous to the first non-nucleic acid moiety.

[0259] Circular polynucleotides, formulations and compositions comprising circular polynucleotides, and methods of making, using and administering circular polynucleotides are also described in co-pending US Provisional Application No 61/873,010, filed September 3, 2013, entitled Circular Polynucleotides, and US Provisional Application No. 61/877,527, filed September 13, 2013, entitled Circular Polynucleotides.

Multimers of Polynucleotides

[0260] According to the present invention, multiple distinct chimeric polynucleotides and/or IVT polynucleotides may be linked together through the 3'-end using nucleotides which are modified at the 3'-terminus. Chemical conjugation may be used to control the stoichiometry of delivery into cells. For example, the glyoxylate cycle enzymes, isocitrate lyase and malate synthase, may be supplied into cells at a 1:1 ratio to alter cellular fatty acid metabolism. This ratio may be controlled by chemically linking chimeric polynucleotides and/or IVT polynucleotides using a 3'-azido terminated nucleotide on one polynucleotides species and a C5-ethynyl or alkynyl-containing nucleotide on the opposite polynucleotide species. The modified nucleotide is added post-transcriptionally using terminal transferase (New England Biolabs, Ipswich, MA) according to the manufacturer's protocol. After the addition of the 3'-modified nucleotide, the two polynucleotides species may be combined in an aqueous solution, in the presence or absence of copper, to form a new covalent linkage via a click chemistry mechanism as described in the literature.

[0261] In another example, more than two chimeric polynucleotides and/or IVT polynucleotides may be linked together using a functionalized linker molecule. For example, a functionalized saccharide molecule may be chemically modified to contain multiple chemical reactive groups (SH-, NH_2-, N_3, etc...) to react with the cognate moiety on a 3'-functionalized mRNA molecule (i.e., a 3'-maleimide ester, 3'-NHS-ester, alkynyl). The number of reactive groups on the modified saccharide can be controlled in a stoichiometric fashion to directly control the stoichiometric ratio of conjugated chimeric polynucleotides and/or IVT polynucleotides.

[0262] In one embodiment, the chimeric polynucleotides and/or IVT polynucleotides may be linked together in a pattern. The pattern may be a simple alternating pattern such as $CD[CD]_x$ where each "C" and each "D" represent a chimeric polynucleotide, IVT polynucleotide, different chimeric polynucleotides or different IVT polynucleotides. The pattern may repeat x number of times, where x= 1-300. It is to be understood that the antigens of the NAVs of the present invention may be encoded by such linked polynucleotides, as described herein. Paterns may also be alternating multiples such as $CCDD[CCDD]_x$ (an alternating double multiple) or $CCCDDD[CCCDDD]_x$ (an alternating triple multiple) pattern. The alternating double multiple or alternating triple multiple may repeat x number of times, where x= 1-300.

Conjugates and Combinations of Polynucleotides

[0263] In order to further enhance protein production, polynucleotides of the present invention can be designed to be conjugated to other polynucleotides, dyes, intercalating agents (*e.g.* acridines), cross-linkers (*e.g.* psoralene, mitomycin C), porphyrins (TPPC4, texaphyrin, Sapphyrin), polycyclic aromatic hydrocarbons (*e.g.*, phenazine, dihydrophenazine), artificial endonucleases (*e.g.* EDTA), alkylating agents, phosphate, amino, mercapto, PEG (*e.g.*, PEG-40K), MPEG, $[MPEG]_2$, polyamino, alkyl, substituted alkyl, radiolabeled markers, enzymes, haptens (*e.g.* biotin), transport/absorption facilitators (*e.g.*, aspirin, vitamin E, folic acid), synthetic ribonucleases, proteins, e.g., glycoproteins, or peptides, e.g., molecules having a specific affinity for a co-ligand, or antibodies *e.g.*, an antibody, that binds to a specified cell type such as a cancer cell, endothelial cell, or bone cell, hormones and hormone receptors, non-peptidic species, such as lipids, lectins, carbohydrates, vitamins, cofactors, or a drug.

[0264] In a preferred embodiment, the polynucleotides of the present invention which encode an antigen are conjugated to one or more dendritic cell markers.

[0265] Conjugation may result in increased stability and/or half life and may be particularly useful in targeting the polynucleotides to specific sites in the cell, tissue or organism.

[0266] According to the present invention, the polynucleotides may be administered with, conjugated to or further encode one or more of RNAi agents, siRNAs, shRNAs, miRNAs, miRNA binding sites, antisense RNAs, ribozymes, catalytic DNA, tRNA, RNAs that induce triple helix formation, aptamers or vectors, and the like.

Bifunctional Polynucleotides

[0267] In one embodiment of the invention NAVs may comprise bifunctional polynucleotides (e.g., bifunctional IVT polynucleotides, bifunctional chimeric polynucleotides or bifunctional circular polynucleotides). As the name implies, bifunctional polynucleotides are those having or capable of at least two functions. These molecules may also by convention be referred to as multi-functional. It is to be understood that the NAV polynucleotides of the present invention may be conjugated to other molecules or agents, as described *supra*.

[0268] The multiple functionalities of bifunctional polynucleotides may be encoded by the NAV (the function may not manifest until the encoded product is translated) or may be a property of the polynucleotide itself. It may be structural or chemical. Bifunctional modified polynucleotides may comprise a function that is covalently or electrostatically associated with the polynucleotides. Further, the two functions may be provided in the context of a complex of a chimeric polynucleotide and another molecule.

Noncoding Polynucleotides

[0269] As described herein, provided are polynucleotides having sequences that are partially or substantially not translatable, e.g., having a noncoding region. As one non-limiting example, the noncoding region may be the first region of the IVT polynucleotide or the circular polynucleotide. Alternatively, the noncoding region may be a region other than the first region. As another non-limiting example, the noncoding region may be the A, B and/or C region of the chimeric polynucleotide.

[0270] Such molecules are generally not translated, but can exert an effect on the immune response or protein production by one or more of binding to and sequestering one or more translational machinery components such as a ribosomal protein or a transfer RNA (tRNA), thereby effectively reducing protein expression in the cell or modulating one or more pathways or cascades in a cell which in turn alters protein levels. The polynucleotide may contain or encode one or more long noncoding RNA (lncRNA, or lincRNA) or portion thereof, a small nucleolar RNA (sno-RNA), micro RNA (miRNA), small interfering RNA (siRNA) or Piwi-interacting RNA (piRNA). Examples of such lncRNA molecules and RNAi constructs designed to target such lncRNA any of which may be encoded in the polynucleotides are taught in International Publication, WO2012/018881 A2.

[0271] According to the present invention, the polynucleotide encodes one or more polypeptides of interest or fragments thereof. Such polypeptide of interest may include, but is not limited to, whole polypeptides, a plurality of polypeptides or fragments of polypeptides, which independently may be encoded by one or more regions or parts or the whole of a polynucleotide. As used herein, the term "polypeptides of interest" refer to any polypeptide which is selected to be encoded within, or whose function is affected by, the polnucleotides of the present invention. The peptides or polypetides of the invention are antigenic.

[0272] As used herein, "polypeptide" means a polymer of amino acid residues (natural or unnatural) linked together most often by peptide bonds. The term, as used herein, refers to proteins, polypeptides, and peptides of any size, structure, or function. The polypeptides of interest are antigens encoded by the polynucleotides as described herein.

[0273] In some instances the polypeptide encoded is smaller than about 50 amino acids and the polypeptide is then termed a peptide. If the polypeptide is a peptide, it will be at least about 2, 3, 4, or at least 5 amino acid residues long. Thus, polypeptides include gene products, naturally occurring polypeptides, synthetic polypeptides, homologs, orthologs, paralogs, fragments and other equivalents, variants, and analogs of the foregoing. A polypeptide may be a single molecule or may be a multi-molecular complex such as a dimer, trimer or tetramer. They may also comprise single chain or multichain polypeptides such as antibodies or insulin and may be associated or linked. Most commonly disulfide linkages are found in multichain polypeptides. The term polypeptide may also apply to amino acid polymers in which one or more amino acid residues are an artificial chemical analogue of a corresponding naturally occurring amino acid.

[0274] The term "polypeptide variant" refers to molecules which differ in their amino acid sequence from a native or reference sequence. The amino acid sequence variants may possess substitutions, deletions, and/or insertions at certain positions within the amino acid sequence, as compared to a native or reference sequence. Ordinarily, variants will possess at least about 50% identity (homology) to a native or reference sequence, and preferably, they will be at least about 80%, more preferably at least about 90% identical (homologous) to a native or reference sequence.

[0275] In some embodiments "variant mimics" are provided. As used herein, the term "variant mimic" is one which contains one or more amino acids which would mimic an activated sequence. For example, glutamate may serve as a mimic for phosphoro-threonine and/or phosphoro-serine. Alternatively, variant mimics may result in deactivation or in an inactivated product containing the mimic, e.g., phenylalanine may act as an inactivating substitution for tyrosine; or alanine may act as an inactivating substitution for serine.

[0276] "Homology" as it applies to amino acid sequences is defined as the percentage of residues in the candidate amino acid sequence that are identical with the residues in the amino acid sequence of a second sequence after aligning the sequences and introducing gaps, if necessary, to achieve the maximum percent homology. Methods and computer

programs for the alignment are well known in the art. It is understood that homology depends on a calculation of percent identity but may differ in value due to gaps and penalties introduced in the calculation.

[0277] By "homologs" as it applies to polypeptide sequences means the corresponding sequence of other species having substantial identity to a second sequence of a second species

[0278] "Analogs" is meant to include polypeptide variants which differ by one or more amino acid alterations, e.g., substitutions, additions or deletions of amino acid residues that still maintain one or more of the properties of the parent or starting polypeptide.

[0279] The present invention contemplates several types of compositions which are polypeptide based including variants and derivatives These include substitutional, insertional, deletion and covalent variants and derivatives The term "derivative" is used synonymously with the term "variant" but generally refers to a molecule that has been modified and/or changed in any way relative to a reference molecule or starting molecule.

[0280] As such, polynucleotides encoding peptides or polypeptides containing substitutions, insertions and/or additions, deletions and covalent modifications with respect to reference sequences, in particular the polypeptide sequences disclosed herein, are included within the scope of this invention. For example, sequence tags or amino acids, such as one or more lysines, can be added to the peptide sequences of the invention (e.g., at the N-terminal or C-terminal ends). Sequence tags can be used for peptide purification or localization Lysines can be used to increase peptide solubility or to allow for biotinylation. Alternatively, amino acid residues located at the carboxy and amino terminal regions of the amino acid sequence of a peptide or protein may optionally be deleted providing for truncated sequences Certain amino acids (e.g., C-terminal or N-terminal residues) may alternatively be deleted depending on the use of the sequence, as for example, expression of the sequence as part of a larger sequence which is soluble, or linked to a solid support.

[0281] "Substitutional variants" when referring to polypeptides are those that have at least one amino acid residue in a native or starting sequence removed and a different amino acid inserted in its place at the same position. The substitutions may be single, where only one amino acid in the molecule has been substituted, or they may be multiple, where two or more amino acids have been substituted in the same molecule.

[0282] As used herein the term "conservative amino acid substitution" refers to the substitution of an amino acid that is normally present in the sequence with a different amino acid of similar size, charge, or polarity. Examples of conservative substitutions include the substitution of a non-polar (hydrophobic) residue such as isoleucine, valine and leucine for another non-polar residue. Likewise, examples of conservative substitutions include the substitution of one polar (hydrophilic) residue for another such as between arginine and lysine, between glutamine and asparagine, and between glycine and serine. Additionally, the substitution of a basic residue such as lysine, arginine or histidine for another, or the substitution of one acidic residue such as aspartic acid or glutamic acid for another acidic residue are additional examples of conservative substitutions Examples of non-conservative substitutions include the substitution of a non-polar (hydrophobic) amino acid residue such as isoleucine, valine, leucine, alanine, methionine for a polar (hydrophilic) residue such as cysteine, glutamine, glutamic acid or lysine and/or a polar residue for a non-polar residue.

[0283] "Insertional variants" when referring to polypeptides are those with one or more amino acids inserted immediately adjacent to an amino acid at a particular position in a native or starting sequence. "Immediately adjacent" to an amino acid means connected to either the alpha-carboxy or alpha-amino functional group of the amino acid.

[0284] "Deletional variants" when referring to polypeptides are those with one or more amino acids in the native or starting amino acid sequence removed. Ordinarily, deletional variants will have one or more amino acids deleted in a particular region of the molecule.

[0285] "Covalent derivatives" when referring to polypeptides include modifications of a native or starting protein with an organic proteinaceous or non-proteinaceous derivatizing agent, and/or post-translational modifications. Covalent modifications are traditionally introduced by reacting targeted amino acid residues of the protein with an organic derivatizing agent that is capable of reacting with selected side-chains or terminal residues, or by harnessing mechanisms of post-translational modifications that function in selected recombinant host cells. The resultant covalent derivatives are useful in programs directed at identifying residues important for biological activity, for immunoassays, or for the preparation of anti-protein antibodies for immunoaffinity purification of the recombinant glycoprotein. Such modifications are within the ordinary skill in the art and are performed without undue experimentation.

[0286] Certain post-translational modifications are the result of the action of recombinant host cells on the expressed polypeptide. Glutaminyl and asparaginyl residues are frequently post-translationally deamidated to the corresponding glutamyl and aspartyl residues. Alternatively, these residues are deamidated under mildly acidic conditions. Either form of these residues may be present in the polypeptides produced in accordance with the present invention.

[0287] Other post-translational modifications include hydroxylation of proline and lysine, phosphorylation of hydroxyl groups of seryl or threonyl residues, methylation of the alpha-amino groups of lysine, arginine, and histidine side chains (T. E. Creighton, Proteins: Structure and Molecular Properties, W.H. Freeman & Co., San Francisco, pp. 79-86 (1983)).

[0288] "Features" when referring to polypeptides are defined as distinct amino acid sequence-based components of a molecule. Features of the polypeptides encoded by the polynucleotides of the present invention include surface manifestations, local conformational shape, folds, loops, half-loops, domains, half-domains, sites, termini or any com-

bination thereof.

[0289] As used herein when referring to polypeptides the term "surface manifestation" refers to a polypeptide based component of a protein appearing on an outermost surface.

[0290] As used herein when referring to polypeptides the term "local conformational shape" means a polypeptide based structural manifestation of a protein which is located within a definable space of the protein.

[0291] As used herein when referring to polypeptides the term "fold" refers to the resultant conformation of an amino acid sequence upon energy minimization. A fold may occur at the secondary or tertiary level of the folding process. Examples of secondary level folds include beta sheets and alpha helices. Examples of tertiary folds include domains and regions formed due to aggregation or separation of energetic forces. Regions formed in this way include hydrophobic and hydrophilic pockets, and the like.

[0292] As used herein the term "turn" as it relates to protein conformation means a bend which alters the direction of the backbone of a peptide or polypeptide and may involve one, two, three or more amino acid residues.

[0293] As used herein when referring to polypeptides the term "loop" refers to a structural feature of a polypeptide which may serve to reverse the direction of the backbone of a peptide or polypeptide Where the loop is found in a polypeptide and only alters the direction of the backbone, it may comprise four or more amino acid residues. Oliva et al. have identified at least 5 classes of protein loops (J. Mol Biol 266 (4): 814-830; 1997). Loops may be open or closed. Closed loops or "cyclic" loops may comprise 2, 3, 4, 5, 6, 7, 8, 9, 10 or more amino acids between the bridging moieties. Such bridging moieties may comprise a cysteine-cysteine bridge (Cys-Cys) typical in polypeptides having disulfide bridges or alternatively bridging moieties may be non-protein based such as the dibromozylyl agents used herein.

[0294] As used herein when referring to polypeptides the term "half-loop" refers to a portion of an identified loop having at least half the number of amino acid resides as the loop from which it is derived. It is understood that loops may not always contain an even number of amino acid residues. Therefore, in those cases where a loop contains or is identified to comprise an odd number of amino acids, a half-loop of the odd-numbered loop will comprise the whole number portion or next whole number portion of the loop (number of amino acids of the loop/2+/-0.5 amino acids). For example, a loop identified as a 7 amino acid loop could produce half-loops of 3 amino acids or 4 amino acids (7/2=3.5+/-0.5 being 3 or 4).

[0295] As used herein when referring to polypeptides the term "domain" refers to a motif of a polypeptide having one or more identifiable structural or functional characteristics or properties (e.g., binding capacity, serving as a site for protein-protein interactions).

[0296] As used herein when referring to polypeptides the term "half-domain" means a portion of an identified domain having at least half the number of amino acid resides as the domain from which it is derived. It is understood that domains may not always contain an even number of amino acid residues. Therefore, in those cases where a domain contains or is identified to comprise an odd number of amino acids, a half-domain of the odd-numbered domain will comprise the whole number portion or next whole number portion of the domain (number of amino acids of the domain/2+/-0.5 amino acids). For example, a domain identified as a 7 amino acid domain could produce half-domains of 3 amino acids or 4 amino acids (7/2=3.5+/-0.5 being 3 or 4). It is also understood that sub-domains may be identified within domains or half-domains, these subdomains possessing less than all of the structural or functional properties identified in the domains or half domains from which they were derived. It is also understood that the amino acids that comprise any of the domain types herein need not be contiguous along the backbone of the polypeptide (i.e., nonadjacent amino acids may fold structurally to produce a domain, half-domain or subdomain).

[0297] As used herein when referring to polypeptides the terms "site" as it pertains to amino acid based embodiments is used synonymously with "amino acid residue" and "amino acid side chain." A site represents a position within a peptide or polypeptide that may be modified, manipulated, altered, derivatized or varied within the polypeptide based molecules of the present invention.

[0298] As used herein the terms "termini" or "terminus" when referring to polypeptides refers to an extremity of a peptide or polypeptide. Such extremity is not limited only to the first or final site of the peptide or polypeptide but may include additional amino acids in the terminal regions. The polypeptide based molecules of the present invention may be characterized as having both an N-terminus (terminated by an amino acid with a free amino group (NH2)) and a C-terminus (terminated by an amino acid with a free carboxyl group (COOH)). Proteins of the invention are in some cases made up of multiple polypeptide chains brought together by disulfide bonds or by non-covalent forces (multimers, oligomers). These sorts of proteins will have multiple N- and C-termini. Alternatively, the termini of the polypeptides may be modified such that they begin or end, as the case may be, with a non-polypeptide based moiety such as an organic conjugate.

[0299] Once any of the features have been identified or defined as a desired component of a polypeptide to be encoded by a polynucleotide of the invention, any of several manipulations and/or modifications of these features may be performed by moving, swapping, inverting, deleting, randomizing or duplicating. Furthermore, it is understood that manipulation of features may result in the same outcome as a modification to the molecules of the invention. For example, a manipulation which involved deleting a domain would result in the alteration of the length of a molecule just as modification of a nucleic acid to encode less than a full length molecule would.

[0300] Modifications and manipulations can be accomplished by methods known in the art such as, but not limited to, site directed mutagenesis or *a priori* incorporation during chemical synthesis. The resulting modified molecules may then be tested for activity using *in vitro* or *in vivo* assays such as those described herein or any other suitable screening assay known in the art.

[0301] According to the present invention, the polypeptides may comprise a consensus sequence which is discovered through rounds of experimentation. As used herein a "consensus" sequence is a single sequence which represents a collective population of sequences allowing for variability at one or more sites.

[0302] As recognized by those skilled in the art, protein fragments, functional protein domains, and homologous proteins are also considered to be within the scope of polypeptides of interest of this invention. For example, provided herein is any protein fragment (meaning a polypeptide sequence at least one amino acid residue shorter than a reference polypeptide sequence but otherwise identical) of a reference protein 10, 20, 30, 40, 50, 60, 70, 80, 90,100 or greater than 100 amino acids in length In another example, any protein that includes a stretch of about 20, about 30, about 40, about 50, or about 100 amino acids which are about 40%, about 50%, about 60%, about 70%, about 80%, about 90%, about 95%, or about 100% identical to any of the sequences described herein can be utilized in accordance with the invention. In certain embodiments, a polypeptide to be utilized in accordance with the invention includes 2, 3, 4, 5, 6, 7, 8, 9, 10, or more mutations as shown in any of the sequences provided or referenced herein.

[0303] Reference molecules (polypeptides or polynucleotides) may share a certain identity with the designed molecules (polypeptides or polynucleotides). The term "identity" as known in the art, refers to a relationship between the sequences of two or more peptides, polypeptides or polynucleotides, as determined by comparing the sequences. In the art, identity also means the degree of sequence relatedness between them as determined by the number of matches between strings of two or more amino acid residues or nucleosides. Identity measures the percent of identical matches between the smaller of two or more sequences with gap alignments (if any) addressed by a particular mathematical model or computer program (i.e., "algorithms"). Identity of related peptides can be readily calculated by known methods. Such methods include, but are not limited to, those described in Computational Molecular Biology, Lesk, A. M., ed., Oxford University Press, New York, 1988; Biocomputing: Informatics and Genome Projects, Smith, D. W., ed., Academic Press, New York, 1993; Computer Analysis of Sequence Data, Part 1, Griffin, A. M., and Griffin, H. G., eds, Humana Press, New Jersey, 1994; Sequence Analysis in Molecular Biology, von Heinje, G., Academic Press, 1987; Sequence Analysis Primer, Gribskov, M. and Devereux, J., eds, M. Stockton Press, New York, 1991; and Carillo et al., SIAM J. Applied Math. 48,1073 (1988).

[0304] In some embodiments, the encoded polypeptide variant may have the same or a similar activity as the reference polypeptide. Alternatively, the variant may have an altered activity (e.g., increased or decreased) relative to a reference polypeptide. Generally, variants of a particular polynucleotide or polypeptide of the invention will have at least about 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% but less than 100% sequence identity to that particular reference polynucleotide or polypeptide as determined by sequence alignment programs and parameters described herein and known to those skilled in the art. Such tools for alignment include those of the BLAST suite (Stephen F. Altschul, Thomas L. Madden, Alejandro A. Schaffer, Jinghui Zhang, Zheng Zhang, Webb Miller, and David J. Lipman (1997), "Gapped BLAST and PSI-BLAST: a new generation of protein database search programs", Nucleic Acids Res. 25:3389-3402.) Other tools are described herein, specifically in the definition of "Identity."

[0305] Default parameters in the BLAST algorithm include, for example, an expect threshold of 10, Word size of 28, Match/Mismatch Scores 1, -2, Gap costs Linear. Any filter can be applied as well as a selection for species specific repeats, e.g., Homo sapiens.

Cell-Penetrating Polypeptides

[0306] The polynucleotides disclosed herein (*e.g.*, antigen-encoding polynucleotides featured in the NAVs of the invention), may also encode one or more cell-penetrating polypeptides. As used herein, "cell-penetrating polypeptide" or CPP refers to a polypeptide which may facilitate the cellular uptake of molecules. A cell-penetrating polypeptide of the present invention may contain one or more detectable labels. The polypeptides may be partially labeled or completely labeled throughout. The polynucleotides may encode the detectable label completely, partially or not at all. The cell-penetrating peptide may also include a signal sequence. As used herein, a "signal sequence" refers to a sequence of amino acid residues bound at the amino terminus of a nascent protein during protein translation. The signal sequence may be used to signal the secretion of the cell-penetrating polypeptide.

[0307] In one embodiment, the polynucleotides may also encode a fusion protein. The fusion protein may be created by operably linking a charged protein to a therapeutic protein. As used herein, "operably linked" refers to the therapeutic protein and the charged protein being connected in such a way to permit the expression of the complex when introduced into the cell. As used herein, "charged protein" refers to a protein that carries a positive, negative or overall neutral electrical charge. Preferably, the therapeutic protein may be covalently linked to the charged protein in the formation of

the fusion protein. The ratio of surface charge to total or surface amino acids may be approximately 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8 or 0.9.

[0308] The cell-penetrating polypeptide encoded by the polynucleotides may form a complex after being translated. The complex may comprise a charged protein linked, e.g. covalently linked, to the cell-penetrating polypeptide. "Therapeutic protein" refers to a protein that, when administered to a cell has a therapeutic, diagnostic, and/or prophylactic effect and/or elicits a desired biological and/or pharmacological effect.

[0309] In one embodiment, the cell-penetrating polypeptide may comprise a first domain and a second domain. The first domain may comprise a supercharged polypeptide. The second domain may comprise a protein-binding partner. As used herein, "protein-binding partner" includes, but is not limited to, antibodies and functional fragments thereof, scaffold proteins, or peptides. The cell-penetrating polypeptide may further comprise an intracellular binding partner for the protein-binding partner. The cell-penetrating polypeptide may be capable of being secreted from a cell where the polynucleotides may be introduced. The cell-penetrating polypeptide may also be capable of penetrating the first cell.

[0310] In a further embodiment, the cell-penetrating polypeptide is capable of penetrating a second cell. The second cell may be from the same area as the first cell, or it may be from a different area. The area may include, but is not limited to, tissues and organs. The second cell may also be proximal or distal to the first cell.

[0311] In one embodiment, the polynucleotides may encode a cell-penetrating polypeptide which may comprise a protein-binding partner. The protein binding partner may include, but is not limited to, an antibody, a supercharged antibody or a functional fragment. The polynucleotides may be introduced into the cell where a cell-penetrating polypeptide comprising the protein-binding partner is introduced.

Anti-Microbial and Anti-viral Polypeptides

[0312] The polynucleotides of the present invention (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention) may be designed to be co-administered with, a polynucleotide encoding one or more antimicrobial peptides (AMP) or antiviral peptides (AVP). AMPs and AVPs have been isolated and described from a wide range of animals such as, but not limited to, microorganisms, invertebrates, plants, amphibians, birds, fish, and mammals (Wang et al., Nucleic Acids Res. 2009; 37 (Database issue):D933-7). For example, anti-microbial polypeptides are described in Antimicrobial Peptide Database (http://aps.unmc.edu/AP/main.php; Wang et al., Nucleic Acids Res. 2009; 37 (Database issue):D933-7), CAMP: Collection of Anti-Microbial Peptides (http://www.bicnirrh.res.in/antimicrobial/); Thomas et al., Nucleic Acids Res. 2010, 38 (Database issue):D774-80), US 5221732, US 5447914, US 5519115, US 5607914, US 5714577, US 5734015, US 5798336, US 5821224, US 5849490, US 5856127, US 5905187, US 5994308, US 5998374, US 6107460, US 6191254, US 6211148, US 6300489, US 6329504, US 6399370, US 6476189, US 6478825, US 6492328, US 6514701, US 6573361, US 6573361, US 6576755, US 6605698, US 6624140, US 6638531, US 6642203, US 6653280, US 6696238, US 6727066, US 6730659, US 6743598, US 6743769, US 6747007, US 6790833, US 6794490, US 6818407, US 6835536, US 6835713, US 6838435, US 6872705, US 6875907, US 6884776, US 6887847, US 6906035, US 6911524, US 6936432, US 7001924, US 7071293, US 7078380, US 7091185, US 7094759, US 7166769, US 7244710, US 7314858, and US 7582301.

[0313] The anti-microbial polypeptides described herein may block cell fusion and/or viral entry by one or more enveloped viruses (*e.g.*, HIV, HCV). For example, the anti-microbial polypeptide can comprise or consist of a synthetic peptide corresponding to a region, *e.g.*, a consecutive sequence of at least about 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, or 60 amino acids of the transmembrane subunit of a viral envelope protein, *e.g.*, HIV-1 gp120 or gp41. The amino acid and nucleotide sequences of HIV-1 gp120 or gp41 are described in, *e.g.*, Kuiken *et al.,* (2008). *"HIV Sequence Compendium,"* Los Alamos National Laboratory.

[0314] In some embodiments, the anti-microbial polypeptide may have at least about 75%, 80%, 85%, 90%, 95%, 100% sequence homology to the corresponding viral protein sequence. In some embodiments, the anti-microbial polypeptide may have at least about 75%, 80%, 85%, 90%, 95%, or 100% sequence homology to the corresponding viral protein sequence.

[0315] In other embodiments, the anti-microbial polypeptide may comprise or consist of a synthetic peptide corresponding to a region, *e.g.*, a consecutive sequence of at least about 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, or 60 amino acids of the binding domain of a capsid binding protein. In some embodiments, the anti-microbial polypeptide may have at least about 75%, 80%, 85%, 90%, 95%, or 100% sequence homology to the corresponding sequence of the capsid binding protein.

[0316] The anti-microbial polypeptides described herein may block protease dimerization and inhibit cleavage of viral proproteins (e.g., HIV Gag-pol processing) into functional proteins thereby preventing release of one or more enveloped viruses (*e.g.*, HIV, HCV). In some embodiments, the anti-microbial polypeptide may have at least about 75%, 80%, 85%, 90%, 95%, 100% sequence homology to the corresponding viral protein sequence.

[0317] In other embodiments, the anti-microbial polypeptide can comprise or consist of a synthetic peptide corresponding to a region, *e.g.*, a consecutive sequence of at least about 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, or 60 amino acids

of the binding domain of a protease binding protein. In some embodiments, the anti-microbial polypeptide may have at least about 75%, 80%, 85%, 90%, 95%, 100% sequence homology to the corresponding sequence of the protease binding protein.

[0318] The anti-microbial polypeptides described herein can include an *in vitro*-evolved polypeptide directed against a viral pathogen.

Anti-Microbial Polypeptides

[0319] Anti-microbial polypeptides (AMPs) are small peptides of variable length, sequence and structure with broad spectrum activity against a wide range of microorganisms including, but not limited to, bacteria, viruses, fungi, protozoa, parasites, prions, and tumor/cancer cells. (See, *e.g.*, Zaiou, J Mol Med, 2007; 85:317). It has been shown that AMPs have broad-spectrum of rapid onset of killing activities, with potentially low levels of induced resistance and concomitant broad anti-inflammatory effects.

[0320] In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may be under 10kDa, *e.g.*, under 8kDa, 6kDa, 4kDa, 2kDa, or 1kDa. In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) consists of from about 6 to about 100 amino acids, *e.g.*, from about 6 to about 75 amino acids, about 6 to about 50 amino acids, about 6 to about 25 amino acids, about 25 to about 100 amino acids, about 50 to about 100 amino acids, or about 75 to about 100 amino acids. In certain embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may consist of from about 15 to about 45 amino acids. In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) is substantially cationic.

[0321] In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may be substantially amphipathic. In certain embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may be substantially cationic and amphipathic. In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may be cytostatic to a Gram-positive bacterium. In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may be cytotoxic to a Gram-positive bacterium. In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may be cytostatic and cytotoxic to a Gram-positive bacterium. In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may be cytostatic to a Gram-negative bacterium. In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may be cytotoxic to a Gram-negative bacterium. In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may be cytostatic and cytotoxic to a Gram-positive bacterium. In some embodiments, the anti-microbial polypeptide may be cytostatic to a virus, fungus, protozoan, parasite, prion, or a combination thereof. In some embodiments, the anti-microbial polypeptide may be cytotoxic to a virus, fungus, protozoan, parasite, prion, or a combination thereof. In certain embodiments, the anti-microbial polypeptide may be cytostatic and cytotoxic to a virus, fungus, protozoan, parasite, prion, or a combination thereof. In some embodiments, the anti-microbial polypeptide may be cytotoxic to a tumor or cancer cell (*e.g.*, a human tumor and/or cancer cell). In some embodiments, the anti-microbial polypeptide may be cytostatic to a tumor or cancer cell (*e.g.*, a human tumor and/or cancer cell). In certain embodiments, the anti-microbial polypeptide may be cytotoxic and cytostatic to a tumor or cancer cell (*e.g.*, a human tumor or cancer cell). In some embodiments, the anti-microbial polypeptide (*e.g.*, an anti-bacterial polypeptide) may be a secreted polypeptide.

[0322] In some embodiments, the anti-microbial polypeptide comprises or consists of a defensin. Exemplary defensins include, but are not limited to, α-defensins (*e.g.*, neutrophil defensin 1, defensin alpha 1, neutrophil defensin 3, neutrophil defensin 4, defensin 5, defensin 6), β-defensins (*e.g.*, beta-defensin 1, beta-defensin 2, beta-defensin 103, beta-defensin 107, beta-defensin 110, beta-defensin 136), and θ-defensins. In other embodiments, the anti-microbial polypeptide comprises or consists of a cathelicidin (*e.g.*, hCAP18).

Anti-Viral Polypeptides

[0323] Anti-viral polypeptides (AVPs) are small peptides of variable length, sequence and structure with broad spectrum activity against a wide range of viruses. See, *e.g.*, Zaiou, J Mol Med, 2007; 85:317. It has been shown that AVPs have a broad-spectrum of rapid onset of killing activities, with potentially low levels of induced resistance and concomitant broad anti-inflammatory effects. In some embodiments, the anti-viral polypeptide is under 10kDa, *e.g.*, under 8kDa, 6kDa, 4kDa, 2kDa, or 1kDa. In some embodiments, the anti-viral polypeptide comprises or consists of from about 6 to about 100 amino acids, *e.g.*, from about 6 to about 75 amino acids, about 6 to about 50 amino acids, about 6 to about 25 amino acids, about 25 to about 100 amino acids, about 50 to about 100 amino acids, or about 75 to about 100 amino acids. In certain embodiments, the anti-viral polypeptide comprises or consists of from about 15 to about 45 amino acids. In some embodiments, the anti-viral polypeptide is substantially cationic. In some embodiments, the anti-viral polypeptide is substantially amphipathic. In certain embodiments, the anti-viral polypeptide is substantially cationic and amphipathic. In some embodiments, the anti-viral polypeptide is cytostatic to a virus. In some embodiments, the anti-viral polypeptide is cytotoxic to a virus. In some embodiments, the anti-viral polypeptide is cytostatic and cytotoxic to a virus. In some

embodiments, the anti-viral polypeptide is cytostatic to a bacterium, fungus, protozoan, parasite, prion, or a combination thereof. In some embodiments, the anti-viral polypeptide is cytotoxic to a bacterium, fungus, protozoan, parasite, prion or a combination thereof. In certain embodiments, the anti-viral polypeptide is cytostatic and cytotoxic to a bacterium, fungus, protozoan, parasite, prion, or a combination thereof. In some embodiments, the anti-viral polypeptide is cytotoxic to a tumor or cancer cell (*e.g.*, a human cancer cell). In some embodiments, the anti-viral polypeptide is cytostatic to a tumor or cancer cell (*e.g.*, a human cancer cell). In certain embodiments, the anti-viral polypeptide is cytotoxic and cytostatic to a tumor or cancer cell (*e.g.*, a human cancer cell). In some embodiments, the anti-viral polypeptide is a secreted polypeptide.

Cytotoxic Nucleosides

[0324] In one embodiment, the polynucleotides of the present invention (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention) may incorporate one or more cytotoxic nucleosides. For example, cytotoxic nucleosides may be incorporated into polynucleotides such as bifunctional modified RNAs or mRNAs. Cytotoxic nucleoside anti-cancer agents include, but are not limited to, adenosine arabinoside, cytarabine, cytosine arabinoside, 5-fluorouracil, fludarabine, floxuridine, FTORAFUR® (a combination of tegafur and uracil), tegafur ((RS)-5-fluoro-1-(tetrahydrofuran-2-yl)pyrimidine-2,4(1H,3H)-dione), and 6-mercaptopurine.

[0325] A number of cytotoxic nucleoside analogues are in clinical use, or have been the subject of clinical trials, as anticancer agents. Examples of such analogues include, but are not limited to, cytarabine, gemcitabine, troxacitabine, decitabine, tezacitabine, 2'-deoxy-2'-methylidenecytidine (DMDC), cladribine, clofarabine, 5-azacytidine, 4'-thio-aracytidine, cyclopentenylcytosine and 1-(2-C-cyano-2-deoxy-beta-D-arabino-pentofuranosyl)-cytosine. Another example of such a compound is fludarabine phosphate. These compounds may be administered systemically and may have side effects which are typical of cytotoxic agents such as, but not limited to, little or no specificity for tumor cells over proliferating normal cells.

[0326] A number of prodrugs of cytotoxic nucleoside analogues are also reported in the art. Examples include, but are not limited to, N4-behenoyl-1-beta-D-arabinofuranosylcytosine, N4-octadecyl-1-beta-D-arabinofuranosylcytosine, N4-palmitoyl-1-(2-C-cyano-2-deoxy-beta-D-arabino-pentofuranosyl) cytosine, and P-4055 (cytarabine 5'-elaidic acid ester). In general, these prodrugs may be converted into the active drugs mainly in the liver and systemic circulation and display little or no selective release of active drug in the tumor tissue. For example, capecitabine, a prodrug of 5'-deoxy-5-fluorocytidine (and eventually of 5-fluorouracil), is metabolized both in the liver and in the tumor tissue. A series of capecitabine analogues containing "an easily hydrolysable radical under physiological conditions" has been claimed by Fujiu et al. (U.S. Pat. No. 4,966,891). The series described by Fujiu includes N4 alkyl and aralkyl carbamates of 5'-deoxy-5-fluorocytidine and the implication that these compounds will be activated by hydrolysis under normal physiological conditions to provide 5'-deoxy-5-fluorocytidine.

[0327] A series of cytarabine N4-carbamates has been by reported by Fadl et al (Pharmazie. 1995, 50, 382-7) in which compounds were designed to convert into cytarabine in the liver and plasma. WO 2004/041203 discloses prodrugs of gemcitabine, where some of the prodrugs are N4-carbamates. These compounds were designed to overcome the gastrointestinal toxicity of gemcitabine and were intended to provide gemcitabine by hydrolytic release in the liver and plasma after absorption of the intact prodrug from the gastrointestinal tract. Nomura et al (Bioorg Med. Chem. 2003, 11, 2453-61) have described acetal derivatives of 1-(3-C-efhynyl-β-D-ribo-penfofaranosyl) cytosine which, on bioreduction, produced an intermediate that required further hydrolysis under acidic conditions to produce a cytotoxic nucleoside compound.

[0328] Cytotoxic nucleotides which may be chemotherapeutic also include, but are not limited to, pyrazolo [3,4-D]-pyrimidines, allopurinol, azathioprine, capecitabine, cytosine arabinoside, fluorouracil, mercaptopurine, 6-thioguanine, acyclovir, ara-adenosine, ribavirin, 7-deaza-adenosine, 7-deaza-guanosine, 6-aza-uracil, 6-aza-cytidine, thymidine ribonucleotide, 5-bromodeoxyuridine, 2-chloro-purine, and inosine, or combinations thereof.

Polynucleotides having Untranslated Regions (UTRs)

[0329] The polynucleotides of the present invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) may comprise one or more regions or parts which act or function as an untranslated region. Where polynucleotides are designed to encode at least one polypeptide of interest, the polynucleotides may comprise one or more of these untranslated regions

[0330] By definition, wild type untranslated regions (UTRs) of a gene are transcribed but not translated. In mRNA, the 5'UTR starts at the transcription start site and continues to the start codon but does not include the start codon, whereas, the 3'UTR starts immediately following the stop codon and continues until the transcriptional termination signal. There is growing body of evidence about the regulatory roles played by the UTRs in terms of stability of the nucleic acid molecule and translation. The regulatory features of a UTR can be incorporated into the polynucleotides of the present invention

to, among other things, enhance the stability of the molecule. The specific features can also be incorporated to ensure controlled down-regulation of the transcript in case they are misdirected to undesired organs sites.

[0331] Tables 19 and 20 provide a listing of exemplary UTRs which may be utilized in the polynucleotides of the present invention. Shown in Table 19 is a listing of a 5'-untranslated region of the invention. Variants of 5' UTRs may be utilized wherein one or more nucleotides are added or removed to the termini, including A, T, C or G.

Table 19. 5'-Untranslated Regions

5' UTR Identifier	Name/ Description	Sequence	SEQ ID NO.
5UTR-001	Upstream UTR	GGGAAATAAGAGAGAAAAGAAGAGTAAGAAGAAATATAAGAGCCAC C	2007
5UTR-002	Upstream UTR	GGGAGATCAGAGAGAAAAGAAGAGTAAGAAGAAATATAAGAGCCAC C	2008
5UTR-003	Upstream UTR	GGAATAAAAGTCTCAACACAACATATACAAAACAAACGAATCTCAAG CAATCAAGCATTCTACTTCTATTGCAGCAATTTAAATCATTTCTTTTAA AGCAAAAGCAATTTTCTGAAAATTTTCACCATTTACGAACGATAGCA AC	2009
5UTR-004	Upstream UTR	GGGAGACAAGCUUGGCAUUCCGGUACUGUUGGUAAAGCCACC	2010
5UTR-005	Upstream UTR	GGGAGATCAGAGAGAAAAGAAGAGTAAGAAGAAATATAAGAGCCAC C	2011
5UTR-006	Upstream UTR	GGAATAAAAGTCTCAACACAACATATACAAAACAAACGAATCTCAAG CAATCAAGCATTCTACTTCTATTGCAGCAATTTAAATCATTTCTTTTAA AGCAAAAGCAATTTTCTGAAAATTTTCACCATTTACGAACGATAGCA AC	2012
5UTR-007	Upstream UTR	GGGAGACAAGCUUGGCAUUCCGGUACUGUUGGUAAAGCCACC	2013
5UTR-008	Upstream UTR	GGGAATTAACAGAGAAAAGAAGAGTAAGAAGAAATATAAGAGCCAC C	2014
5UTR-009	Upstream UTR	GGGAAATTAGACAGAAAAGAAGAGTAAGAAGAAATATAAGAGCCAC C	2015
5UTR-010	Upstream UTR	GGGAAATAAGAGAGTAAAGAACAGTAAGAAGAAATATAAGAGCCAC C	2016
5UTR-011	Upstream UTR	GGGAAAAAAGAGAGAAAAGAAGACTAAGAAGAAATATAAGAGCCAC C	2017
5UTR-012	Upstream UTR	GGGAAATAAGAGAGAAAAGAAGAGTAAGAAGATATATAAGAGCCAC C	2018
5UTR-013	Upstream UTR	GGGAAATAAGAGACAAAACAAGAGTAAGAAGAAATATAAGAGCCAC C	2019
5UTR-014	Upstream UTR	GGGAAATTAGAGAGTAAAGAACAGTAAGTAGAATTAAAAGAGCCAC C	2020
5UTR-015	Upstream UTR	GGGAAATAAGAGAGAATAGAAGAGTAAGAAGAAATATAAGAGCCAC C	2021
5UTR-016	Upstream UTR	GGGAAATAAGAGAGAAAAGAAGAGTAAGAAGAAAATTAAGAGCCAC C	2022
5UTR-017	Upstream UTR	GGGAAATAAGAGAGAAAAGAAGAGTAAGAAGAAATTTAAGAGCCAC C	2023

[0332] Shown in Table 20 is a listing of 3'-untranslated regions of the invention. Variants of 3' UTRs may be utilized wherein one or more nucleotides are added or removed to the termini, including A, T, C or G.

Table 20. 3'-Untranslated Regions

3' UTR Identifier	Name/ Description	Sequence	SEQ ID NO.
3UTR-001	Creatine Kinase	GCGCCTGCCCACCTGCCACCGACTGCTGGAACCCAGCCAGTGGGAGGGCCTG GCCCACCAGAGTCCTGCTCCCTCACTCCTCGCCCCGCCCCCTGTCCCAGAGTC CCACCTGGGGGCTCTCTCCACCCTTCTCAGAGTTCCAGTTTCAACCAGAGTTCC AACCAATGGGCTCCATCCTCTGGATTCTGGCCAATGAAATATCTCCCTGGCAGG GTCCTCTTCTTTTCCCAGAGCTCCACCCCAACCAGGAGCTCTAGTTAATGGAGA GCTCCCAGCACACTCGGAGCTTGTGCTTTGTCTCCACGCAAAGCGATAAATAAA AGCATTGGTGGCCTTTGGTCTTTGAATAAAGCCTGAGTAGGAAGTCTAGA	2024
3UTR-002	Myoglobin	GCCCCTGCCGCTCCCACCCCCACCCATCTGGGCCCCGGGTTCAAGAGAGAGC GGGGTCTGATCTCGTGTAGCCATATAGAGTTTGCTTCTGAGTGTCTGCTTTGTT TAGTAGAGGTGGGCAGGAGGAGCTGAGGGGCTGGGGCTGGGGTGTTGAAGTT GGCTTTGCATGCCCAGCGATGCGCCTCCCTGTGGGATGTCATCACCCTGGGAA CCGGGAGTGGCCCTTGGCTCACTGTGTTCTGCATGGTTTGGATCTGAATTAATT GTCCTTTCTTCTAAATCCCAACCGAACTTCTTCCAACCTCCAAACTGGCTGTAAC CCCAAATCCAAGCCATTAACTACACCTGACAGTAGCAATTGTCTGATTAATCACT GGCCCCTTGAAGACAGCAGAATGTCCCTTTGCAATGAGGAGGAGATCTGGGCT GGGCGGGCCAGCTGGGGAAGCATTTGACTATCTGGAACTTGTGTGTGCCTCCT CAGGTATGGCAGTGACTCACCTGGTTTTAATAAAACAACCTGCAACATCTCATG GTCTTTGAATAAAGCCTGAGTAGGAAGTCTAGA	2025
3UTR-003	α-actin	ACACACTCCACCTCCAGCACGCGACTTCTCAGGACGACGAATCTTCTCAATGG GGGGGCGGCTGAGCTCCAGCCACCCCGCAGTCACTTTCTTTGTAACAACTTCC GTTGCTGCCATCGTAAACTGACACAGTGTTTATAACGTGTACATACATTAACTTA TTACCTCATTTTGTTATTTTTCGAAACAAAGCCCTGTGGAAGAAAATGGAAAACT TGAAGAAGCATTAAAGTCATTCTGTTAAGCTGCGTAAATGGTCTTTGAATAAAGC CTGAGTAGGAAGTCTAGA	2026
3UTR-004	Albumin	CATCACATTTAAAAGCATCTCAGCCTACCATGAGAATAAGAGAAAGAAAATGAA GATCAAAAGCTTATTCATCTGTTTTTCTTTTTCGTTGGTGTAAAGCCAACACCCT GTCTAAAAAACATAAATTTCTTTAATCATTTTGCCTCTTTTCTCTGTGCTTCAATT AATAAAAAATGGAAAGAATCTAATAGAGTGGTACAGCACTGTTATTTTTCAAAGA TGTGTTGCTATCCTGAAAATTCTGTAGGTTCTGTGGAAGTTCCAGTGTTCTCTCT TATTCCACTTCGGTAGAGGATTTCTAGTTTCTTGTGGGCTAATTAAATAAATCAT TAATACTCTTCTAATGGTCTTTGAATAAAGCCTGAGTAGGAAGTCTAGA	2027
3UTR-005	α-globin	GCTGCCTTCTGCGGGGCTTGCCTTCTGGCCATGCCCTTCTTCTCTCCCTTGCA CCTGTACCTCTTGGTCTTTGAATAAAGCCTGAGTAGGAAGGCGGCCGCTCGAG	2028

		CATGCATCTAGA	
3UTR-006	G-CSF	GCCAAGCCCTCCCCATCCCATGTATTTATCTCTATTTAATATTTATGTCTATTTAA GCCTCATATTTAAAGACAGGGAAGAGCAGAACGGAGCCCCAGGCCTCTGTGTC CTTCCCTGCATTTCTGAGTTTCATTCTCCTGCCTGTAGCAGTGAGAAAAAGCTC CTGTCCTCCCATCCCCTGGACTGGGAGGTAGATAGGTAAATACCAAGTATTTAT TACTATGACTGCTCCCCAGCCCTGGCTCTGCAATGGGCACTGGGATGAGCCGC TGTGAGCCCCTGGTCCTGAGGGTCCCCACCTGGGACCCTTGAGAGTATCAGGT CTCCCACGTGGGAGACAAGAAATCCCTGTTTAATATTTAAACAGCAGTGTTCCC CATCTGGGTCCTTGCACCCCTCACTCTGGCCTCAGCCGACTGCACAGCGGCCC CTGCATCCCCTTGGCTGTGAGGCCCCTGGACAAGCAGAGGTGGCCAGAGCTG GGAGGCATGGCCCTGGGGTCCCACGAATTTGCTGGGGAATCTCGTTTTTCTTC TTAAGACTTTTGGGACATGGTTTGACTCCCGAACATCACCGACGCGTCTCCTGT TTTTCTGGGTGGCCTCGGGACACCTGCCCTGCCCCCACGAGGGTCAGGACTG TGACTCTTTTTAGGGCCAGGCAGGTGCCTGGACATTTGCCTTGCTGGACGGGG ACTGGGGATGTGGGAGGGAGCAGACAGGAGGAATCATGTCAGGCCTGTGTGT GAAAGGAAGCTCCACTGTCACCCTCCACCTCTTCACCCCCCACTCACCAGTGT CCCCTCCACTGTCACATTGTAACTGAACTTCAGGATAATAAAGTGTTTGCCTCC ATGGTCTTTGAATAAAGCCTGAGTAGGAAGGCGGCCGCTCGAGCATGCATCTA GA	2029
3UTR-007	Col1a2; collagen, type I, alpha 2	ACTCAATCTAAATTAAAAAAGAAAGAAATTTGAAAAAACTTTCTCTTTGCCATTTC TTCTTCTTCTTTTTTAACTGAAAGCTGAATCCTTCCATTTCTTCTGCACATCTACT TGCTTAAATTGTGGGCAAAAGAGAAAAAGAAGGATTGATCAGAGCATTGTGCAA TACAGTTTCATTAACTCCTTCCCCCGCTCCCCCAAAAATTTGAATTTTTTTTTCAA CACTCTTACACCTGTTATGGAAAATGTCAACCTTTGTAAGAAAACCAAAATAAAA ATTGAAAAATAAAAACCATAAACATTTGCACCACTTGTGGCTTTTGAATATCTTC CACAGAGGGAAGTTTAAAACCCAAACTTCCAAAGGTTTAAACTACCTCAAAACA CTTTCCCATGAGTGTGATCCACATTGTTAGGTGCTGACCTAGACAGAGATGAAC TGAGGTCCTTGTTTTGTTTTGTTCATAATACAAAGGTGCTAATTAATAGTATTTCA GATACTTGAAGAATGTTGATGGTGCTAGAAGAATTTGAGAAGAAATACTCCTGT ATTGAGTTGTATCGTGTGGTGTATTTTTTAAAAAATTTGATTTAGCATTCATATTT TCCATCTTATTCCCAATTAAAAGTATGCAGATTATTTGCCCAAATCTTCTTCAGAT TCAGCATTTGTTCTTTGCCAGTCTCATTTTCATCTTCTTCCATGGTTCCACAGAA GCTTTGTTTCTTGGGCAAGCAGAAAAATTAAATTGTACCTATTTTGTATATGTGA GATGTTTAAATAAATTGTGAAAAAAATGAAATAAAGCATGTTTGGTTTTCCAAAA GAACATAT	2030
3UTR-008	Col6a2; collagen, type VI, alpha 2	CGCCGCCGCCCGGGCCCCGCAGTCGAGGGTCGTGAGCCCACCCCGTCCATG GTGCTAAGCGGGCCCGGGTCCCACACGGCCAGCACCGCTGCTCACTCGGACG ACGCCCTGGGCCTGCACCTCTCCAGCTCCTCCCACGGGGTCCCCGTAGCCCC GGCCCCCGCCCAGCCCCAGGTCTCCCCAGGCCCTCCGCAGGCTGCCCGGCC TCCCTCCCCCTGCAGCCATCCCAAGGCTCCTGACCTACCTGGCCCCTGAGCTC TGGAGCAAGCCCTGACCCAATAAAGGCTTTGAACCCAT	2031
3UTR-009	RPN1; ribophorin I	GGGGCTAGAGCCCTCTCCGCACAGCGTGGAGACGGGGCAAGGAGGGGGGTT ATTAGGATTGGTGGTTTTGTTTTGCTTTGTTTAAAGCCGTGGGAAAATGGCACA ACTTTACCTCTGTGGGAGATGCAACACTGAGAGCCAAGGGGTGGGAGTTGGGA TAATTTTTATATAAAAGAAGTTTTTCCACTTTGAATTGCTAAAAGTGGCATTTTTC CTATGTGCAGTCACTCCTCTCATTTCTAAAATAGGGACGTGGCCAGGCACGGT GGCTCATGCCTGTAATCCCAGCACTTTGGGAGGCCGAGGCAGGCGGCTCACG AGGTCAGGAGATCGAGACTATCCTGGCTAACACGGTAAAACCCTGTCTCTACTA AAAGTACAAAAAATTAGCTGGGCGTGGTGGTGGGCACCTGTAGTCCCAGCTAC TCGGGAGGCTGAGGCAGGAGAAAGGCATGAATCCAAGAGGCAGAGCTTGCAG TGAGCTGAGATCACGCCATTGCACTCCAGCCTGGGCAACAGTGTTAAGACTCT GTCTCAAATATAAATAAATAAATAAATAAATAAATAAATAAATAAAAATAAAGCGA GATGTTGCCCTCAAA	2032
3UTR-010	LRP1; low density lipoprotein receptor-related protein 1	GGCCCTGCCCCGTCGGACTGCCCCCAGAAAGCCTCCTGCCCCCTGCCAGTGA AGTCCTTCAGTGAGCCCCTCCCCAGCCAGCCCTTCCCTGGCCCCGCCGGATGT ATAAATGTAAAAATGAAGGAATTACATTTTATATGTGAGCGAGCAAGCCGGCAA GCGAGCACAGTATTATTTCTCCATCCCCTCCCTGCCTGCTCCTTGGCACCCCCA TGCTGCCTTCAGGGAGACAGGCAGGGAGGGCTTGGGGCTGCACCTCCTACCC	2033

		TCCCACCAGAACGCACCCCACTGGGAGAGCTGGTGGTGCAGCCTTCCCCTCC CTGTATAAGACACTTTGCCAAGGCTCTCCCCTCTCGCCCCATCCCTGCTTGCCC GCTCCCACAGCTTCCTGAGGGCTAATTCTGGGAAGGGAGAGTTCTTTGCTGCC CCTGTCTGGAAGACGTGGCTCTGGGTGAGGTAGGCGGGAAAGGATGGAGTGT TTTAGTTCTTGGGGGAGGCCACCCCAAACCCCAGCCCCAACTCCAGGGGCACC TATGAGATGGCCATGCTCAACCCCCCTCCCAGACAGGCCCTCCCTGTCTCCAG GGCCCCCACCGAGGTTCCCAGGGCTGGAGACTTCCTCTGGTAAACATTCCTCC AGCCTCCCCTCCCCTGGGGACGCCAAGGAGGTGGGCCACACCCAGGAAGGGA AAGCGGGCAGCCCCGTTTTGGGGACGTGAACGTTTTAATAATTTTTGCTGAATT CCTTTACAACTAAATAACACAGATATTGTTATAAATAAAATTGT	
3UTR-011	Nnt1; cardiotrophin-like cytokine factor 1	ATATTAAGGATCAAGCTGTTAGCTAATAATGCCACCTCTGCAGTTTTGGGAACA GGCAAATAAAGTATCAGTATACATGGTGATGTACATCTGTAGCAAAGCTCTTGG AGAAAATGAAGACTGAAGAAAGCAAAGCAAAAACTGTATAGAGAGATTTTTCAA AAGCAGTAATCCCTCAATTTTAAAAAAGGATTGAAAATTCTAAATGTCTTTCTGT GCATATTTTTTGTGTTAGGAATCAAAAGTATTTTATAAAAGGAGAAAGAACAGCC TCATTTTAGATGTAGTCCTGTTGGATTTTTTATGCCTCCTCAGTAACCAGAAATG TTTTAAAAAACTAAGTGTTTAGGATTTCAAGACAACATTATACATGGCTCTGAAA TATCTGACACAATGTAAACATTGCAGGCACCTGCATTTTATGTTTTTTTTTTCAAC AAATGTGACTAATTTGAAACTTTTATGAACTTCTGAGCTGTCCCCTTGCAATTCA ACCGCAGTTTGAATTAATCATATCAAATCAGTTTTAATTTTTTAAATTGTACTTCA GAGTCTATATTTCAAGGGCACATTTTCTCACTACTATTTTAATACATTAAAGGACT AAATAATCTTTCAGAGATGCTGGAAACAAATCATTTGCTTTATATGTTTCATTAGA ATACCAATGAAACATACAACTTGAAAATTAGTAATAGTATTTTTGAAGATCCCATT TCTAATTGGAGATCTCTTTAATTTCGATCAACTTATAATGTGTAGTACTATATTAA GTGCACTTGAGTGGAATTCAACATTTGACTAATAAAATGAGTTCATCATGTTGGC AAGTGATGTGGCAATTATCTCTGGTGACAAAAGAGTAAAATCAAATATTTCTGCC TGTTACAAATATCAAGGAAGACCTGCTACTATGAAATAGATGACATTAATCTGTC TTCACTGTTTATAATACGGATGGATTTTTTTTCAAATCAGTGTGTGTTTTGAGGT CTTATGTAATTGATGACATTTGAGAGAAATGGTGGCTTTTTTTAGCTACCTCTTT GTTCATTTAAGCACCAGTAAAGATCATGTCTTTTTATAGAAGTGTAGATTTTCTTT GTGACTTTGCTATCGTGCCTAAAGCTCTAAATATAGGTGAATGTGTGATGAATA CTCAGATTATTTGTCTCTCTATATAATTAGTTTGGTACTAAGTTTCTCAAAAAATT ATTAACACATGAAAGACAATCTCTAAACCAGAAAAAGAAGTAGTACAAATTTTGT TACTGTAATGCTCGCGTTTAGTGAGTTTAAAACACACAGTATCTTTTGGTTTTAT AATCAGTTTCTATTTTGCTGTGCCTGAGATTAAGATCTGTGTATGTGTGTGTGTG TGTGTGTGCGTTTGTGTGTTAAAGCAGAAAAGACTTTTTTAAAAGTTTTAAGTGA TAAATGCAATTTGTTAATTGATCTTAGATCACTAGTAAACTCAGGGCTGAATTAT ACCATGTATATTCTATTAGAAGAAAGTAAACACCATCTTTATTCCTGCCCTTTTTC TTCTCTCAAAGTAGTTGTAGTTATATCTAGAAAGAAGCAATTTTGATTTCTTGAAA AGGTAGTTCCTGCACTCAGTTTAAACTAAAAATAATCATACTTGGATTTTATTTAT TTTTGTCATAGTAAAAATTTTAATTTATATATATTTTTATTTAGTATTATCTTATTCT TTGCTATTTGCCAATCCTTTGTCATCAATTGTGTTAAATGAATTGAAAATTCATGC CCTGTTCATTTTATTTTACTTTATTGGTTAGGATATTTAAAGGATTTTTGTATATAT AATTTCTTAAATTAATATTCCAAAAGGTTAGTGGACTTAGATTATAAATTATGGCA AAAATCTAAAAACAACAAAAATGATTTTTATACATTCTATTTCATTATTCCTCTTTT TCCAATAAGTCATACAATTGGTAGATATGACTTATTTTATTTTTGTATTATTCACT ATATCTTTATGATATTTAAGTATAAATAATTAAAAAAATTTATTGTACCTTATAGTC TGTCACCAAAAAAAAAAAATTATCTGTAGGTAGTGAAATGCTAATGTTGATTTGT CTTTAAGGGCTTGTTAACTATCCTTTATTTTCTCATTTGTCTTAAATTAGGAGTTT GTGTTTAAATTACTCATCTAAGCAAAAAATGTATATAAATCCCATTACTGGGTATA TACCCAAAGGATTATAAATCATGCTGCTATAAAGACACATGCACACGTATGTTTA TTGCAGCACTATTCACAATAGCAAAGACTTGGAACCAACCCAAATGTCCATCAA TGATAGACTTGATTAAGAAAATGTGCACATATACACCATGGAATACTATGCAGC CATAAAAAAGGATGAGTTCATGTCCTTTGTAGGGACATGGATAAAGCTGGAAAC CATCATTCTGAGCAAACTATTGCAAGGACAGAAAACCAAACACTGCATGTTCTC ACTCATAGGTGGGAATTGAACAATGAGAACACTTGGACACAAGGTGGGGAACA CCACACACCAGGGCCTGTCATGGGGTGGGGGGAGTGGGGAGGGATAGCATTA GGAGATATACCTAATGTAAATGATGAGTTAATGGGTGCAGCACACCAACATGGC	2034

		ACATGTATACATATGTAGCAAACCTGCACGTTGTGCACATGTACCCTAGAACTT AAAGTATAATTAAAAAAAAAAAGAAAACAGAAGCTATTTATAAAGAAGTTATTTGC TGAAATAAATGTGATCTTTCCCATTAAAAAAATAAAGAAATTTTGGGGTAAAAAA ACACAATATATTGTATTCTTGAAAAATTCTAAGAGAGTGGATGTGAAGTGTTCTC ACCACAAAAGTGATAACTAATTGAGGTAATGCACATATTAATTAGAAAGATTTTG TCATTCCACAATGTATATATACTTAAAAATATGTTATACACAATAAATACATACAT TAAAAAATAAGTAAATGTA	
3UTR-012	Col6a1; collagen, type VI, alpha 1	CCCACCCTGCACGCCGGCACCAAACCCTGTCCTCCCACCCCTCCCCACTCATC ACTAAACAGAGTAAAATGTGATGCGAATTTTCCCGACCAACCTGATTCGCTAGA TTTTTTTTAAGGAAAAGCTTGGAAAGCCAGGACACAACGCTGCTGCCTGCTTTG TGCAGGGTCCTCCGGGGGCTCAGCCCTGAGTTGGCATCACCTGCGCAGGGCCC TCTGGGGCTCAGCCCTGAGCTAGTGTCACCTGCACAGGGCCCTCTGAGGCTCA GCCCTGAGCTGGCGTCACCTGTGCAGGGCCCTCTGGGGCTCAGCCCTGAGCT GGCCTCACCTGGGTTCCCCACCCCGGGCTCTCCTGCCCTGCCCTCCTGCCCG CCCTCCCTCCTGCCTGCGCAGCTCCTTCCCTAGGCACCTCTGTGCTGCATCCC ACCAGCCTGAGCAAGACGCCCTCTCGGGGCCTGTGCCGCACTAGCCTCCCTC TCCTCTGTCCCCATAGCTGGTTTTTCCCACCAATCCTCACCTAACAGTTACTTTA CAATTAAACTCAAAGCAAGCTCTTCTCCTCAGCTTGGGGCAGCCATTGGCCTCT GTCTCGTTTTGGGAAACCAAGGTCAGGAGGCCGTTGCAGACATAAATCTCGGC GACTCGGCCCCGTCTCCTGAGGGTCCTGCTGGTGACCGGCCTGGACCTTGGC CCTACAGCCCTGGAGGCCGCTGCTGACCAGCACTGACCCCGACCTCAGAGAG TACTCGCAGGGGCGCTGGCTGCACTCAAGACCCTCGAGATTAACGGTGCTAAC CCCGTCTGCTCCTCCCTCCCGCAGAGACTGGGGCCTGGACTGGACATGAGAG CCCCTTGGTGCCACAGAGGGCTGTGTCTTACTAGAAACAACGCAAACCTCTCC TTCCTCAGAATAGTGATGTGTTCGACGTTTTATCAAAGGCCCCCTTTCTATGTTC ATGTTAGTTTTGCTCCTTCTGTGTTTTTTTCTGAACCATATCCATGTTGCTGACTT TTCCAAATAAAGGTTTTCACTCCTCTC	2035
3UTR-013	Calr; calreticulin	AGAGGCCTGCCTCCAGGGCTGGACTGAGGCCTGAGCGCTCCTGCCGCAGAGC TGGCCGCGCCAAATAATGTCTCTGTGAGACTCGAGAACTTTCATTTTTTTCCAG GCTGGTTCGGATTTGGGGTGGATTTTGGTTTTGTTCCCCTCCTCCACTCTCCCC CACCCCCTCCCCGCCCTTTTTTTTTTTTTTTTTTAAACTGGTATTTTATCTTTGAT TCTCCTTCAGCCCTCACCCCTGGTTCTCATCTTTCTTGATCAACATCTTTTCTTG CCTCTGTCCCCTTCTCTCATCTCTTAGCTCCCCTCCAACCTGGGGGGCAGTGG TGTGGAGAAGCCACAGGCCTGAGATTTCATCTGCTCTCCTTCCTGGAGCCCAG AGGAGGGCAGCAGAAGGGGGTGGTGTCTCCAACCCCCCAGCACTGAGGAAGA ACGGGGCTCTTCTCATTTCACCCCTCCCTTTCTCCCCTGCCCCCAGGACTGGG CCACTTCTGGGTGGGGCAGTGGGTCCCAGATTGGCTCACACTGAGAATGTAAG AACTACAAACAAAATTTCTATTAAATTAAATTTTGTGTCTCC	2036
3UTR-014	Col1a1; collagen, type I, alpha 1	CTCCCTCCATCCCAACCTGGCTCCCTCCCACCCAACCAACTTTCCCCCCAACC CGGAAACAGACAAGCAACCCAAACTGAACCCCCTCAAAAGCCAAAAAATGGGA GACAATTTCACATGGACTTTGGAAAATATTTTTTTCCTTTGCATTCATCTCTCAAA CTTAGTTTTTATCTTTGACCAACCGAACATGACCAAAAACCAAAAGTGCATTCAA CCTTACCAAAAAAAAAAAAAAAAAAAAGAATAAATAAATAACTTTTTAAAAAAGGAA GCTTGGTCCACTTGCTTGAAGACCCATGCGGGGGTAAGTCCCTTTCTGCCCGT TGGGCTTATGAAACCCCAATGCTGCCCTTTCTGCTCCTTTCTCCACACCCCCCT TGGGGCCTCCCCTCCACTCCTTCCCAAATCTGTCTCCCCAGAAGACACAGGAA ACAATGTATTGTCTGCCCAGCAATCAAAGGCAATGCTCAAACACCCAAGTGGCC CCCACCCTCAGCCCGCTCCTGCCCGCCCAGCACCCCCAGGCCCTGGGGGACC TGGGGTTCTCAGACTGCCAAAGAAGCCTTGCCATCTGGCGCTCCCATGGCTCT TGCAACATCTCCCCTTCGTTTTTGAGGGGGTCATGCCGGGGGAGCCACCAGCC CCTCACTGGGTTCGGAGGAGAGTCAGGAAGGGCCACGACAAAGCAGAAACAT CGGATTTGGGGAACGCGTGTCAATCCCTTGTGCCGCAGGGCTGGGCGGGAGA GACTGTTCTGTTCCTTGTGTAACTGTGTTGCTGAAAGACTACCTCGTTCTTGTCT TGATGTGTCACCGGGGCAACTGCCTGGGGGCGGGGATGGGGGCAGGGTGGA AGCGGCTCCCCATTTTATACCAAAGGTGCTACATCTATGTGATGGGTGGGGTG GGGAGGGAATCACTGGTGCTATAGAAATTGAGATGCCCCCCCAGGCCAGCAAA TGTTCCTTTTTGTTCAAAGTCTATTTTTATTCCTTGATATTTTTCTTTTTTTTTTTT TTTTTTGTGGATGGGGACTTGTGAATTTTTCTAAAGGTGCTATTTAACATGGGAG	2037

		GAGAGCGTGTGCGGCTCCAGCCCAGCCCGCTGCTCACTTTCCACCCTCTCTCC ACCTGCCTCTGGCTTCTCAGGCCTCTGCTCTCCGACCTCTCTCCTCTGAAACCC TCCTCCACAGCTGCAGCCCATCCTCCCGGCTCCCTCCTAGTCTGTCCTGCGTC CTCTGTCCCCGGGTTTCAGAGACAACTTCCCAAAGCACAAAGCAGTTTTTCCCC CTAGGGGTGGGAGGAAGCAAAAGACTCTGTACCTATTTTGTATGTGTATAATAA TTTGAGATGTTTTTAATTATTTTGATTGCTGGAATAAAGCATGTGGAAATGACCC AAACATAATCCGCAGTGGCCTCCTAATTTCCTTCTTTGGAGTTGGGGGAGGGGT AGACATGGGGAAGGGGCTTTGGGGTGATGGGCTTGCCTTCCATTCCTGCCCTT TCCCTCCCCACTATTCTCTTCTAGATCCCTCCATAACCCCACTCCCCTTTCTCTC ACCCTTCTTATACCGCAAACCTTTCTACTTCCTCTTTCATTTTCTATTCTTGCAAT TTCCTTGCACCTTTTCCAAATCCTCTTCTCCCCTGCAATACCATACAGGCAATCC ACGTGCACAACACACACACACACTCTTCACATCTGGGGTTGTCCAAACCTCATA CCCACTCCCCTTCAAGCCCATCCACTCTCCACCCCCTGGATGCCCTGCACTTG GTGGCGGTGGGATGCTCATGGATACTGGGAGGGTGAGGGGAGTGGAACCCGT GAGGAGGACCTGGGGGCCTCTCCTTGAACTGACATGAAGGGTCATCTGGCCTC TGCTCCCTTCTCACCCACGCTGACCTCCTGCCGAAGGAGCAACGCAACAGGAG AGGGGTCTGCTGAGCCTGGCGAGGGTCTGGGAGGGACCAGGAGGAAGGCGT GCTCCCTGCTCGCTGTCCTGGCCCTGGGGGAGTGAGGGAGACAGACACCTGG GAGAGCTGTGGGGAAGGCACTCGCACCGTGCTCTTGGGAAGGAAGGAGACCT GGCCCTGCTCACCACGGACTGGGTGCCTCGACCTCCTGAATCCCCAGAACACA ACCCCCCTGGGCTGGGGTGGTCTGGGGAACCATCGTGCCCCCGCCTCCCGCC TACTCCTTTTTAAGCTT	
3UTR-015	Plod1; procollagen-lysine, 2-oxoglutarate 5-dioxygenase 1	TTGGCCAGGCCTGACCCTCTTGGACCTTTCTTCTTTGCCGACAACCACTGCCCA GCAGCCTCTGGGACCTCGGGGTCCCAGGGAACCCAGTCCAGCCTCCTGGCTG TTGACTTCCCATTGCTCTTGGAGCCACCAATCAAAGAGATTCAAAGAGATTCCT GCAGGCCAGAGGCGGAACACACCTTTATGGCTGGGGCTCTCCGTGGTGTTCT GGACCCAGCCCCTGGAGACACCATTCACTTTTACTGCTTTGTAGTGACTCGTGC TCTCCAACCTGTCTTCCTGAAAAACCAAGGCCCCCTTCCCCCACCTCTTCCATG GGGTGAGACTTGAGCAGAACAGGGGCTTCCCCAAGTTGCCCAGAAAGACTGTC TGGGTGAGAAGCCATGGCCAGAGCTTCTCCCAGGCACAGGTGTTGCACCAGG GACTTCTGCTTCAAGTTTTGGGGTAAAGACACCTGGATCAGACTCCAAGGGCT GCCCTGAGTCTGGGACTTCTGCCTCCATGGCTGGTCATGAGAGCAAACCGTAG TCCCCTGGAGACAGCGACTCCAGAGAACCTCTTGGGAGACAGAAGAGGCATCT GTGCACAGCTCGATCTTCTACTTGCCTGTGGGGAGGGGAGTGACAGGTCCACA CACCACACTGGGTCACCCTGTCCTGGATGCCTCTGAAGAGAGGGACAGACCGT CAGAAACTGGAGAGTTTCTATTAAAGGTCATTTAAACCA	2038
3UTR-016	Nucb1; nucleobindin 1	TCCTCCGGGACCCCAGCCCTCAGGATTCCTGATGCTCCAAGGCGACTGATGGG CGCTGGATGAAGTGGCACAGTCAGCTTCCCTGGGGGCTGGTGTCATGTTGGG CTCCTGGGGCGGGGGCACGGCCTGGCATTTCACGCATTGCTGCCACCCCAGG TCCACCTGTCTCCACTTTCACAGCCTCCAAGTCTGTGGCTCTTCCCTTCTGTCC TCCGAGGGGCTTGCCTTCTCTCGTGTCCAGTGAGGTGCTCAGTGATCGGCTTA ACTTAGAGAAGCCCGCCCCCTCCCCTTCTCCGTCTGTCCCAAGAGGGTCTGCT CTGAGCCTGCGTTCCTAGGTGGCTCGGCCTCAGCTGCCTGGGTTGTGGCCGC CCTAGCATCCTGTATGCCCACAGCTACTGGAATCCCCGCTGCTGCTCCGGGCC AAGCTTCTGGTTGATTAATGAGGGCATGGGGTGGTCCCTCAAGACCTTCCCCT ACCTTTTGTGGAACCAGTGATGCCTCAAAGACAGTGTCCCCTCCACAGCTGGG TGCCAGGGGCAGGGGATCCTCAGTATAGCCGGTGAACCCTGATACCAGGAGC CTGGGCCTCCCTGAACCCCTGGCTTCCAGCCATCTCATCGCCAGCCTCCTCCT GGACCTCTTGGCCCCCAGCCCCTTCCCCACACAGCCCCAGAAGGGTCCCAGA GCTGACCCCACTCCAGGACCTAGGCCCAGCCCCTCAGCCTCATCTGGAGCCC CTGAAGACCAGTCCCACCCACCTTTCTGGCCTCATCTGACACTGCTCCGCATC CTGCTGTGTGTCCTGTTCCATGTTCCGGTTCCATCCAAATACACTTTCTGGAAC AAA	2039
3UTR-017	α-globin	GCTGGAGCCTCGGTGGCCATGCTTCTTGCCCCTTGGGCCTCCCCCCAGCCCC TCCTCCCCTTCCTGCACCCGTACCCCCGTGGTCTTTGAATAAAGTCTGAGTGG GCGGC	2040

5' UTR and Translation Initiation

[0333] Natural 5'UTRs bear features which play roles in translation initiation They harbor signatures like Kozak sequences which are commonly known to be involved in the process by which the ribosome initiates translation of many genes. Kozak sequences have the consensus CCR(A/G)CCAUGG, where R is a purine (adenine or guanine) three bases upstream of the start codon (AUG), which is followed by another 'G'. 5'UTR also have been known to form secondary structures which are involved in elongation factor binding

[0334] By engineering the features typically found in abundantly expressed genes of specific target organs, one can enhance the stability and protein production of the polynucleotides of the invention. For example, introduction of 5' UTR of liver-expressed mRNA, such as albumin, serum amyloid A, Apolipoprotein A/B/E, transferrin, alpha fetoprotein, erythropoietin, or Factor VIII, could be used to enhance expression of a nucleic acid molecule, such as a polynucleotides, in hepatic cell lines or liver. Likewise, use of 5' UTR from other tissue-specific mRNA to improve expression in that tissue is possible for muscle (MyoD, Myosin, Myoglobin, Myogenin, Herculin), for endothelial cells (Tie-1, CD36), for myeloid cells (CIEBP, AML1, G-CSF, GM-CSF, CD11b, MSR, Fr-1, i-NOS), for leukocytes (CD45, CD18), for adipose tissue (CD36, GLUT4, ACRP30, adiponectin) and for lung epithelial cells (SP-A/B/C/D).

[0335] Untranslated regions useful in the design and manufacture of polynucleotides (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention) include, but are not limited, to those disclosed in co-pending, co-owned US Provisional Application (USSN) 61/829,372, US Provisional Application 61/829,372 (USSN), and International Application, PCT/US14/21522 filed March 7, 2014.

[0336] Other non-UTR sequences may also be used as regions or subregions within the polynucleotides. For example, introns or portions of introns sequences may be incorporated into regions of the polynucleotides of the invention. Incorporation of intronic sequences may increase protein production as well as polynucletoide levels.

[0337] Combinations of features may be included in flanking regions and may be contained within other features. For example, the ORF may be flanked by a 5' UTR which may contain a strong Kozak translational initiation signal and/or a 3' UTR which may include an oligo(dT) sequence for templated addition of a poly-A fail. 5'UTR may comprise a first polynucleotide fragment and a second polynucleotide fragment from the same and/or different genes such as the 5'UTRs described in US Patent Application Publication No. 20100293625.

[0338] Co-pending, co-owned US Provisional Application (USSN) 61/829,372 and US Provisional Application 61/829,372 (USSN) provides a listing of exemplary UTRs which may be utilized in the polynucleotide of the present invention as flanking regions. Variants of 5' or 3' UTRs may be utilized wherein one or more nucleotides are added or removed to the termini, including A, T, C or G.

[0339] It should be understood that any UTR from any gene may be incorporated into the regions of the polynucleotide. Furthermore, multiple wild-type UTRs of any known gene may be utilized. It is also within the scope of the present invention to provide artificial UTRs which are not variants of wild type regions. These UTRs or portions thereof may be placed in the same orientation as in the transcript from which they were selected or may be altered in orientation or location Hence a 5' or 3' UTR may be inverted, shortened, lengthened, made with one or more other 5' UTRs or 3' UTRs. As used herein, the term "altered" as it relates to a UTR sequence, means that the UTR has been changed in some way in relation to a reference sequence. For example, a 3' or5' UTR may be altered relative to a wild type or native UTR by the change in orientation or location as taught above or may be altered by the inclusion of additional nucleotides, deletion of nucleotides, swapping or transposition of nucleotides. Any of these changes producing an "altered" UTR (whether 3' or 5') comprise a variant UTR.

[0340] In one embodiment, a double, triple or quadruple UTR such as a 5' or 3' UTR may be used. As used herein, a "double" UTR is one in which two copies of the same UTR are encoded either in series or substantially in series. For example, a double beta-globin 3' UTR may be used as described in US Patent publication 20100129877.

[0341] It is also within the scope of the present invention to have patterned UTRs. As used herein "patterned UTRs" are those UTRs which reflect a repeating or alternating pattern, such as ABABAB or AABBAABBAABB or ABCABCABC or variants thereof repeated once, twice, or more than 3 times. In these patterns, each letter, A, B, or C represent a different UTR at the nucleotide level.

[0342] In one embodiment, flanking regions are selected from a family of transcripts whose proteins share a common function, structure, feature of property. For example, polypeptides of interest may belong to a family of proteins which are expressed in a particular cell, tissue or at some time during development. The UTRs from any of these genes may be swapped for any other UTR of the same or different family of proteins to create a new polynucleotide. As used herein, a "family of proteins" is used in the broadest sense to refer to a group of two or more polypeptides of interest which share at least one function, structure, feature, localization, origin, or expression pattern.

[0343] The untranslated region may also include translation enhancer elements (TEE). As a non-limiting example, the TEE may include those described in US Application No. 20090226470.

3' UTR and the AU Rich Elements

[0344] Natural or wild type 3' UTRs are known to have stretches of Adenosines and Uridines embedded in them. These AU rich signatures are particularly prevalent in genes with high rates of turnover. Based on their sequence features and functional properties, the AU rich elements (AREs) can be separated into three classes (Chen et al, 1995): Class I AREs contain several dispersed copies of an AUUUA motif within U-rich regions. C-Myc and MyoD contain class I AREs. Class II AREs possess two or more overlapping UUAUUUA(U/A)(U/A) nonamers. Molecules containing this type of AREs include GM-CSF and TNF-a. Class III ARES are less well defined. These U rich regions do not contain an AUUUA motif. c-Jun and Myogenin are two well-studied examples of this class. Most proteins binding to the AREs are known to destabilize the messenger, whereas members of the ELAV family, most notably HuR, have been documented to increase the stability of mRNA. HuR binds to AREs of all the three classes. Engineering the HuR specific binding sites into the 3' UTR of nucleic acid molecules will lead to HuR binding and thus, stabilization of the message *in vivo.*

[0345] Introduction, removal or modification of 3' UTR AU rich elements (AREs) can be used to modulate the stability of polynucleotides of the invention (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention). When engineering specific polynucleotides, one or more copies of an ARE can be introduced to make polynucleotides of the invention less stable and thereby curtail translation and decrease production of the resultant protein. Likewise, AREs can be identified and removed or mutated to increase the intracellular stability and thus increase translation and production of the resultant protein. Transfection experiments can be conducted in relevant cell lines, using polynucleotides of the invention and protein production can be assayed at various time points post-transfection. For example, cells can be transfected with different ARE-engineering molecules and by using an ELISA kit to the relevant protein and assaying protein produced at 6 hour, 12 hour, 24 hour, 48 hour, and 7 days post-transfection.

microRNA Binding Sites

[0346] microRNAs (or miRNA) are 19-25 nucleotide long noncoding RNAs that bind to the 3'UTR of nucleic acid molecules and down-regulate gene expression either by reducing nucleic acid molecule stability or by inhibiting translation. The polynucleotides of the invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) may comprise one or more microRNA target sequences, microRNA seqences, or microRNA seeds. Such sequences may correspond to any known microRNA such as those taught in US Publication US2005/0261218 and US Publication US2005/0059005.

[0347] A microRNA sequence comprises a "seed" region, i.e., a sequence in the region of positions 2-8 of the mature microRNA, which sequence has perfect Watson-Crick complementarity to the miRNA target sequence. A microRNA seed may comprise positions 2-8 or 2-7 of the mature microRNA. In some embodiments, a microRNA seed may comprise 7 nucleotides (e.g., nucleotides 2-8 of the mature microRNA), wherein the seed-complementary site in the corresponding miRNA target is flanked by an adenine (A) opposed to microRNA position 1. In some embodiments, a microRNA seed may comprise 6 nucleotides (e.g., nucleotides 2-7 of the mature microRNA), wherein the seed-complementary site in the corresponding miRNA target is flanked byan adenine (A) opposed to microRNA position 1. See for example, Grimson A, Farh KK, Johnston WK, Garrett-Engele P, Lim LP, Bartel DP; Mol Cell. 2007 Jul 6;27(1):91-105. The bases of the microRNA seed have complete complementarity with the target sequence. By engineering microRNA target sequences into the polynucleotides (e.g., in a 3'UTR like region or other region) of the invention one can target the molecule for degradation or reduced translation, provided the microRNA in question is available. This process will reduce the hazard of off target effects upon nucleic acid molecule delivery. Identification of microRNA, microRNA target regions, and their expression patterns and role in biology have been reported (Bonauer et al., Curr Drug Targets 2010 11:943-949; Anand and Cheresh Curr Opin Hematol 2011 18:171-176; Contreras and Rao Leukemia 2012 26:404-413 (2011 Dec 20. doi: 10.1038/leu.2011.356); Bartel Cell 2009 136:215-233; Landgraf et al, Cell, 2007 129:1401-1414).

[0348] For example, if the nucleic acid molecule is an mRNA and is not intended to be delivered to the liver but ends up there, then miR-122, a microRNA abundant in liver, can inhibit the expression of the gene of interest if one or multiple target sites of miR-122 are engineered into the 3' UTR region of the polynucleotides. Introduction of one or multiple binding sites for different microRNA can be engineered to further decrease the longevity, stability, and protein translation of polynucleotides.

[0349] As used herein, the term "microRNA site" refers to a microRNA target site or a microRNA recognition site, or any nucleotide sequence to which a microRNA binds or associates. It should be understood that "binding" may follow traditional Watson-Crick hybridization rules or may reflect any stable association of the microRNA with the target sequence at or adjacent to the microRNA site.

[0350] Conversely, for the purposes of the polynucleotides of the present invention, microRNA binding sites can be engineered out of (i.e. removed from) sequences in which they occur, e.g., in order to increase protein expression in specific tissues. For example, miR-122 binding sites may be removed to improve protein expression in the liver. Regulation of expression in multiple tissues can be accomplished through introduction or removal or one or several microRNA

binding sites.

[0351] Examples of tissues where microRNA are known to regulate mRNA, and thereby protein expression, include, but are not limited to, liver (miR-122), muscle (miR-133, miR-206, miR-208), endothelial cells (miR-17-92, miR-126), myeloid cells (miR-142-3p, miR-142-5p, miR-16, miR-21, miR-223, miR-24, miR-27), adipose tissue (let-7, miR-30c), heart (miR-1d, miR-149), kidney (miR-192, miR-194, miR-204), and lung epithelial cells (let-7, miR-133, miR-126). MicroRNA can also regulate complex biological processes such as angiogenesis (miR-132) (Anand and Cheresh Curr Opin Hematol 2011 18:171-176).

[0352] Expression profiles, microRNA and cell lines useful in the present invention include those taught in for example,U.S. Provisional Application Nos 61/857,436 and 61/857,304 each filed July 23, 2013.

[0353] In the polynucleotides of the present invention, binding sites for microRNAs that are involved in such processes may be removed or introduced, in order to tailor the expression of the polynucleotides expression to biologically relevant cell types or to the context of relevant biological processes A listing of microRNA, miR sequences and miR binding sites is listed in Table 9 of U.S. Provisional Application No. 61/753,661 filed January 17, 2013, in Table 9 of U.S. Provisional Application No. 61/754,159 filed January 18, 2013, and in Table 7 of U.S. Provisional Application No. 61/758,921 filed January 31, 2013.

[0354] Examples of use of microRNA to drive tissue or disease-specific gene expression are listed (Getner and Naldini, Tissue Antigens. 2012, 80:393-403). In addition, microRNA seed sites can be incorporated into mRNA to decrease expression in certain cells which results in a biological improvement. An example of this is incorporation of miR-142 sites into a UGT1A1-expressing lentiviral vector. The presence of miR-142 seed sites reduced expression in hematopoietic cells, and as a consequence reduced expression in antigen-presentating cells, leading to the absence of an immune response against the virally expressed UGT1A1 (Schmitt et al., Gastroenterology 2010; 139:999-1007; Gonzalez-Asequinolaza et al. Gastroenterology 2010, 139:726-729) . Incorporation of miR-142 sites into modified mRNA could not only reduce expression of the encoded protein in hematopoietic cells, but could also reduce or abolish immune responses to the mRNA-encoded protein Incorporation of miR-142 seed sites (one or multiple) into mRNA would be important in the case of treatment of patients with complete protein deficiencies (UGT1A1 type I, LDLR-deficient patients, CRIM-negative Pompe patients, etc.).

[0355] Lastly, through an understanding of the expression patterns of microRNA in different cell types, polynucleotides (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention) can be engineered for more targeted expression in specific cell types or only under specific biological conditions. Through introduction of tissue-specific microRNA binding sites, polynucleotides could be designed that would be optimal for protein expression in a tissue or in the context of a biological condition.

[0356] Transfection experiments can be conducted in relevant cell lines, using engineered polynucleotides and protein production can be assayed at various time points post-transfection. For example, cells can be transfected with different microRNA binding site-engineering polynucleotides and by using an ELISA kit to the relevant protein and assaying protein produced at 6 hour, 12 hour, 24 hour, 48 hour, 72 hour and 7 days post-transfection. *In vivo* experiments can also be conducted using microRNA-binding site-engineered molecules to examine changes in tissue-specific expression of formulated polynucleotides.

Regions having a 5'Cap

[0357] The 5' cap structure of a natural mRNA is involved in nuclear export, increasing mRNA stability and binds the mRNA Cap Binding Protein (CBP), which is responsibile for mRNA stability in the cell and translation competency through the association of CBP with poly(A) binding protein to form the mature cyclic mRNA species. The cap further assists the removal of 5' proximal introns removal during mRNA splicing.

[0358] Endogenous mRNA molecules may be 5'-end capped generating a 5'-ppp-5'-triphosphate linkage between a terminal guanosine cap residue and the 5'-terminal transcribed sense nucleotide of the mRNA molecule. This 5'-guanylate cap may then be methylated to generate an N7-methyl-guanylate residue. The ribose sugars of the terminal and/or anteterminal transcribed nucleotides of the 5' end of the mRNA may optionally also be 2'-O-methylated. 5'-decapping through hydrolysis and cleavage of the guanylate cap structure may target a nucleic acid molecule, such as an mRNA molecule, for degradation

[0359] In some embodiments, polynucleotides (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention) may be designed to incorporate a cap moiety. Modifications to the polynucleotides of the present invention may generate a non-hydrolyzable cap structure preventing decapping and thus increasing mRNA half-life. Because cap structure hydrolysis requires cleavage of 5'-ppp-5' phosphorodiester linkages, modified nucleotides may be used during the capping reaction. For example, a Vaccinia Capping Enzyme from New England Biolabs (Ipswich, MA) may be used with α-thio-guanosine nucleotides according to the manufacturer's instructions to create a phosphorothioate linkage in the 5'-ppp-5' cap. Additional modified guanosine nucleotides may be used such as α-methyl-phosphonate and seleno-phosphate nucleotides.

[0360] Additional modifications include, but are not limited to, 2'-O-methylation of the ribose sugars of 5'-terminal and/or 5'-anteterminal nucleotides of the polynucleotide (as mentioned above) on the 2'-hydroxyl group of the sugar ring. Multiple distinct 5'-cap structures can be used to generate the 5'-cap of a nucleic acid molecule, such as a polynucleotide which functions as an mRNA molecule.

[0361] Cap analogs, which herein are also referred to as synthetic cap analogs, chemical caps, chemical cap analogs, or structural or functional cap analogs, differ from natural (i.e. endogenous, wild-type or physiological) 5'-caps in their chemical structure, while retaining cap function. Cap analogs may be chemically (i.e. non-enzymatically) or enzymatically synthesized and/or linked to the polynucleotides of the invention.

[0362] For example, the Anti-Reverse Cap Analog (ARCA) cap contains two guanines linked by a 5'-5'-triphosphate group, wherein one guanine contains an N7 methyl group as well as a 3'-O-methyl group (i.e., N7,3'-O-dimethyl-guanosine-5'-triphosphate-5'-guanosine (m^7G-3'mppp-G; which may equivaliently be designated 3' O-Me-m7G(5')ppp(5')G). The 3'-O atom of the other, unmodified, guanine becomes linked to the 5'-terminal nucleotide of the capped polynucleotide. The N7- and 3'-O-methlyated guanine provides the terminal moiety of the capped polynucleotide.

[0363] Another exemplary cap is mCAP, which is similar to ARCA but has a 2'-O-methyl group on guanosine (i.e., N7,2'-O-dimethyl-guanosine-5'-triphosphate-5'-guanosine, m^7Gm-ppp-G).

[0364] In one embodiment, the cap is a dinucleotide cap analog. As a non-limiting example, the dinucleotide cap analog may be modified at different phosphate positions with a boranophosphate group or a phophoroselenoate group such as the dinucleotide cap analogs described in US Patent No. US 8,519,110.

[0365] In another embodiment, the cap is a cap analog is a N7-(4-chlorophenoxyethyl) substituted dicucleotide form of a cap analog known in the art and/or described herein. Non-limiting examples of a N7-(4-chlorophenoxyethyl) substituted dicucleotide form of a cap analog include a N7-(4-chlorophenoxyethyl)-G(5')ppp(5')G and a N7-(4-chlorophenoxyethyl)-$m^{3'-O}$G(5')ppp(5')G cap analog (See e.g., the various cap analogs and the methods of synthesizing cap analogs described in Kore et al. Bioorganic & Medicinal Chemistry 2013 21:4570-4574). In another embodiment, a cap analog of the present invention is a 4-chloro/bromophenoxyethyl analog.

[0366] While cap analogs allow for the concomitant capping of a polynucleotide or a region thereof, in an in vitro transcription reaction, up to 20% of transcripts can remain uncapped. This, as well as the structural differences of a cap analog from an endogenous 5'-cap structures of nucleic acids produced by the endogenous, cellular transcription machinery, may lead to reduced translational competency and reduced cellular stability.

[0367] Polynucleotides of the invention may also be capped post-manufacture (whether IVT or chemical synthesis), using enzymes, in order to generate more authentic 5'-cap structures. As used herein, the phrase "more authentic" refers to a feature that closely mirrors or mimics, either structurally or functionally, an endogenous or wild type feature. That is, a "more authentic" feature is better representative of an endogenous, wild-type, natural or physiological cellular function and/or structure as compared to synthetic features or analogs, etc., of the prior art, or which outperforms the corresponding endogenous, wild-type, natural or physiological feature in one or more respects. Non-limiting examples of more authentic 5'cap structures of the present invention are those which, among other things, have enhanced binding of cap binding proteins, increased half life, reduced susceptibility to 5' endonucleases and/or reduced 5'decapping, as compared to synthetic 5'cap structures known in the art (or to a wild-type, natural or physiological 5'cap structure). For example, recombinant Vaccinia Virus Capping Enzyme and recombinant 2'-O-methyltransferase enzyme can create a canonical 5'-5'-triphosphate linkage between the 5'-terminal nucleotide of a polynucleotide and a guanine cap nucleotide wherein the cap guanine contains an N7 methylation and the 5'-terminal nucleotide of the mRNA contains a 2'-O-methyl. Such a structure is termed the Cap1 structure. This cap results in a higher translational-competency and cellular stability and a reduced activation of cellular pro-inflammatory cytokines, as compared, e.g., to other 5'cap analog structures known in the art. Cap structures include, but are not limited to, 7mG(5')ppp(5')N,pN2p (cap 0), 7mG(5')ppp(5')NlmpNp (cap 1), and 7mG(5')-ppp(5')NlmpN2mp (cap 2).

[0368] As a non-limiting example, capping chimeric polynucleotides post-manufacture may be more efficient as nearly 100% of the chimeric polynucleotides may be capped. This is in contrast to ~80% when a cap analog is linked to a chimeric polynucleotide in the course of an in vitro transcription reaction

[0369] According to the present invention, 5' terminal caps may include endogenous caps or cap analogs. According to the present invention, a 5' terminal cap may comprise a guanine analog. Useful guanine analogs include, but are not limited to, inosine, N1-methyl-guanosine, 2'fluoro-guanosine, 7-deaza-guanosine, 8-oxo-guanosine, 2-amino-guanosine, LNA-guanosine, and 2-azido-guanosine.

Viral Sequences

[0370] Additional viral sequences such as, but not limited to, the translation enhancer sequence of the barley yellow dwarf virus (BYDV-PAV), the Jaagsiekte sheep retrovirus (JSRV) and/or the Enzootic nasal tumor virus (See e.g., International Pub. No. WO2012129648) can be engineered and inserted in the polynucleotides of the invention and can stimulate the translation of the construct *in vitro* and *in vivo*. Transfection experiments can be conducted in relevant cell

lines at and protein production can be assayed by ELISA at 12hr, 24hr, 48hr, 72 hr and day 7 post-transfection.

IRES Sequences

[0371] Further, provided are polynucleotides (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) which may contain an internal ribosome entry site (IRES). First identified as a feature Picorna virus RNA, IRES plays an important role in initiating protein synthesis in absence of the 5' cap structure. An IRES may act as the sole ribosome binding site, or may serve as one of multiple ribosome binding sites of an mRNA. Polynucleotides containing more than one functional ribosome binding site may encode several peptides or polypeptides that are translated independently by the ribosomes ("multicistronic nucleic acid molecules"). When polynucleotides are provided with an IRES, further optionally provided is a second translatable region. Examples of IRES sequences that can be used according to the invention include without limitation, those from picornaviruses (e.g. FMDV), pest viruses (CFFV), polio viruses (PV), encephalomyocarditis viruses (ECMV), foot-and-mouth disease viruses (FMDV), hepatitis C viruses (HCV), classical swine fever viruses (CSFV), murine leukemia virus (MLV), simian immune deficiency viruses (SIV) or cricket paralysis viruses (CrPV).

Poly-A tails

[0372] During RNA processing, a long chain of adenine nucleotides (poly-A fail) may be added to a polynucleotide such as an mRNA molecule in order to increase stability. Immediately after transcription, the 3' end of the transcript may be cleaved to free a 3' hydroxyl. Then poly-A polymerase adds a chain of adenine nucleotides to the RNA The process, called polyadenylation, adds a poly-A tail that can be between, for example, approximately 80 to approximately 250 residues long (SEQ ID NO:2279), including approximately 80, 90, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240 or 250 residues long.

[0373] PolyA tails may also be added after the construct is exported from the nucleus.

[0374] According to the present invention, terminal groups on the poly A tail may be incorporated for stabilization into polynucleotides of the invention (*e.g.*, antigen-encoding polynucleotides featured in the RNAVs of the invention). Polynucleotides of the present invention may incude des-3' hydroxyl tails. They may also include structural moieties or 2'-Omethyl modifications as taught by Junjie Li, et al. (Current Biology, Vol. 15, 1501-1507, August 23, 2005).

[0375] The polynucleotides of the present invention (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention) may be desiged to encode transcripts with alternative polyA tail structures including histone mRNA. According to Norbury, "Terminal uridylation has also been detected on human replication-dependent histone mRNAs. The turnover of these mRNAs is thought to be important for the prevention of potentially toxic histone accumulation following the completion or inhibition of chromosomal DNA replication. These mRNAs are distinguished by their lack of a 3' poly(A) tail, the function of which is instead assumed by a stable stem-loop structure and its cognate stem-loop binding protein (SLBP); the latter carries out the same functions as those of PABP on polyadenylated mRNAs" (Norbury, "Cytoplasmic RNA: a case of the tail wagging the dog," Nature Reviews Molecular Cell Biology; AOP, published online 29 August 2013; doi:10.1038/nrm3645).

[0376] Unique poly-A tail lengths provide certain advantages to the polynucleotides of the present invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention).

[0377] Generally, the length of a poly-A tail, when present, is greater than 30 nucleotides (SEQ ID NO:2280) in length In another embodiment, the poly-A fail is greater than 35 nucleotides (SEQ ID NO:2281) in length (e.g., at least or greater than about 35, 40, 45, 50, 55, 60, 70, 80, 90, 100, 120, 140, 160, 180, 200, 250, 300, 350, 400, 450, 500, 600, 700, 800, 900, 1,000, 1,100, 1,200, 1,300, 1,400, 1,500, 1,600, 1,700, 1,800, 1,900, 2,000, 2,500, and 3,000 nucleotides). In some embodiments, the polynucleotide or region thereof includes from about 30 to about 3,000 nucleotides (e.g., from 30 to 50, from 30 to 100, from 30 to 250, from 30 to 500, from 30 to 750, from 30 to 1,000, from 30 to 1,500, from 30 to 2,000, from 30 to 2,500, from 50 to 100, from 50 to 250, from 50 to 500, from 50 to 750, from 50 to 1,000, from 50 to 1,500, from 50 to 2,000, from 50 to 2,500, from 50 to 3,000, from 100 to 500, from 100 to 750, from 100 to 1,000, from 100 to 1,500, from 100 to 2,000, from 100 to 2,500, from 100 to 3,000, from 500 to 750, from 500 to 1,000, from 500 to 1,500, from 500 to 2,000, from 500 to 2,500, from 500 to 3,000, from 1,000 to 1,500, from 1,000 to 2,000, from 1,000 to 2,500, from 1,000 to 3,000, from 1,500 to 2,000, from 1,500 to 2,500, from 1,500 to 3,000, from 2,000 to 3,000, from 2,000 to 2,500, and from 2,500 to 3,000).

[0378] In one embodiment, the poly-A fail is designed relative to the length of the overall polynucleotide or the length of a particular region of the polynucleotide. This design may be based on the length of a coding region, the length of a particular feature or region or based on the length of the ultimate product expressed from the polynucleotides.

[0379] In this context the poly-A tail may be 10, 20, 30, 40, 50, 60, 70, 80, 90, or 100% greater in length than the polynucleotide or feature thereof. The poly-A fail may also be designed as a fraction of the polynucleotides to which it belongs. In this context, the poly-A tail may be 10, 20, 30, 40, 50, 60, 70, 80, or 90% or more of the total length of the

construct, a construct region or the total length of the construct minus the poly-A tail. Further, engineered binding sites and conjugation of polynucleotides for Poly-A binding protein may enhance expression.

[0380] Additionally, multiple distinct polynucleotides may be linked together via the PABP (Poly-A binding protein) through the 3'-end using modified nucleotides at the 3'-terminus of the poly-A tail. Transfection experiments can be conducted in relevant cell lines at and protein production can be assayed by ELISA at 12hr, 24hr, 48hr, 72 hr and day 7 post-transfection.

[0381] In one embodiment, the polynucleotides of the present invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) are designed to include a polyA-G Quartet region. The G-quartet is a cyclic hydrogen bonded array of four guanine nucleotides that can be formed by G-rich sequences in both DNA and RNA. In this embodiment, the G-quartet is incorporated at the end of the poly-A tail. The resultant polynucleotide is assayed for stability, protein production and other parameters including half-life at various time points It has been discovered that the polyA-G quartet results in protein production from an mRNA equivalent to at least 75% of that seen using a poly-A tail of 120 nucleotides (SEQ ID NO: 2282) alone.

Start codon region

[0382] In some embodiments, the polynucleotides of the present invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) may have regions that are analogous to or function like a start codon region.

[0383] In one embodiment, the translation of a polynucleotide may initiate on a codon which is not the start codon AUG. Translation of the polynucleotide may initiate on an alternative start codon such as, but not limited to, ACG, AGG, AAG, CTG/CUG, GTG/GUG, ATA/AUA, ATT/AUU, TTG/UUG (see Touriol et al. Biology of the Cell 95 (2003) 169-178 and Matsuda and Mauro PLoS ONE, 2010 5:11). As a non-limiting example, the translation of a polynucleotide begins on the alternative start codon ACG. As another non-limiting example, polynucleotide translation begins on the alternative start codon CTG or CUG. As yet another non-limiting example, the translation of a polynucleotide begins on the alternative start codon GTG or GUG.

[0384] Nucleotides flanking a codon that initiates translation such as, but not limited to, a start codon or an alternative start codon, are known to affect the translation efficiency, the length and/or the structure of the polynucleotide. (See e.g., Matsuda and Mauro PLoS ONE, 2010 5:11). Masking any of the nucleotides flanking a codon that initiates translation may be used to alter the position of translation initiation, translation efficiency, length and/or structure of a polynucleotide.

[0385] In one embodiment, a masking agent may be used near the start codon or alternative start codon in order to mask or hide the codon to reduce the probability of translation initiation at the masked start codon or alternative start codon. Non-limiting examples of masking agents include antisense locked nucleic acids (LNA) polynucleotides and exon-junction complexes (EJCs) (See e.g., Matsuda and Mauro describing masking agents LNA polynucleotides and EJCs (PLoS ONE, 2010 5:11);.

[0386] In another embodiment, a masking agent may be used to mask a start codon of a polynucleotide in order to increase the likelihood that translation will initiate on an alternative start codon.

[0387] In one embodiment, a masking agent may be used to mask a first start codon or alternative start codon in order to increase the chance that translation will initiate on a start codon or alternative start codon downstream to the masked start codon or alternative start codon.

[0388] In one embodiment, a start codon or alternative start codon may be located within a perfect complement for a miR binding site. The perfect complement of a miR binding site may help control the translation, length and/or structure of the polynucleotide similar to a masking agent. As a non-limiting example, the start codon or alternative start codon may be located in the middle of a perfect complement for a miR-122 binding site. The start codon or alternative start codon may be located after the first nucleotide, second nucleotide, third nucleotide, fourth nucleotide, fifth nucleotide, sixth nucleotide, seventh nucleotide, eighth nucleotide, ninth nucleotide, tenth nucleotide, eleventh nucleotide, twelfth nucleotide, thirteenth nucleotide, fourteenth nucleotide, fifteenth nucleotide, sixteenth nucleotide, seventeenth nucleotide, eighteenth nucleotide, nineteenth nucleotide, twentieth nucleotide or twenty-first nucleotide

[0389] In another embodiment, the start codon of a polynucleotide may be removed from the polynucleotide sequence in order to have the translation of the polynucleotide begin on a codon which is not the start codon. Translation of the polynucleotide may begin on the codon following the removed start codon or on a downstream start codon or an alternative start codon. In a non-limiting example, the start codon ATG or AUG is removed as the first 3 nucleotides of the polynucleotide sequence in order to have translation initiate on a downstream start codon or alternative start codon. The polynucleotide sequence where the start codon was removed may further comprise at least one masking agent for the downstream start codon and/or alternative start codons in order to control or attempt to control the initiation of translation, the length of the polynucleotide and/or the structure of the polynucleotide.

Stop Codon Region

[0390] In one embodiment, the polynucleotides of the present invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) may include at least two stop codons before the 3' untranslated region (UTR). The stop codon may be selected from TGA, TAA and TAG. In one embodiment, the polynucleotides of the present invention include the stop codon TGA and one additional stop codon. In a further embodiment the addition stop codon may be TAA. In another embodiment, the polynucleotides of the present invention include three stop codons.

Signal Sequences

[0391] The polynucleotides of the invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) may also encode additional features which facilitate trafficking of the polypeptides to therapeutically relevant sites. One such feature which aids in protein trafficking is the signal sequence. As used herein, a "signal sequence" or "signal peptide" is a polynucleotide or polypeptide, respectively, which is from about 9 to 200 nucleotides (3-60 amino acids) in length which is incorporated at the 5' (or N-terminus) of the coding region or polypeptide encoded, respectively. Addition of these sequences result in trafficking of the encoded polypeptide to the endoplasmic reticulum through one or more secretory pathways. Some signal peptides are cleaved from the protein by signal peptidase after the proteins are transported.

[0392] Additional signal sequences which may be utilized in the present invention include those taught in, for example, databases such as those found at http://www.signalpeptide.de/ or http://proline.bic.nus.edu.sg/spdb/. Those described in US Patents 8,124,379; 7,413,875 and 7,385,034 are also within the scope of the invention.

Protein Cleavage Signals and Sites

[0393] In exemplary embodiments, polypeptides of the invention (i.e., antigen polypeptides) may include various protein cleavage signals and/or sites.

[0394] In one embodiment, the polypeptides of the present invention may include at least one protein cleavage signal containing at least one protein cleavage site. The protein cleavage site may be located at the N-terminus, the C-terminus, at any space between the N- and the C- termini such as, but not limited to, half-way between the N- and C-termini, between the N-terminus and the half way point, between the half way point and the C-terminus, and combinations thereof.

[0395] In one embodiment, the polynucleotides of the present invention may be engineered such that the polynucleotide contains at least one encoded protein cleavage signal. The encoded protein cleavage signal may be located in any region including but not limited to before the start codon, after the start codon, before the coding region, within the coding region such as, but not limited to, half way in the coding region, between the start codon and the half way point, between the half way point and the stop codon, after the coding region, before the stop codon, between two stop codons, after the stop codon and combinations thereof.

[0396] In one embodiment, the polynucleotides of the present invention may include at least one encoded protein cleavage signal containing at least one protein cleavage site. The encoded protein cleavage signal may include, but is not limited to, a proprotein convertase (or prohormone convertase), thrombin and/or Factor Xa protein cleavage signal.

[0397] As a non-limiting example, U.S. Pat. No. 7,374,930 and U.S. Pub. No. 20090227660 use a furin cleavage site to cleave the N-terminal methionine of GLP-1 in the expression product from the Golgi apparatus of the cells. In one embodiment, the polypeptides of the present invention include at least one protein cleavage signal and/or site with the proviso that the polypeptide is not GLP-1.

Insertions and Substitutions

[0398] In exemplary embodiments, polynucleotides of the invention (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention) can include various substitutions and/or insertions.

[0399] In one embodiment, the 5'UTR of the polynucleotide may be replaced by the insertion of at least one region and/or string of nucleosides of the same base. The region and/or string of nucleotides may include, but is not limited to, at least 3, at least 4, at least 5, at least 6, at least 7 or at least 8 nucleotides and the nucleotides may be natural and/or unnatural. As a non-limiting example, the group of nucleotides may include 5-8 adenine, cytosine, thymine, a string of any of the other nucleotides disclosed herein and/or combinations thereof.

[0400] In one embodiment, the 5'UTR of the polynucleotide may be replaced by the insertion of at least two regions and/or strings of nucleotides of two different bases such as, but not limited to, adenine, cytosine, thymine, any of the other nucleotides disclosed herein and/or combinations thereof. For example, the 5'UTR may be replaced by inserting 5-8 adenine bases followed by the insertion of 5-8 cytosine bases. In another example, the 5'UTR may be replaced by inserting 5-8 cytosine bases followed by the insertion of 5-8 adenine bases.

[0401] In one embodiment, the polynucleotide may include at least one substitution and/or insertion downstream of the transcription start site which may be recognized by an RNA polymerase. As a non-limiting example, at least one substitution and/or insertion may occur downstream the transcription start site by substituting at least one nucleic acid in the region just downstream of the transcription start site (such as, but not limited to, +1 to +6). Changes to region of nucleotides just downstream of the transcription start site may affect initiation rates, increase apparent nucleotide triphosphate (NTP) reaction constant values, and increase the dissociation of short transcripts from the transcription complex curing initial transcription (Brieba et al, Biochemistry (2002) 41: 5144-5149). The modification, substitution and/or insertion of at least one nucleoside may cause a silent mutation of the sequence or may cause a mutation in the amino acid sequence.

[0402] In one embodiment, the polynucleotide may include the substitution of at least 1, at least 2, at least 3, at least 4, at least 5, at least 6, at least 7, at least 8, at least 9, at least 10, at least 11, at least 12 or at least 13 guanine bases downstream of the transcription start site.

[0403] In one embodiment, the polynucleotide may include the substitution of at least 1, at least 2, at least 3, at least 4, at least 5 or at least 6 guanine bases in the region just downstream of the transcription start site. As a non-limiting example, if the nucleotides in the region are GGGAGA the guanine bases may be substituted by at least 1, at least 2, at least 3 or at least 4 adenine nucleotides. In another non-limiting example, if the nucleotides in the region are GGGAGA the guanine bases may be substituted by at least 1, at least 2, at least 3 or at least 4 cytosine bases. In another non-limiting example, if the nucleotides in the region are GGGAGA the guanine bases may be substituted by at least 1, at least 2, at least 3 or at least 4 thymine, and/or any of the nucleotides described herein.

[0404] In one embodiment, the polynucleotide may include at least one substitution and/or insertion upstream of the start codon. For the purpose of clarity, one of skill in the art would appreciate that the start codon is the first codon of the protein coding region whereas the transcription start site is the site where transcription begins. The polynucleotide may include, but is not limited to, at least 1, at least 2, at least 3, at least 4, at least 5, at least 6, at least 7 or at least 8 substitutions and/or insertions of nucleotide bases. The nucleotide bases may be inserted or substituted at 1, at least 1, at least 2, at least 3, at least 4 or at least 5 locations upstream of the start codon. The nucleotides inserted and/or substituted may be the same base (e.g., all A or all C or all T or all G), two different bases (e.g., A and C, A and T, or C and T), three different bases (e.g., A, C and T or A, C and T) or at least four different bases. As a non-limiting example, the guanine base upstream of the coding region in the polynucleotide may be substituted with adenine, cytosine, thymine, or any of the nucleotides described herein. In another non-limiting example the substitution of guanine bases in the polynucleotide may be designed so as to leave one guanine base in the region downstream of the transcription start site and before the start codon (see Esvelt et al. Nature (2011) 472(7344):499-503). As a non-limiting example, at least 5 nucleotides may be inserted at 1 location downstream of the transcription start site but upstream of the start codon and the at least 5 nucleotides may be the same base type.

Incorporating Post Transcriptional Control Modulators

[0405] In one embodiment, the polynucleotides of the present invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) may include at least one post transcriptional control modulator. These post transcriptional control modulators may be, but are not limited to, small molecules, compounds and regulatory sequences. As a non-limiting example, post transcriptional control may be achieved using small molecules identified by PTC Therapeutics Inc. (South Plainfield, NJ) using their GEMS™ (Gene Expression Modulation by Small-Moleclues) screening technology.

[0406] The post transcriptional control modulator may be a gene expression modulator which is screened by the method detailed in or a gene expression modulator described in International Publication No. WO2006022712. Methods identifying RNA regulatory sequences involved in translational control are described in International Publication No. WO2004067728; methods identifying compounds that modulate untranslated region dependent expression of a gene are described in International Publication No. WO2004065561.

[0407] In one embodiment, the polynucleotides of the present invention may include at least one post transcriptional control modulator is located in the 5' and/or the 3' untranslated region of the polynucleotides of the present invention.

[0408] In another embodiment, the polynucleotides of the present invention may include at least one post transcription control modulator to modulate premature translation termination. The post transcription control modulators may be compounds described in or a compound found by methods outlined in International Publication Nso. WO2004010106, WO2006044456, WO2006044682, WO2006044503 and WO2006044505. As a non-limiting example, the compound may bind to a region of the 28S ribosomal RNA in order to modulate premature translation termination (See e.g., WO2004010106).

[0409] In one embodiment, polynucleotides of the present invention may include at least one post transcription control modulator to alter protein expression. As a non-limiting example, the expression of VEGF may be regulated using the compounds described in or a compound found by the methods described in International Publication Nos. WO2005118857, WO2006065480, WO2006065479 and WO2006058088.

[0410] The polynucleotides of the present invention may include at least one post transcription control modulator to control translation. In one embodiment, the post transcription control modulator may be a RNA regulatory sequence. As a non-limiting example, the RNA regulatory sequence may be identified by the methods described in International Publication No. WO2006071903.

Codon Optimization

[0411] The polynucleotides contained in the NAVs of the invention, their regions or parts or subregions may be codon optimized. Codon optimization methods are known in the art and may be useful in efforts to achieve one or more of several goals. These goals include to match codon frequencies in target and host organisms to ensure proper folding, bias GC content to increase mRNA stability or reduce secondary structures, minimize tandem repeat codons or base runs that may impair gene construction or expression, customize transcriptional and translational control regions, insert or remove protein trafficking sequences, remove/add post translation modification sites in encoded protein (e.g. glycosylation sites), add, remove or shuffle protein domains, insert or delete restriction sites, modify ribosome binding sites and mRNA degradation sites, to adjust translational rates to allow the various domains of the protein to fold properly, or to reduce or eliminate problem secondary structures within the polynucleotide. Codon optimization tools, algorithms and services are known in the art, non-limiting examples include services from GeneArt (Life Technologies), DNA2.0 (Menlo Park CA) and/or proprietary methods. In one embodiment, the ORF sequence is optimized using optimization algorithms. Codon options for each amino acid are given in Table 21.

Table 21. Codon Options

Amino Acid	Single Letter Code	Codon Options
Isoleucine	I	ATT, ATC, ATA
Leucine	L	CTT, CTC, CTA, CTG, TTA, TTG
Valine	V	GTT, GTC, GTA, GTG
Phenylalanine	F	TTT, TTC
Methionine	M	ATG
Cysteine	C	TGT, TGC
Alanine	A	GCT, GCC, GCA, GCG
Glycine	G	GGT, GGC, GGA, GGG
Proline	P	CCT, CCC, CCA, CCG
Threonine	T	ACT, ACC, ACA, ACG
Serine	S	TCT, TCC, TCA, TCG, AGT, AGC
Tyrosine	Y	TAT, TAC
Tryptophan	W	TGG
Glutamine	Q	CAA, CAG
Asparagine	N	AAT, AAC
Histidine	H	CAT, CAC
Glutamic acid	E	GAA, GAG
Aspartic acid	D	GAT, GAC
Lysine	K	AAA, AAG
Arginine	R	CGT, CGC, CGA, CGG, AGA, AGG
Selenocysteine	Sec	UGA in mRNA in presence of Selenocystein insertion element (SECIS)
Stop codons	Stop	TAA, TAG, TGA

[0412] Features, which may be considered beneficial in some embodiments of the present invention, may be encoded by regions of the polynucleotide and such regions may be upstream (5') or downstream (3') to a region which encodes a polypeptide. These regions may be incorporated into the polynucleotide before and/or after codon optimization of the protein encoding region or open reading frame (ORF). It is not required that a polynucleotide contain both a 5' and 3' flanking region. Examples of such features include, but are not limited to, untranslated regions (UTRs), Kozak sequences, an oligo(dT) sequence, and detectable tags and may include multiple cloning sites which may have XbaI recognition

[0413] In some embodiments, a 5' UTR and/or a 3' UTR region may be provided as flanking regions Multiple 5' or 3' UTRs may be included in the flanking regions and may be the same or of different sequences Any portion of the flanking regions, including none, may be codon optimized and any may independently contain one or more different structural

or chemical modifications, before and/or after codon optimization.

[0414] After optimization (if desired), the polynucleotides components are reconstituted and transformed into a vector such as, but not limited to, plasmids, viruses, cosmids, and artificial chromosomes. For example, the optimized polynucleotide may be reconstituted and transformed into chemically competent *E. coli,* yeast, neurospora, maize, drosophila, etc. where high copy plasmid-like or chromosome structures occur by methods described herein.

[0415] Synthetic polynucleotides and their nucleic acid analogs play an important role in the research and studies of biochemical processes. Various enzyme-assisted and chemical-based methods have been developed to synthesize polynucleotides and nucleic acids, in particular, polynucleotides and nucleic acids featured in the NAVs of the invention, as described infra.

Synthesis: Enzymatic Methods

In Vitro Transcription-enzymatic synthesis

[0416] cDNA encoding the polynucleotides described herein may be transcribed using an *in vitro* transcription (IVT) system. The system typically comprises a transcription buffer, nucleotide triphosphates (NTPs), an RNase inhibitor and a polymerase. The NTPs may be manufactured in house, may be selected from a supplier, or may be synthesized as described herein. The NTPs may be selected from, but are not limited to, those described herein including natural and unnatural (modified) NTPs. The polymerase may be selected from, but is not limited to, T7 RNA polymerase, T3 RNA polymerase and mutant polymerases such as, but not limited to, polymerases able to incorporate polynucleotides (e.g., modified nucleic acids).

RNA Polymerases useful for synthesis

[0417] Any number of RNA polymerases or variants may be used in the synthesis of the polynucleotides of the present invention.

[0418] RNA polymerases may be modified by inserting or deleting amino acids of the RNA polymerase sequence. As a non-limiting example, the RNA polymerase may be modified to exhibit an increased ability to incorporate a 2'-modified nucleotide triphosphate compared to an unmodified RNA polymerase (see International Publication WO2008078180 and U.S. Patent 8,101,385).

[0419] Variants may be obtained by evolving an RNA polymerase, optimizing the RNA polymerase amino acid and/or nucleic acid sequence and/or by using other methods known in the art. As a non-limiting example, T7 RNA polymerase variants may be evolved using the continuous directed evolution system set out by Esvelt et al. (Nature (2011) 472(7344):499-503) where clones of T7 RNA polymerase may encode at least one mutation such as, but not limited to, lysine at position 93 substituted for threonine (K93T), I4M, A7T, E63V, V64D, A65E, D66Y, T76N, C125R, S128R, A136T, N165S, G175R, H176L, Y178H, F182L, L196F, G198V, D208Y, E222K, S228A, Q239R, T243N, G259D, M267I, G280C, H300R, D351A, A354S, E356D, L360P, A383V, Y385C, D388Y, S397R, M401T, N410S, K450R, P451T, G452V, E484A, H523L, H524N, G542V, E565K, K577E, K577M, N601S, S684Y, L699I, K713E, N748D, Q754R, E775K, A827V, D851N or L864F. As another non-limiting example, T7 RNA polymerase variants may encode at least mutation as described in U.S. Pub. Nos. 20100120024 and 20070117112. Variants of RNA polymerase may also include, but are not limited to, substitutional variants, conservative amino acid substitution, insertional variants, deletional variants and/or covalent derivatives

[0420] In one embodiment, the polynucleotide may be designed to be recognized by the wild type or variant RNA polymerases. In doing so, the polynucleotide may be modified to contain sites or regions of sequence changes from the wild type or parent polynucleotide.

[0421] Polynucleotide or nucleic acid synthesis reactions may be carried out by enzymatic methods utilizing polymerases. Polymerases catalyze the creation of phosphodiester bonds between nucleotides in a polynucleotide or nucleic acid chain. Currently known DNA polymerases can be divided into different families based on amino acid sequence comparison and crystal structure analysis. DNA polymerase I (pol I) or A polymerase family, including the Klenow fragments of *E. Coli, Bacillus* DNA polymerase I, *Thermus aquaticus* (Taq) DNA polymerases, and the T7 RNA and DNA polymerases, is among the best studied of these families. Another large family is DNA polymerase α (pol α) or B polymerase family, including all eukaryotic replicating DNA polymerases and polymerases from phages T4 and RB69. Although they employ similar catalytic mechanism, these families of polymerases differ in substrate specificity, substrate analog-incorporating efficiency, degree and rate for primer extension, mode of DNA synthesis, exonuclease activity, and sensitivity against inhibitors.

[0422] DNA polymerases are also selected based on the optimum reaction conditions they require, such as reaction temperature, pH, and template and primer concentrations. Sometimes a combination of more than one DNA polymerases is employed to achieve the desired DNA fragment size and synthesis efficiency. For example, Cheng et al. increase pH,

add glycerol and dimethyl sulfoxide, decrease denaturation times, increase extension times, and utilize a secondary thermostable DNA polymerase that possesses a 3' to 5' exonuclease activity to effectively amplify long targets from cloned inserts and human genomic DNA (Cheng et al., PNAS, Vol. 91, 5695-5699 (1994)). RNA polymerases from bacteriophage T3, T7, and SP6 have been widely used to prepare RNAs for biochemical and biophysical studies. RNA polymerases, capping enzymes, and poly-A polymerases are disclosed in the copending application No. PCT/US2013/054635 (M032).

[0423] In one instance, the RNA polymerase which may be used in the synthesis of the chimeric polynucleotides described herein is a Syn5 RNA polymerase (see Zhu et al. Nucleic Acids Research 2013). The Syn5 RNA polymerase was recently characterized from marine cyanophage Syn5 by Zhu et al. where they also identified the promoter sequence (see Zhu et al. Nucleic Acids Research 2013). Zhu et al. found that Syn5 RNA polymerase catalyzed RNA synthesis over a wider range of temperatures and salinity as compared to T7 RNA polymerase. Additionally, the requirement for the initiating nucleotide at the promoter was found to be less stringent for Syn5 RNA polymerase as compared to the T7 RNA polymerase making Syn5 RNA polymerase promising for RNA synthesis.

[0424] In one instance, a Syn5 RNA polymerase may be used in the synthesis of the chimeric polynucleotides described herein. As a non-limiting example, a Syn5 RNA polymerase may be used in the synthesis of the chimeric polynucleotide requiring a precise 3'-termini.

[0425] In one instance, a Syn5 promoter may be used in the synthesis of the chimeric polynucleotides. As a non-limiting example, the Syn5 promoter may be 5'-ATTGGGCACCCGTAAGGG-3' (SEQ ID NO:2041) as described by Zhu et al. (Nucleic Acids Research 2013).

[0426] In one instance, a Syn5 RNA polymerase may be used in the synthesis of chimeric polynucleotides comprising at least one chemical modification described herein and/or known in the art. (see e.g., the incorporation of pseudo-UTP and 5Me-CTP described in Zhu et al. Nucleic Acids Research 2013).

[0427] In one instance, the chimeric polynucleotides described herein may be synthesized using a Syn5 RNA polymerase which has been purified using modified and improved purification procedure described by Zhu et al. (Nucleic Acids Research 2013).

[0428] Various tools in genetic engineering are based on the enzymatic amplification of a target gene which acts as a template. For the study of sequences of individual genes or specific regions of interest and other research needs, it is necessary to generate multiple copies of a target gene from a small sample of polynucleotides or nucleic acids. Such methods may be applied in the manufacture of the polynucleotides of the disclosure.

[0429] Polymerase chain reaction (PCR) has wide applications in rapid amplification of a target gene, as well as genome mapping and sequencing. The key components for synthesizing DNA comprise target DNA molecules as a template, primers complementary to the ends of target DNA strands, deoxynucleoside triphosphates (dNTPs) as building blocks, and a DNA polymerase. As PCR progresses through denaturation, annealing and extension steps, the newly produced DNA molecules can act as a template for the next circle of replication, achieving exponentially amplification of the target DNA PCR requires a cycle of heating and cooling for denaturation and annealing. Variations of the basic PCR include, but are not limited to, asymmetric PCR (See e.g., Innis et al., PNAS, vol. 85, 9436-9440 (1988)), inverse PCR (see e.g., Ochman et al., Genetics, vol. 120(3), 621-623, (1988)), and reverse transcription PCR (RT-PCR) (see e.g., Freeman et al., BioTechniques, vol. 26(1), 112-22,124-5 (1999)). In RT-PCR, a single stranded RNA is the desired target and is converted to a double stranded DNA first by reverse transcriptase.

[0430] A variety of isothermal *in vitro* nucleic acid amplification techniques have been developed as alternatives or complements of PCR. For example, strand displacement amplification (SDA) is based on the ability of a restriction enzyme to form a nick (Walker et al., PNAS, vol. 89, 392-396 (1992)). A restriction enzyme recognition sequence is inserted into an annealed primer sequence. Primers are extended by a DNA polymerase and dNTPs to form a duplex. Only one strand of the duplex is cleaved by the restriction enzyme. Each single strand chain is then available as a template for subsequent synthesis SDA does not require the complicated temperature control cycle of PCR.

[0431] Nucleic acid sequence-based amplification (NASBA), also called transcription mediated amplification (TMA), is also an isothermal amplification method that utilizes a combination of DNA polymerase, reverse transcriptase, RNAse H, and T7 RNA polymerase (Compton, Nature, vol. 350, 91-92 (1991)). A target RNA is used as a template and a reverse transcriptase synthesizes its complementary DNA strand. RNAse H hydrolyzes the RNA template, making space for a DNA polymerase to synthesize a DNA strand complementary to the first DNA strand which is complementary to the RNA target, forming a DNA duplex. T7 RNA polymerase continuously generates complementary RNA strands of this DNA duplex. These RNA strands act as templates for new cycles of DNA synthesis, resulting in amplification of the target gene.

[0432] Rolling-circle amplification (RCA) amplifies a single stranded circular polynucleotide and involves numerous rounds of isothermal enzymatic synthesis where Φ29 DNA polymerase extends a primer by continuously progressing around the polynucleotide circle to replicate its sequence over and over again. Therefore, a linear copy of the circular template is achieved. A primer can then be annealed to this linear copy and its complementary chain can be synthesized (Lizardi et al., Nature Genetics, vol. 19, 225-232 (1998)). A single stranded circular DNA can also serve as a template

for RNA synthesis in the presence of an RNA polymerase (Daubendiek et al., JACS, vol. 117, 7818-7819 (1995)). An inverse rapid amplification of cDNA ends (RACE) RCA is described by Polidoros et al. (BioTechniques, vol. 41, 35-42 (2006)). A messenger RNA (mRNA) is reverse transcribed into cDNA, followed by RNAse H treatment to separate the cDNA. The cDNA is then circularized by CircLigase into a circular DNA. The amplification of the resulting circular DNA is achived with RCA.

[0433] Any of the foregoing methods may be utilized in the manufacture of one or more regions of the polynucleotides of the present invention (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention).

[0434] Assembling polynucleotides or nucleic acids (*e.g.*, antigen-encoding polynucleotides or nucleic acids) by a ligase is also widely used. DNA or RNA ligases promote intermolecular ligation of the 5' and 3' ends of polynucleotide chains through the formation of a phosphodiester bond. Ligase chain reaction (LCR) is a promising diagnosing technique based on the principle that two adjacent polynucleotide probes hybridize to one strand of a target gene and couple to each other by a ligase. If a target gene is not present, or if there is a mismatch at the target gene, such as a single-nucleotide polymorphism (SNP), the probes cannot ligase (Wiedmann et al., PCR Methods and Application, vol.3 (4), s51-s64 (1994)). LCR may be combined with various amplification techniques to increase sensitivity of detection or to increase the amount of products if it is used in synthesizing polynucleotides and nucleic acids.

[0435] Several library preparation kits for nucleic acids are now commercially available. They include enzymes and buffers to convert a small amount of nucleic acid samples into an indexed library for downstream applications. For example, DNA fragments may be placed in a NEBNEXT® ULTRA™ DNA Library Prep Kit by NEWENGLAND BIOLABS® for end preparation, ligation, size selection, clean-up, PCR amplification and final clean-up.

[0436] Continued development is going on to improvement the amplification techniques. For example, US Pat. 8,367,328 to Asada et al. teaches utilizing a reaction enhancer to increase the efficiency of DNA synthesis reactions by DNA polymerases. The reaction enhancer comprises an acidic substance or cationic complexes of an acidic substance. US Pat. 7.384,739 to Kitabayashi et al. teaches a carboxylate ion-supplying substance that promotes enzymatic DNA synthesis, wherein the carboxylate ioin-supplying substance is selected from oxalic acid, malonic acid, esters of oxalic acid, esters of malonic acid, salts of malonic acid, and esters of maleic acid. US Pat. 7,378,262 to Sobek et al. discloses an enzyme composition to increase fidelity of DNA amplifications. The composition comprises one enzyme with 3' exonuclease activity but no polymerase activity and another enzyme that is a polymerase. Both of the enzymes are thermostable and are reversibly modified to be inactive at lower temperatures.

[0437] US Pat. No. 7,550,264 to Getts et al. teaches multiple round of synthesis of sense RNA molecules are performed by attaching oligodeoxynucleotides tails onto the 3' end of cDNA molecules and initiating RNA transcription using RNA polymerase. US Pat. Publication No. 2013/0183718 to Rohayem teaches RNA synthesis by RNA-dependent RNA polymerases (RdRp) displaying an RNA polymerase activity on single-stranded DNA templates,. Oligonucleotides with non-standard nucleotides may be synthesized with enzymatic polymerization by contacting a template compring non-standard nucleotides with a mixture of nucleotides that are complementary to the nucleotides of the template as disclosed in US Pat. No. 6,617,106 to Benner.

Synthesis: Solid-phase chemical synthesis

[0438] Chimeric polynucleotides or circular polynucleotides of the present invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) may be manufactured in whole or in part using solid phase techniques.

[0439] Solid-phase chemical synthesis of polynucleotides or nucleic acids is an automated method wherein molecules are immobilized on a solid support and synthesized step by step in a reactant solution. Impurities and excess reagents are washed away and no purification is required after each step. The automation of the process is amenable on a computer-controlled solid-phase synthesizer. Solid-phase synthesis allows rapid production of polynucleotides or nucleic acids in a relatively large scale that leads to the commercial availability of some polynucleotides or nucleic acids. Furthermore, it is useful in site-specific introduction of chemical modifications in the polynucleotide or nucleic acid sequences. It is an indispensable tool in designing modified derivatives of natural nucleic acids.

[0440] In automated solid-phase synthesis, the chain is synthesized in 3' to 5' direction. The hydroxyl group in the 3' end of a nucleoside is tethered to a solid support via a chemically cleavable or light-cleavable linker. Activated nucleoside monomers, such as 2'-deoxynucleosides (dA, dC, dG and T), ribonucleosides (A, C, G, and U), or chemically modified nucleosides, are added to the support-bound nucleoside sequentially. Currently most widely utilized monomers are the 3'-phophoramidite derivatives of nucleoside building blocks. The 3' phosphorus atom of the activated monomer couples with the 5' oxygen atom of the support-bound nucleoside to form a phosphite triester. To prevent side reactions, all functional groups not involved in the coupling reaction, such as the 5' hydroxyl group, the hydroxyl group on the 3' phosphorus atom, the 2' hydroxyl group in ribonucleosides monomers, and the amino groups on the purine or pyrimidine bases, are all blocked with protection groups. The next step involves oxidation of the phosphite triester to form a phosphate triester or phosphotriester, where the phosphorus atom is pentavalent. The protection group on the 5' hydroxyl group at the end of the growing chain is then removed, ready to couple with an incoming activated monomer building block.

At the end of the synthesis, a cleaving agent such as ammonia or ammonium hydroxide is added to remove all the protecting groups and release the polynucleotide chains from the solid support. Light may also be applied to cleave the polynucleotide chain. The product can then be further purified with high pressure liquid chromatography (HPLC) or electrophoresis.

[0441] In solid-phase synthesis, the polynucleotide chain is covalently bound to the solid support via its 3' hydroxyl group. The solid supports are insoluble particles also called resins, typically 50-200 μm in diameter Many different kinds of resins are now available, as reviewed in "Solid-phase supports for polynucleotide synthesis" by Guzaev (Guzaev, Current Protocols in Nucleic Acid Chemistry, 3.1.1-3.1.60 (2013)). The most common materials for the resins include highly cross-linked polystyrene beads and controlled pore glass (CPG) beads. The surface of the beads may be treated to have functional groups, such as amino or aminomethyl groups that can be used as anchoring points for linkers to tether nucleosides. They can be implemented in columns, multi-well plates, microarrays or microchips. The column-based format allows relatively large scale synthesis of the polynucleotides or nucleic acids. The resins are held between filters in columns that enable all reagents and solvents to pass through freely. Multi-well plates, microarrays, or microchips are designed specifically for cost-effective small scale synthesis. Up to a million polynucleotides can be produced on a single microarray chip. However, the error rates of microchip-based synthesis are higher than traditional column-based methods (Borovkov et al., Nucleic Acids Research, vol. 38(19), e180 (2010)). Multi-well plates allow parallel synthesis of polynucleotides or nucleic acids with different sequences simultaneously (Sindelar, et al., Nucleic Acids Research, vol. 23, 982-987 (1995)). The loading on the solid supports is limited In addition, as the extension progresses, the morphology and bulkiness of the growing chains on the solid supports might hinder the incoming monomers from reacting with the terminal group of the growing chains. Therefore, the number of monomers that can be added to the growing chain is also limited

[0442] Linkers are attached to the solid support for further extension of the chain. They are stable to all the reagents used in the synthesis process, except in the end of the synthesis when the chain is detached from the solid support. Solid supports with a specific nucleoside linker, i.e., A, C, dT, G, or U, can be used to prepare polynucleotides with A, C, T, G, or U as the first nucleotide in the sequence, respectively. Universal solid supports with non-nucleoside linkers can be used for all polynucleotide sequences (US Pat. 6,653,468 to Guzaev et al.). Various non-nucleoside linkers have been developed for universal supports, a lot of them with two vicinal hydroxyl groups. For example, a succinyl group is a frequently used linker.

[0443] As used herein, a linker refers to a group of atoms, e.g., 10-1,000 atoms, and can be comprised of the atoms or groups such as, but not limited to, carbon, amino, alkylamino, oxygen, sulfur, sulfoxide, sulfonyl, carbonyl, and imine. The linker can be attached to a modified nucleoside or nucleotide on the nucleobase or sugar moiety. A linker may be nucleic acid based or non-nucleosidic. The linker may be of sufficient length as to not interfere with incorporation into a nucleic acid sequence. The linker can be used for any useful purpose, such as to form multimers (e.g., through linkage of two or more chimeric polynucleotides molecules) or conjugates, as well as to administer a therapeutic molecule or incorporate a label, as described herein. Examples of chemical groups that can be incorporated into the linker include, but are not limited to, alkyl, alkenyl, alkynyl, amido, amino, ether, thioether, ester, alkylene, heteroalkylene, aryl, or heterocyclyl, each of which can be optionally substituted, as described herein. Examples of linkers include, but are not limited to, unsaturated alkanes, polyethylene glycols (e.g., ethylene or propylene glycol monomeric units, e.g., diethylene glycol, dipropylene glycol, triethylene glycol, tripropylene glycol, tetraethylene glycol, or tetraethylene glycol), and dextran polymers and derivatives thereof, Other examples include, but are not limited to, cleavable moieties within the linker, such as, for example, a disulfide bond (-S-S-) or an azo bond (-N=N-), which can be cleaved using a reducing agent or photolysis. Non-limiting examples of a selectively cleavable bond include an amido bond can be cleaved for example by the use of tris(2-carboxyethyl)phosphine (TCEP), or other reducing agents, and/or photolysis, as well as an ester bond can be cleaved for example by acidic or basic hydrolysis.)

[0444] Besides the functional groups on the activated monomer and the growing chain needed for the coupling reaction to extend the chain, all other functional groups need to be protected to avoid side reactions. The conditions for protection and deprotection, and the selection of appropriate protecting groups can be readily determined by one skilled in the art. The chemistry of protecting groups can be found and/or described, for example, in Greene, et al.(Protective Groups in Organic Synthesis, 2d. Ed., Wiley & Sons, 1991.) For example, the 5' hydroxyl group on the activated nucleoside phosphoramidite monomers may be protected with 4,4'-dimethoxytrityil (DMT) and the hydroxyl group on the phosphorus atom may be protected with 2-cyanoethyl. The exocyclic amino groups on the A, C, G bases may be protected with acyl groups.

[0445] In a solid-phase synthesis system, the reactivity of the activated monomers is important, because of the heterogeneity of the media. A majority of solid-phase synthesis uses phosphoramidite nucleosides, the mechanism of which is discussed above. Another activated monomer example is nucleoside *H*-phosphonates (Abramova, Molecules, vol. 18, 1063-1075 (2013)). A large excess of reagents, such as monomers, oxidizing agents, and deprotection agents, is required in order to ensure high yields in the solid-phase synthesis system.

[0446] Scientific studies and research are going on to further improve the solid-phase synthesis method. For example,

instead of the well-established 3'-to-5' synthesis, US Pat. No. 8,309,707 and US Pat. Publication No. 2013/0072670 to Srivastava et al. disclosed a 5'-to-3' synthesis of RNA utilizing a novel phosphoramidite and a novel nucleoside derivative, thereby allowing easy modifications of the synthetic RNA at the 3' end. PCT application WO2013123125 to Church et al. describes assembly of a target nucleic acid sequence from a plurality of subsequences, wherein resins with the subsequences are placed in an emulsion droplet. The subsequences are cleaved off the resins and assemble within the emulsion droplet. To reduce the cost of solid supports, a reusable CPG solid support has been developed with a hydroquinone-O,O'-diacetic acid linker (Q-linker) (Pon et al., Nucleic Acid Research, vol. 27, 1531-1538 (1999)).

[0447] New protecting groups for solid-phase synthesis have also been developed. Nagat et al. has successfully synthesized 110-nt-long RNA with the sequence of a candidate precursor microRNA by using 2-cyanoethoxymethyl (CEM) as the 2'-hydroxy protecting group (Shiba et al., Nucleic Acids Research, vol. 35, 3287-3296 (2007)). Also with CEM as 2'-O-protecting group, a 130-nt mRNA has been synthesized encoding a 33-amino acid peptide that includes the sequence of glucagon-like peptide-1 (GLP-1). The biological activity of the artificial 130-nt mRNA is shown by producing GLP-1 in a cell-free protein synthesis system and in Chinese hamster ovary (CHO) cells (Nagata et al., Nucleic Acids Research, vol. 38(21), 7845-7857 (2010)). Novel protecting groups for solid-phase synthesis monomers include, but are not limited to, carbonate protecting group disclosed in US Pat. No. 8,309,706 to Dellinger et al., orthoester-type 2' hydroxyl protecting group and an acyl carbonate-type hydroxyl protecting group disclosed in US Pat. No. 8,242,258 to Dellinger et al., 2'-hydroxyl thiocarbon protecting group disclosed in US Pat. No. 8,202,983 to Dellinger et al., 2'-silyl containing thiocarbonate protecting group disclosed in US Pat. No. 7,999,087 to Dellinger et al., 9-fluorenylmethoxycarbonyl (FMOS) derivatives as an amino protecting group disclosed in US Pat. No. 7,667,033 to Alvarado, fluoride-labile 5'silyl protecting group disclosed in US Pat. No. 5,889,136 to Scaringe et al., and pixyl protecting groups disclosed in US Pat. Publication No. 2008/0119645 to Griffey et al.. US Pat. Publication No. 2011/0275793 to Debart et al. teaches RNA synthesis using a protecting group of the hyoxyls in position 2' of the ribose that can be removed by a base. Novel solid supports include polymers made from monomers comprising protected hydroxypolyC_{2-4}alkyleneoxy chain attached to a polymerizable unit taught in US Pat. No. 7.476,709 to Moody et al..

Synthesis: Liquid Phase Chemical Synthesis

[0448] The synthesis of chimeric polynucleotides or circular polynucleotides of the present invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) by the sequential addition of monomer building blocks may be carried out in a liquid phase. A covalent bond is formed between the monomers or between a terminal functional group of the growing chain and an incoming monomer. Functional groups not involved in the reaction must be temporarily protected. After the addition of each monomer building block, the reaction mixture has to be purified before adding the next monomer building block. The functional group at one terminal of the chain has to be deprotected to be able to react with the next monomer building blocks. A liquid phase synthesis is labor- and time-consuming and cannot not be automated. Despite the limitations, liquid phase synthesis is still useful in preparing short polynucleotides in a large scale. Because the system is homogenous, it does not require a large excess of reagents and is cost-effective in this respect.

Synthesis: Combination of Synthetic Methods

[0449] The synthetic methods discussed above each has its own advantages and limitations. Attempts have been conducted to combine these methods to overcome the limitations. Such combinations of methods are within the scope of the present disclosure.

[0450] Short polynucleotide chains with 2-4 nucleotides may be prepared in liquid phase followed by binding to a solid support for extension reactions by solid phase synthesis. A high efficiency liquid phase (HELP) synthesis is developed that uses monomethyl ether of polyethylene glycol (MPEG) beads as a support for the monomer building blocks. MPEG is soluble in methylene chloride and pyridine solvents but precipitates in a diethyl ether solvent. By choosing an appropriate solvent, the coupling reaction between monomers or between a growing chain and an incoming monomer bound on MPEG can be carried out in a homogenous liquid phase system. The mixture can then be washed with a diethyl ether solvent to easily precipitate and purify the product (Bonora et al., Nucleic Acids Research, vol. 18, 3155-3159 (1990)). US Pat. No. 8,304,532 to Adamo et al. teaches a solution phase oligonucleotide synthesis where at least some of the reagents are solid supported.

[0451] The use of solid-phase or liquid-phase chemical synthesis in combination with enzymatic ligation provides an efficient way to generate long chain polynucleotides that cannot be obtained by chemical synthesis alone. Moore and Sharp describe preparing RNA fragments 10- to 20-nt long by chemical synthesis, to which site-specific modifications may be introduced, annealing the fragments to a cDNA bridge, and then assemble the fragments with T4 DNA ligase (Moore et al., Science, vol. 256, 992-997 (1992)).

[0452] A solid-phase synthesizer may produce enough polynucleotides or nucleic acids with good purity to preform PCR and other amplification techniques. Agilent Technologies have developed microarrays that are commercially avail-

able. Polynucleotides may be synthesized on a microarray substrate, cleaved by a strong base or light, followed by PCR amplification to generate a library of polynucleotides (Cleary et al., Nature Methods, vol. 1(3), 241-247 (2004)).

Synthesis: Small Region Synthesis

[0453] Regions or subregions of the polynucleotides of the present invention (e.g., antigen-encoding polynucleotides featured in the NAVs of the invention) may comprise small RNA molecules such as siRNA, and therefore may be synthesized in the same manner. There are several methods for preparing siRNA, such as chemical synthesis using appropriately protected ribonucleoside phosphoramidites, *in vitro* transcription, siRNA expression vectors, and PCR expression cassettes. Sigma-Aldrich® is one of the siRNA suppliers and synthesizes their siRNA using ribonucleoside phosphoramidite monomers protected at the 2' position with a t-butylmethylsilyl (TBDMS) group. The solid-phase chemical synthesis is carried out with SIGMA-ALDRICH®'s Ultra Fast Parallel Synthesis (UFPS) and Ultra Fast Parallel Deprotection (UFPD) to achieve high coupling efficiency and fast deprotection. The final siRNA products may be purified with HPLC or PAGE. Such methods may be used to synthesize regions or subregions of chimeric polynucleotides.

[0454] *In vitro* transcription and expression from a vector or a PCR-generated siRNA cassette require appropriate templates to produce siRNAs. The commercially available AMBIONO SILENCER® siRNA construction kit produces siRNA by *in vitro* transcription of DNA templates and contains the enzymes, buffers, primers needed. Such methods may be used to synthesize regions or subregions of chimeric polynucleotides.

Synthesis: Ligation of Polynucleotide Regions or Subregions

[0455] Polynucleotides of the invention (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention) such as chimeric polynucleotides and/or circular polynucleotides may be prepared by ligation of one or more regions or subregions.

[0456] Ligation is an indispensable tool for assembling polynucleotide or nucleic acid fragments into larger constructs DNA fragments can be joined by a ligase catalyzed reaction to create recombinant DNA with different functions. Two oligodeoxynucleotides, one with a 5' phosphoryl group and another with a free 3' hydroxyl group, serve as substrates for a DNA ligase. Oligodexoynucleotides with fluorescent or chemiluminescent labels may also serve as DNA ligase substrates (Martinelli et al., Clinical Chemistry, vol. 42, 14-18 (1996)). RNA ligases such as T4 RNA ligase catalyze the formation of a phosphodiester bond between two single stranded oligoribonucleotides or RNA fragments. Copies of large DNA constructs have been synthesized with a combination of polynucleotide fragments, thermostable DNA polymerases, and DNA ligases. US Pat. Publication No. 2009/0170090 to Ignatov et al. discloses improving PCT, especially enhancing yield of a long distance PCR and/or a low copy DNA template PCR amplication, by using a DNA ligase in addition to a DNA polymerase.

[0457] Ligases may be used with other enzymes to prepare desired chimeric polynucleotide or nucleic acid molecules and to perform genome analysis. For example, ligation-mediated selective PCR amplification is disclosed in EP Pat. Pub. No. 0735144 to Kato. Complementary DNAs (cDNAs) reverse-transcribed from tissue- or cell-derived RNA or DNA are digested into fragments with type IIS restriction enzymes. Biotinylated adapter sequences are attached to the fragments by *E. coli* DNA ligases. The biotin-labeled DNA fragments are then immobilized onto streptavidin-coated beads for downstream analysis.

[0458] A ligation splint or a ligation splint oligo is an oligonucleotide that is used to provide an annealing site or a ligation template for joining two ends of one nucleic acid, i.e., intramolecular joining, or two ends of two nucleic acids, i.e., intermolecular joining, using a ligase or another enzyme with ligase activity. The ligation splint holds the ends adjacent to each other and creates a ligation junction between the 5'-phosphorylated and a 3'-hydroxylated ends that are to be ligated.

[0459] In one instance, a splint-mediated ligation or splint ligation method may be used to synthesize the chimeric polynucleotides described herein. The chimeric polynucleotide may be assembled using a method that does not rely on the presence of restriction endonuclease cleavage sites such as the method described in International Patent Publication No. WO2012138453. Splint-mediated ligation allows for the rapid synthesis of the construct using controlled concatenation and without the need or with limited need for the introduction of restriction sites at the joining regions. As a non-limiting example, splint ligation may be used to add at least one untranslated region to a coding region of the chimeric polynucleotide. In one instance, splint ligation may be used in combination with other synthesis methods in the synthesis of the chimeric polynucleotides described herein.

[0460] If the 5'-phosphorylated and the 3'-hydroxyl ends of nucleic acids are ligated when the ends are annealed to a ligation splint so that the ends are adjacent, enzymes such as, but not limited to, T4 DNA ligase, AMPLIGASE® DNA Ligase (EPICENTRE® Technologies), *Tth* DNA ligase, *Tfl* DNA ligase, or *Tsc* DNA Ligase (Prokaria) can be used. Farugui IN U.S. Pat. No. 6,368,801 describes that T4 RNA ligase can efficiently ligate ends of DNA molecules that are adjacent to each other when hybridized to an RNA splint. Thus, T4 RNA ligase is a suitable ligase for joining DNA ends with

[0461] a ligation splint oligo comprising RNA or modified RNA. Examples of RNA splints include modified RNA containing 2'-fluorine-CTP (2'-F-dCTP) and 2'-fluorine-UTP (2'-F-dUTP) made using the DURASCRIBE® T7 Transcription Kit (EPICENTRE® Technologies) disclosed in US Pat. No. 8,137,911 and US Pat. Publication 2012/0156679 to Dahl et al. The modified RNA produced from DURASCRIBE®T7 Transcription kit is completely resistant to RNase A digestion. DNA splint and DNA ligase may be used to generate RNA-protein fusions disclosed in US Pat. No. 6,258,558 to Szostak et al..

[0462] For intramolecular ligation of linear ssDNA, US Pat. No et al teaches constructing a 83-nucleotide circle by making linear oligonucleoxynucleotides fragments on a DNA synthetiser followed by ligation with T4 DNA ligase and two 30 nucleotide splint oligonucleotides.

[0463] Circulation of linear sense promoter-containing cDNA is disclosed in US Pat. Publication No. 2012/0156679 to Dahl et al.. THERMOPHAGE™ ssDNA ligase (Prokazyme), which is derived from phage TS2126 that infects *Thermus scotoductus,* catalyzes ATP-dependent intra and inter-molecular ligation of DNA and RNA.

[0464] The solid-phase chemical synthesis method that uses phosphoramidite monomers is limited to produce DNA molecules with short strands. The purity of the DNA products and the yield of reactions become poor when the length exceeds 150 bases. For the synthesis of long polynucleotides in high yields, it is more convenient to use enzymatic ligation method in tandem with chemical synthesis. For example, Moore and Sharp describe preparing RNA fragments 10- to 20-nt long by chemical synthesis, to which site-specific modifications may be introduced, annealing the fragments to a cDNAsplint, and then assemble the fragments with T4 DNA ligase (Moore et al., Science, vol. 256, 992-997 (1992)). Ligation reactions of oligoribonucleotides with T4 RNA ligase and a DNA splint or a polyribonucleotide to generate large, synthetic RNAs are described in Bain et al., Nucleic Acids Research, vol. 20(16), 4372 (1992), Stark et al., RNA, vol. 12, 2014-2019 (2006), and US Pat. Application No. 2005/0130201 to Deras et al.. 5'-cap and 3'-polyA fail are often added by enzymatic addition to an oligonucleotide synthesized with solid-phase methods. As a non-limiting example, a synthetic capped 42-mer mRNA has been synthesized in three fragments enzymatically ligated as described by Iwase et al. (Nucleic Acids Research, vol. 20,1643-1648 (1992)). As another example, a 16.3-kilobase mouse mitochondrial genome has been produced from 600 overlapping 60-mer polynucleotides. The method cycles between *in vitro* recombination and amplification may be repeated until the desired length is reached (Gibson et al., Nature Methods, vol. 7, 901-903 (2010)). The assembly of a 1.08 megabase *Mycoplasma mycoides* JCVI-syn1.0 genome has also been reported. As a non-limiting example, 1080 bp cassettes are produced by assembling polynucleotide fragments chemically generated from a polynucleotide synthesizer. The genome is then assembled in three stages by transformation and homologous recombination in yeast (Gibson, et al., Science, vol. 329, 52-56 (2010)).

[0465] Studies have been conducted to join short DNA fragments with chemical linkers. 'Click' chemistry or 'click' ligation, the cycloaddition reaction between azide and alkyne, has gained a lot of interest because of its advantages such as mild reaction condition, high yields, and inoffensive byproducts. 'Click' chemistry is reviewed by Nwe et al. in Cancer Biotherapy and Radiopharmaceuticals, vol. 24(3), 289-302 (2009). DNA constructs up to 300 bases in length have been produced with click ligation and longer sequences are feasible. Demonstrated with PCR data, various DNA polymerases are able to amplify the synthesized DNA constructs made by click ligation despite the triazole linkers between the fragments resulting from the cycloaddition reaction *In vitro* transcription and rolling circle amplification can also be performed on the synthesized DNA constructs Hairpin ribozymes up to 100 nucleotides in length and cyclic mini-DNA duplexes have also been prepared with click ligation (El-Sagheer et al., Accounts of Chemical Research, vol. 45(8), 1258-1267 (2012)).

[0466] Sequential ligation can be performed on a solid substrate. For example, initial linker DNA molecules modified with biotin at the end are attached to streptavidin-coated beads. The 3'-ends of the linker DNA molecules are complimentary with the 5'-ends of the incoming DNA fragments. The beads are washed and collected after each ligation step and the final linear constructs are released by a meganuclease. This method allows rapid and efficient assembly of genes in an optimized order and orientation (Takita, DNA Research, vol. 20(4), 1-10 (2013)). Labeled polynucleotides synthesized on solid-supports are disclosed in US Pat. Pub. No. 2001/0014753 to Soloveichik et al. and US Pat. Pub. No. 2003/0191303 to Vinayak et al..

Quantification

[0467] In one instance, the polynucleotides of the present invention (i.e., antigen-encoding polynucleotides featured in the NAVs of the invention) may be quantified in exosomes or when derived from one or more bodily fluid. As used herein "bodily fluids" include peripheral blood, serum, plasma, ascites, urine, cerebrospinal fluid (CSF), sputum, saliva, bone marrow, synovial fluid, aqueous humor, amniotic fluid, cerumen, breast milk, broncheoalveolar lavage fluid, semen, prostatic fluid, cowper's fluid or pre-ejaculatory fluid, sweat, fecal matter, hair, tears, cyst fluid, pleural and peritoneal fluid, pericardial fluid, lymph, chyme, chyle, bile, interstitial fluid, menses, pus, sebum, vomit, vaginal secretions, mucosal secretion, stool water, pancreatic juice, lavage fluids from sinus cavities, bronchopulmonary aspirates, blastocyl cavity fluid, and umbilical cord blood. Alternatively, exosomes may be retrieved from an organ selected from the group consisting

of lung, heart, pancreas, stomach, intestine, bladder, kidney, ovary, testis, skin, colon, breast, prostate, brain, esophagus, liver, and placenta.

[0468] In the exosome quantification method, a sample of not more than 2mL is obtained from the subject and the exosomes isolated by size exclusion chromatography, density gradient centrifugation, differential centrifugation, nanomembrane ultrafiltration, immunoabsorbent capture, affinity purification, microfluidic separation, or combinations thereof. In the analysis, the level or concentration of a polynucleotide may be an expression level, presence, absence, truncation or alteration of the administered construct. It is advantageous to correlate the level with one or more clinical phenotypes or with an assay for a human disease biomarker. The assay may be performed using construct specific probes, cytometry, qRT-PCR, real-time PCR, PCR, flow cytometry, electrophoresis, mass spectrometry, or combinations thereof while the exosomes may be isolated using immunohistochemical methods such as enzyme linked immunosorbent assay (ELISA) methods. Exosomes may also be isolated by size exclusion chromatography, density gradient centrifugation, differential centrifugation, nanomembrane ultrafiltration, immunoabsorbent capture, affinity purification, microfluidic separation, or combinations thereof.

[0469] These methods afford the investigator the ability to monitor, in real time, the level of polynucleotides remaining or delivered. This is possible because the polynucleotides of the present invention differ from the endogenous forms due to the structural or chemical modifications.

[0470] In one instance, the polynucleotide may be quantified using methods such as, but not limited to, ultraviolet visible spectroscopy (UV/Vis). A non-limiting example of a UV/Vis spectrometer is a NANODROPO spectrometer (ThermoFisher, Waltham, MA). The quantified polynucleotide may be analyzed in order to determine if the polynucleotide may be of proper size, check that no degradation of the polynucleotide has occurred. Degradation of the polynucleotide may be checked by methods such as, but not limited to, agarose gel electrophoresis, HPLC based purification methods such as, but not limited to, strong anion exchange HPLC, weak anion exchange HPLC, reverse phase HPLC (RP-HPLC), and hydrophobic interaction HPLC (HIC-HPLC), liquid chromatography-mass spectrometry (LCMS), capillary electrophoresis (CE) and capillary gel electrophoresis (CGE).

Purification

[0471] Purification of the polynucleotides of the invention (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention) described herein may include, but is not limited to, polynucleotide clean-up, quality assurance and quality control. Clean-up may be performed by methods known in the arts such as, but not limited to, AGENCOURT® beads (Beckman Coulter Genomics, Danvers, MA), poly-T beads, LNA™ oligo-T capture probes (EXIQON® Inc, Vedbaek, Denmark) or HPLC based purification methods such as, but not limited to, strong anion exchange HPLC, weak anion exchange HPLC, reverse phase HPLC (RP-HPLC), and hydrophobic interaction HPLC (HIC-HPLC). The term "purified" when used in relation to a polynucleotide such as a "purified polynucleotide" refers to one that is separated from at least one contaminant. As used herein, a "contaminant" is any substance which makes another unfit, impure or inferior. Thus, a purified polynucleotide (e.g., DNA and RNA) is present in a form or setting different from that in which it is found in nature, or a form or setting different from that which existed prior to subjecting it to a treatment or purification method.

[0472] A quality assurance and/or quality control check may be conducted using methods such as, but not limited to, gel electrophoresis, UV absorbance, or analytical HPLC.

[0473] In another instance, the polynucleotides may be sequenced by methods including, but not limited to reverse-transcriptase-PCR.

IV. Modifications

[0474] In exemplary embodiments, polynucleotides of the invention (*i.e.*, antigen-encoding polynucleotides featured in the NAVs of the invention) can include various substitutions and/or insertions.

[0475] The data presented in the Examples demonstrate significant enhanced immune responses using the formulations of the invention. The data demonstrated the effectiveness of both chemically modified and unmodified RNA vaccines of the disclosure. Surprisingly, in contrast to prior art reports that it was preferable to use chemically unmodified mRNA formulated in a carrier for the production of vaccines, it was discovered herein that chemically modified mRNA-LNP vaccines required a much lower effective mRNA dose than unmodified mRNA, i.e., tenfold less than unmodified mRNA.

[0476] Additionally, a study described herein involved intravenous (IV), intramuscular (IM), or intradermal (ID) vaccination of mice, followed by challenge with a lethal dose of virus. In addition to all of the vaccinated animals surviving the lethal dose, significantly stronger early immune responses were observed (anti-viral activity via virus neutralization assay and HA inhibition (HAI)) in comparison to protein antigen and other lipid based formulations (lipoplex). The data demonstrated that as early as 1 week after vaccination both groups of animals receiving a chemically modified mRNA-LNP formulation (ID or IM) displayed HAI titers over 40, at 60 and 114, respectively. An HAI titer of greater than 40 is deemed sufficient to protect from a lethal challenge of influenza. The rapid response was unexpected, particularly when compared

to the response seen with protein antigen and mRNA vaccines formulated in other lipid carriers (lipoplex), which at one week and even at three weeks following vaccination continued to show ineffective HAI titers of less than 40.

[0477] At each of the later time points (3 weeks and 5 weeks), the formulations of the invention continued to provide significantly stronger therapeutic responses than did protein antigen. In fact even chemically unmodified mRNA-LNP formulation administered by IV route had enhanced HAI titers with respect to the protein antigen. By week 3, all of the animals receiving an mRNA-LNP formulation by IM or ID administration displayed HAI activity over 40, as compared to protein antigen, which at one week and three weeks continued to show HAI titers of less than 40. By week 5 the chemically modified mRNA-LNP formulation administered by ID route had a surprising HAI activity of greater than 10,000, in contrast to the HAI titer of around 400 for protein antigen at that time point.

[0478] Both the chemically modified and unmodified RNA vaccines produced better immune responses than did mRNA vaccines formulated in a different lipid carrier (lipoplex). At week 5 the non-chemically modified mRNA -lipoplex vaccine produced HAI titers of 197, in comparison to those achieved by the mRNA-LNP formulations of the invention (HAI titers of 635-10,152). At all other time points and for all of the chemically modified mRNA-lipoplex vaccines, none of the HAI titers reached the critical level of greater than 40. Additionally, the mRNA -lipoplex vaccine did not result in any detectable neutralizing activity in the microneutralization activity, even as late as five weeks aftervaccination.

[0479] As used herein in a polynucleotide (such as a chimeric polynucleotide, IVT polynucleotide or a circular polynucleotide), the terms "chemical modification" or, as appropriate, "chemically modified" refer to modification with respect to adenosine (A), guanosine (G), uridine (U), thymidine (T) or cytidine (C) ribo- or deoxyribnucleosides in one or more of their position, pattern, percent or population. Generally, herein, these terms are not intended to refer to the ribonucleotide modifications in naturally occurring 5'-terminal mRNA cap moieties.

[0480] In a polypeptide, the term "modification" refers to a modification as compared to the canonical set of 20 amino acids.

[0481] The modifications may be various distinct modifications In some embodiments, the regions may contain one, two, or more (optionally different) nucleoside or nucleotide modifications In some embodiments, a modified polynucleotide, introduced to a cell may exhibit reduced degradation in the cell, as compared to an unmodified polynucleotide.

[0482] The at least one chemical modification of the present invention is set out in the appended claims.

[0483] Modifications of the polynucleotides of the NAVs which are useful in the present invention include, but are not limted to those in Table 22. Noted in the table are the symbol of the modification, the nucleobase type and whether the modification is naturally occurring or not.

Table 22. Modifications

Name	Symbol	Base	Naturally Occurring
2-methylthio-N6-(cis-hydroxyisopentenyl)adenosine	ms2i6A	A	YES
2-methylthio-N6-methyladenosine	ms2m6A	A	YES
2-methylthio-N6-threonyl carbamoyladenosine	ms2t6A	A	YES
N6-glycinylcarbamoyladenosine	g6A	A	YES
N6-isopentenyladenosine	i6A	A	YES
N6-methyladenosine	m6A	A	YES
N6-threonylcarbamoyladenosine	t6A	A	YES
1,2'-O-dimethyladenosine	m1Am	A	YES

1-methyladenosine	m1A	A	YES
2'-O-methyladenosine	Am	A	YES
2'-O-ribosyladenosine (phosphate)	Ar(p)	A	YES
2-methyladenosine	m2A	A	YES
2-methylthio-N6 isopentenyladenosine	ms2i6A	A	YES
2-methylthio-N6-hydroxynorvalyl carbamoyladenosine	ms2hn6A	A	YES
2'-O-methyladenosine	m6A	A	YES
2'-O-ribosyladenosine (phosphate)	Ar(p)	A	YES
Isopentenyladenosine	Iga	A	YES
N6-(cis-hydroxyisopentenyl)adenosine	io6A	A	YES
N6,2'-O-dimethyladenosine	m6Am	A	YES
N^6,2'-O-dimethyladenosine	m^6Am	A	YES
N6,N6,2'-O-trimethyladenosine	m62Am	A	YES
N6,N6-dimethyladenosine	m62A	A	YES
N6-acetyladenosine	ac6A	A	YES
N6-hydroxynorvalylcarbamoyladenosine	hn6A	A	YES
N6-methyl-N6-threonylcarbamoyladenosine	m6t6A	A	YES
2-methyladenosine	m^2A	A	YES
2-methylthio-N^6-isopentenyladenosine	ms^2i^6A	A	YES
7-deaza-adenosine	--	A	NO
N1-methyl-adenosine	--	A	NO
N6, N6 (dimethyl)adenine	--	A	NO
N6-cis-hydroxy-isopentenyl-adenosine	--	A	NO
α-thio-adenosine	--	A	NO
2 (amino)adenine	--	A	NO
2 (aminopropyl)adenine	--	A	NO
2 (methylthio) N6 (isopentenyl)adenine	--	A	NO
2-(alkyl)adenine	--	A	NO
2-(aminoalkyl)adenine	--	A	NO
2-(aminopropyl)adenine	--	A	NO
2-(halo)adenine	--	A	NO
2-(halo)adenine	--	A	NO
2-(propyl)adenine	--	A	NO
2'-Amino-2'-deoxy-ATP	--	A	NO
2'-Azido-2'-deoxy-ATP	--	A	NO
2'-Deoxy-2'-a-aminoadenosine TP	--	A	NO
2'-Deoxy-2'-a-azidoadenosine TP	--	A	NO
6 (alkyl)adenine	--	A	NO
6 (methyl)adenine	--	A	NO
6-(alkyl)adenine	--	A	NO
6-(methyl)adenine	--	A	NO

7 (deaza)adenine	--	A	NO
8 (alkenyl)adenine	--	A	NO
8 (alkynyl)adenine	--	A	NO
8 (amino)adenine	--	A	NO
8 (thioalkyl)adenine	--	A	NO
8-(alkenyl)adenine	--	A	NO
8-(alkyl)adenine	--	A	NO
8-(alkynyl)adenine	--	A	NO
8-(amino)adenine	--	A	NO
8-(halo)adenine	--	A	NO
8-(hydroxyl)adenine	--	A	NO
8-(thioalkyl)adenine	--	A	NO
8-(thiol)adenine	--	A	NO
8-azido-adenosine	--	A	NO
aza adenine	--	A	NO
deaza adenine	--	A	NO
N6 (methyl)adenine	--	A	NO
N6-(isopentyl)adenine	--	A	NO
7-deaza-8-aza-adenosine	--	A	NO
7-methyladenine	--	A	NO
1-Deazaadenosine TP	--	A	NO
2'Fluoro-N6-Bz-deoxyadenosine TP	--	A	NO
2'-OMe-2-Amino-ATP	--	A	NO
2'O-methyl-N6-Bz-deoxyadenosine TP	--	A	NO
2'-a-Ethynyladenosine TP	--	A	NO
2-aminoadenine	--	A	NO
2-Aminoadenosine TP	--	A	NO
2-Amino-ATP	--	A	NO
2'-a-Trifluoromethyladenosine TP	--	A	NO
2-Azidoadenosine TP	--	A	NO
2'-b-Ethynyladenosine TP	--	A	NO
2-Bromoadenosine TP	--	A	NO
2'-b-Trifluoromethyladenosine TP	--	A	NO
2-Chloroadenosine TP	--	A	NO
2'-Deoxy-2',2'-difluoroadenosine TP	--	A	NO
2'-Deoxy-2'-a-mercaptoadenosine TP	--	A	NO
2'-Deoxy-2'-a-thiomethoxyadenosine TP	--	A	NO

2'-Deoxy-2'-b-aminoadenosine TP	--	A	NO
2'-Deoxy-2'-b-azidoadenosine TP	--	A	NO
2'-Deoxy-2'-b-bromoadenosine TP	--	A	NO
2'-Deoxy-2'-b-chloroadenosine TP	--	A	NO
2'-Deoxy-2'-b-fluoroadenosine TP	--	A	NO
2'-Deoxy-2'-b-iodoadenosine TP	--	A	NO
2'-Deoxy-2'-b-mercaptoadenosine TP	--	A	NO
2'-Deoxy-2'-b-thiomethoxyadenosine TP	--	A	NO
2-Fluoroadenosine TP	--	A	NO
2-Iodoadenosine TP	--	A	NO
2-Mercaptoadenosine TP	--	A	NO
2-methoxy-adenine	--	A	NO
2-methylthio-adenine	--	A	NO
2-Trifluoromethyladenosine TP	--	A	NO
3-Deaza-3-bromoadenosine TP	--	A	NO
3-Deaza-3-chloroadenosine TP	--	A	NO
3-Deaza-3-fluoroadenosine TP	--	A	NO
3-Deaza-3-iodoadenosine TP	--	A	NO
3-Deazaadenosine TP	--	A	NO
4'-Azidoadenosine TP	--	A	NO
4'-Carbocyclic adenosine TP	--	A	NO
4'-Ethynyladenosine TP	--	A	NO
5'-Homo-adenosine TP	--	A	NO
8-Aza-ATP	--	A	NO
8-bromo-adenosine TP	--	A	NO
8-Trifluoromethyladenosine TP	--	A	NO
9-Deazaadenosine TP	--	A	NO
2-aminopurine	--	A/G	NO
7-deaza-2,6-diaminopurine	--	A/G	NO
7-deaza-8-aza-2,6-diaminopurine	--	A/G	NO
7-deaza-8-aza-2-aminopurine	--	A/G	NO
2,6-diaminopurine	--	A/G	NO
7-deaza-8-aza-adenine, 7-deaza-2-aminopurine	--	A/G	NO
2-thiocytidine	s2C	C	YES
3-methylcytidine	m3C	C	YES
5-formylcytidine	f5C	C	YES
5-hydroxymethylcytidine	hm5C	C	YES
5-methylcytidine	m5C	C	YES
N4-acetylcytidine	ac4C	C	YES

2'-O-methylcytidine	Cm	C	YES
2'-O-methylcytidine	Cm	C	YES
5,2'-O-dimethylcytidine	m5 Cm	C	YES
5-formyl-2'-O-methylcytidine	f5Cm	C	YES
Lysidine	k2C	C	YES
N4,2'-O-dimethylcytidine	m4Cm	C	YES
N4-acetyl-2'-O-methylcytidine	ac4Cm	C	YES
N4-methylcytidine	m4C	C	YES
N4,N4-Dimethyl-2'-OMe-Cytidine TP	--	C	YES
4-methylcytidine	--	C	NO
5-aza-cytidine	--	C	NO
Pseudo-iso-cytidine	--	C	NO
pyrrolo-cytidine	--	C	NO
α-thio-cytidine	--	C	NO
2-(thio)cytosine	--	C	NO
2'-Amino-2'-deoxy-CTP	--	C	NO
2'-Azido-2'-deoxy-CTP	--	C	NO
2'-Deoxy-2'-a-aminocytidine TP	--	C	NO
2'-Deoxy-2'-a-azidocytidine TP	--	C	NO
3 (deaza) 5 (aza)cytosine	--	C	NO
3 (methyl)cytosine	--	C	NO
3-(alkyl)cytosine	--	C	NO
3-(deaza) 5 (aza)cytosine	--	C	NO
3-(methyl)cytidine	--	C	NO
4,2'-O-dimethylcytidine	--	C	NO
5 (halo)cytosine	--	C	NO
5 (methyl)cytosine	--	C	NO
5 (propynyl)cytosine	--	C	NO
5 (trifluoromethyl)cytosine	--	C	NO
5-(alkyl)cytosine	--	C	NO
5-(alkynyl)cytosine	--	C	NO
5-(halo)cytosine	--	C	NO
5-(propynyl)cytosine	--	C	NO
5-(trifluoromethyl)cytosine	--	C	NO
5-bromo-cytidine	--	C	NO
5-iodo-cytidine	--	C	NO
5-propynyl cytosine	--	C	NO
6-(azo)cytosine	--	C	NO
6-aza-cytidine	--	C	NO

aza cytosine	--	C	NO
deaza cytosine	--	C	NO
N4 (acetyl)cytosine	--	C	NO
1-methyl-1-deaza-pseudoisocytidine	--	C	NO
1-methyl-pseudoisocytidine	--	C	NO
2-methoxy-5-methyl-cytidine	--	C	NO
2-methoxy-cytidine	--	C	NO
2-thio-5-methyl-cytidine	--	C	NO
4-methoxy-1-methyl-pseudoisocytidine	--	C	NO
4-methoxy-pseudoisocytidine	--	C	NO
4-thio-1-methyl-1-deaza-pseudoisocytidine	--	C	NO
4-thio-1-methyl-pseudoisocytidine	--	C	NO
4-thio-pseudoisocytidine	--	C	NO
5-aza-zebularine	--	C	NO
5-methyl-zebularine	--	C	NO
pyrrolo-pseudoisocytidine	--	C	NO
Zebularine	--	C	NO
(E)-5-(2-Bromo-vinyl)cytidine TP	--	C	NO
2,2'-anhydro-cytidine TP hydrochloride	--	C	NO
2'Fluor-N4-Bz-cytidine TP	--	C	NO
2'Fluoro-N4-Acetyl-cytidine TP	--	C	NO
2'-O-Methyl-N4-Acetyl-cytidine TP	--	C	NO
2'O-methyl-N4-Bz-cytidine TP	--	C	NO
2'-a-Ethynylcytidine TP	--	C	NO
2'-a-Trifluoromethylcytidine TP	--	C	NO
2'-b-Ethynylcytidine TP	--	C	NO
2'-b-Trifluoromethylcytidine TP	--	C	NO
2'-Deoxy-2',2'-difluorocytidine TP	--	C	NO
2'-Deoxy-2'-a-mercaptocytidine TP	--	C	NO
2'-Deoxy-2'-a-thiomethoxycytidine TP	--	C	NO
2'-Deoxy-2'-b-aminocytidine TP	--	C	NO
2'-Deoxy-2'-b-azidocytidine TP	--	C	NO
2'-Deoxy-2'-b-bromocytidine TP	--	C	NO
2'-Deoxy-2'-b-chlorocytidine TP	--	C	NO
2'-Deoxy-2'-b-fluorocytidine TP	--	C	NO
2'-Deoxy-2'-b-iodocytidine TP	--	C	NO
2'-Deoxy-2'-b-mercaptocytidine TP	--	C	NO

2'-Deoxy-2'-b-thiomethoxycytidine TP	--	C	NO
2'-O-Methyl-5-(1-propynyl)cytidine TP	--	C	NO
3'-Ethynylcytidine TP	--	C	NO
4'-Azidocytidine TP	--	C	NO
4'-Carbocyclic cytidine TP	--	C	NO
4'-Ethynylcytidine TP	--	C	NO
5-(1-Propynyl)ara-cytidine TP	--	C	NO
5-(2-Chloro-phenyl)-2-thiocytidine TP	--	C	NO
5-(4-Amino-phenyl)-2-thiocytidine TP	--	C	NO
5-Aminoallyl-CTP	--	C	NO
5-Cyanocytidine TP	--	C	NO
5-Ethynylara-cytidine TP	--	C	NO
5-Ethynylcytidine TP	--	C	NO
5'-Homo-cytidine TP	--	C	NO
5-Methoxycytidine TP	--	C	NO
5-Trifluoromethyl-Cytidine TP	--	C	NO
N4-Amino-cytidine TP	--	C	NO
N4-Benzoyl-cytidine TP	--	C	NO
Pseudoisocytidine	--	C	NO
7-methylguancsine	m7G	G	YES
N2,2'-O-dimethylguanosine	m2Gm	G	YES
N2-methylguancsine	m2G	G	YES
Wyosine	imG	G	YES
1,2'-O-dimethylguanosine	m1Gm	G	YES
1-methylguancsine	m1G	G	YES
2'-O-methylguanosine	Gm	G	YES
2'-O-ribosylguanosine (phosphate)	Gr(p)	G	YES
2'-O-methylguanosine	Gm	G	YES
2'-O-ribosylguanosine (phosphate)	Gr(p)	G	YES
7-aminomethyl-7-deazaguanosine	preQ1	G	YES
7-cyano-7-deazaguanosine	preQ0	G	YES
Archaeosine	G+	G	YES
Methylwyosine	mimG	G	YES
N2,7-dimethylguanosine	m2,7G	G	YES
N2,N2,2'-O-trimethylguanosine	m22Gm	G	YES
N2,N2,7-trimethylguanosine	m2,2,7G	G	YES
N2,N2-dimethylguanosine	m22G	G	YES
N^2,7,2'-O-trimethylguanosine	m^{2,7}Gm	G	YES
6-thio-guanosine	--	G	NO
7-deaza-guanosine	--	G	NO
8-oxo-guanosine	--	G	NO

N1-methyl-guanosine	--	G	NO
α-thio-guanosine	--	G	NO
2 (propyl)guanine	--	G	NO
2-(alkyl)guanine	--	G	NO
2'-Amino-2'-deoxy-GTP	--	G	NO
2'-Azido-2'-deoxy-GTP	--	G	NO
2'-Deoxy-2'-a-aminoguanosine TP	--	G	NO
2'-Deoxy-2'-a-azidoguanosine TP	--	G	NO
6 (methyl)guanine	--	G	NO
6-(alkyl)guanine	--	G	NO
6-(methyl)guanine	--	G	NO
6-methyl-guanosine	--	G	NO
7 (alkyl)guanine	--	G	NO
7 (deaza)guanine	--	G	NO
7 (methyl)guanine	--	G	NO
7-(alkyl)guanine	--	G	NO
7-(deaza)guanine	--	G	NO
7-(methyl)guanine	--	G	NO
8 (alkyl)guanine	--	G	NO
8 (alkynyl)guanine	--	G	NO
8 (halo)guanine	--	G	NO
8 (thioalkyl)guanine	--	G	NO
8-(alkenyl)guanine	--	G	NO
8-(alkyl)guanine	--	G	NO
8-(alkynyl)guanine	--	G	NO
8-(amino)guanine	--	G	NO
8-(halo)guanine	--	G	NO
8-(hydroxyl)guanine	--	G	NO
8-(thioalkyl)guanine	--	G	NO
8-(thiol)guanine	--	G	NO
aza guanine	--	G	NO
deaza guanine	--	G	NO
N (methyl)guanine	--	G	NO
N-(methyl)guanine	--	G	NO
1-methyl-6-thio-guanosine	--	G	NO
6-methoxy-guanosine	--	G	NO
6-thio-7-deaza-8-aza-guanosine	--	G	NO

6-thio-7-deaza-guanosine	--	G	NO
6-thio-7-methyl-guanosine	--	G	NO
7-deaza-8-aza-guanosine	--	G	NO
7-methyl-8-oxo-guanosine	--	G	NO
N2,N2-dimethyl-6-thio-guanosine	--	G	NO
N2-methyl-6-thio-guanosine	--	G	NO
1-Me-GTP	--	G	NO
2'Fluoro-N2-isobutyl-guanosine TP	--	G	NO
2'O-methyl-N2-isobutyl-guanosine TP	--	G	NO
2'-a-Ethynylguanosine TP	--	G	NO
2'-a-Trifluoromethylguanosine TP	--	G	NO
2'-b-Ethynylguanosine TP	--	G	NO
2'-b-Trifluoromethylguanosine TP	--	G	NO
2'-Deoxy-2',2'-difluoroguanosine TP	--	G	NO
2'-Deoxy-2'-a-mercaptoguanosine TP	--	G	NO
2'-Deoxy-2'-a-thiomethoxyguanosine TP	--	G	NO
2'-Deoxy-2'-b-aminoguanosine TP	--	G	NO
2'-Deoxy-2'-b-azidoguanosine TP	--	G	NO
2'-Deoxy-2'-b-bromoguanosine TP	--	G	NO
2'-Deoxy-2'-b-chloroguanosine TP	--	G	NO
2'-Deoxy-2'-b-fluoroguanosine TP	--	G	NO
2'-Deoxy-2'-b-iodoguanosine TP	--	G	NO
2'-Deoxy-2'-b-mercaptoguanosine TP	--	G	NO
2'-Deoxy-2'-b-thiomethoxyguanosine TP	--	G	NO
4'-Azidoguanosine TP	--	G	NO
4'-Carbocyclic guanosine TP	--	G	NO
4'-Ethynylguanosine TP	--	G	NO
5'-Homo-guanosine TP	--	G	NO
8-bromo-guanosine TP	--	G	NO
9-Deazaguanosine TP	--	G	NO
N2-isobutyl-guanosine TP	--	G	NO
1-methylinosine	m1I	I	YES
Inosine	I	I	YES
1,2'-O-dimethylinosine	m1Im	I	YES
2'-O-methylinosine	Im	I	YES
7-methylinosine		I	NO
2'-O-methylinosine	Im	I	YES
Epoxyqueuosine	oQ	Q	YES
galactosyl-queuosine	galQ	Q	YES

Mannosylqueuosine	manQ	Q	YES
Queuosine	Q	Q	YES
allyamino-thymidine	--	T	NO
aza thymidine	--	T	NO
deaza thymidine	--	T	NO
deoxy-thymidine	--	T	NO
2'-O-methyluridine	--	U	YES
2-thiouridine	s2U	U	YES
3-methyluridine	m3U	U	YES
5-carboxymethyluridine	cm5U	U	YES
5-hydroxyuridine	ho5U	U	YES
5-methyluridine	m5U	U	YES
5-taurinomethyl-2-thiouridine	τm5s2U	U	YES
5-taurinomethyluridine	τm5U	U	YES
Dihydrouridine	D	U	YES
Pseudouridine	Ψ	U	YES
(3-(3-amino-3-carboxypropyl)uridine	acp3U	U	YES
1-methyl-3-(3-amino-5-carboxypropyl)pseudouridine	m1acp3Ψ	U	YES
1-methylpseduouridine	m1Ψ	U	YES
1-methyl-pseudouridine	--	U	YES
2'-O-methyluridine	Um	U	YES
2'-O-methylpseudouridine	Ψm	U	YES
2'-O-methyluridine	Um	U	YES
2-thio-2'-O-methyluridine	s2Um	U	YES
3-(3-amino-3-carboxypropyl)uridine	acp3U	U	YES
3,2'-O-dimethyluridine	m3Um	U	YES
3-Methyl-pseudo-Uridine TP	--	U	YES
4-thiouridine	s4U	U	YES
5-(carboxyhydroxymethyl)uridine	chm5U	U	YES
5-(carboxyhydroxymethyl)uridine methyl ester	mchm5U	U	YES
5,2'-O-dimethyluridine	m5Um	U	YES
5,6-dihydro-uridine	--	U	YES
5-aminomethyl-2-thiouridine	nm5s2U	U	YES
5-carbamoylmethyl-2'-O-methyluridine	ncm5Um	U	YES
5-carbamoylmethyluridine	ncm5U	U	YES
5-carboxyhydroxymethyluridine	--	U	YES
5-carboxyhydroxymethyluridine methyl ester	--	U	YES
5-carboxymethylaminomethyl-2'-O-methyluridine	cmnm5Um	U	YES
5-carboxymethylaminomethyl-2-thiouridine	cmnm5s2U	U	YES
5-carboxymethylaminomethyl-2-thiouridine	--	U	YES
5-carboxymethylaminomethyluridine	cmnm5U	U	YES
5-carboxymethylaminomethyluridine	--	U	YES
5-Carbamoylmethyluridine TP	--	U	YES
5-methoxycarbonylmethyl-2'-O-methyluridine	mcm5Um	U	YES
5-methoxycarbonylmethyl-2-thiouridine	mcm5s2U	U	YES

5-methoxycarbonylmethyluridine	mcm5U	U	YES
5-methoxyuridine	mo5U	U	YES
5-methyl-2-thiouridine	m5s2U	U	YES
5-methylaminomethyl-2-selenouridine	mnm5se2U	U	YES
5-methylaminomethyl-2-thiouridine	mnm5s2U	U	YES
5-methylaminomethyluridine	mnm5U	U	YES
5-Methyldihydrouridine	--	U	YES
5-Oxyacetic acid- Uridine TP	--	U	YES
5-Oxyacetic acid-methyl ester-Uridine TP	--	U	YES
N1-methyl-pseudo-uridine	--	U	YES
uridine 5-oxyacetic acid	cmo5U	U	YES
uridine 5-oxyacetic acid methyl ester	mcmo5U	U	YES
3-(3-Amino-3-carboxypropyl)-Uridine TP	--	U	YES
5-(iso-Pentenylaminomethyl)- 2-thiouridine TP	--	U	YES
5-(iso-Pentenylaminomethyl)-2'-O-methyluridine TP	--	U	YES
5-(iso-Pentenylaminomethyl)uridine TP	--	U	YES
5-propynyl uracil	--	U	NO
α-thio-uridine	--	U	NO
1 (aminoalkylamino-carbonylethylenyl)-2(thio)-pseudouracil	--	U	NO
1 (aminoalkylaminocarbonylethylenyl)-2,4-(dithio)pseudouracil	--	U	NO
1 (aminoalkylaminocarbonylethylenyl)-4 (thio)pseudouracil	--	U	NO
1 (aminoalkylaminocarbonylethylenyl)-pseudouracil	--	U	NO
1 (aminocarbonylethylenyl)-2(thio)-pseudouracil	--	U	NO
1 (aminocarbonylethylenyl)-2,4-(dithio)pseudouracil	--	U	NO
1 (aminocarbonylethylenyl)-4 (thio)pseudouracil	--	U	NO
1 (aminocarbonylethylenyl)-pseudouracil	--	U	NO
1 substituted 2(thio)-pseudouracil	--	U	NO
1 substituted 2,4-(dithio)pseudouracil	--	U	NO
1 substituted 4 (thio)pseudouracil	--	U	NO
1 substituted pseudouracil	--	U	NO
1-(aminoalkylamino-carbonylethylenyl)-2-(thio)-pseudouracil	--	U	NO
1-Methyl-3-(3-amino-3-carboxypropyl) pseudouridine TP	--	U	NO
1-Methyl-3-(3-amino-3-carboxypropyl)pseudo-UTP	--	U	NO
1-Methyl-pseudo-UTP	--	U	NO
2 (thio)pseudouracil	--	U	NO
2' deoxy uridine	--	U	NO
2' fluorouridine	--	U	NO
2-(thio)uracil	--	U	NO
2,4-(dithio)psuedouracil	--	U	NO

2' methyl, 2'amino, 2'azido, 2'fluro-guanosine	--	U	NO
2'-Amino-2'-deoxy-UTP	--	U	NO
2'-Azido-2'-deoxy-UTP	--	U	NO
2'-Azido-deoxyuridine TP	--	U	NO
2'-O-methylpseudouridine	--	U	NO
2' deoxy uridine	2' dU	U	NO
2' fluorouridine	--	U	NO
2'-Deoxy-2'-a-aminouridine TP	--	U	NO
2'-Deoxy-2'-a-azidouridine TP	--	U	NO
2-methylpseudouridine	m3Ψ	U	NO
3 (3 amino-3 carboxypropyl)uracil	--	U	NO
4 (thio)pseudouracil	--	U	NO
4-(thio)pseudouracil	--	U	NO
4-(thio)uracil	--	U	NO
4-thiouracil	--	U	NO
5 (1,3-diazole-1-alkyl)uracil	--	U	NO
5 (2-aminopropyl)uracil	--	U	NO
5 (aminoalkyl)uracil	--	U	NO
5 (dimethylaminoalkyl)uracil	--	U	NO
5 (guanidiniumalkyl)uracil	--	U	NO
5 (methoxycarbonylmethyl)-2-(thio)uracil	--	U	NO
5 (methoxycarbonyl-methyl)uracil	--	U	NO
5 (methyl) 2 (thio)uracil	--	U	NO
5 (methyl) 2,4 (dithio)uracil	--	U	NO
5 (methyl) 4 (thio)uracil	--	U	NO
5 (methylaminomethyl)-2 (thio)uracil	--	U	NO
5 (methylaminomethyl)-2,4 (dithio)uracil	--	U	NO
5 (methylaminomethyl)-4 (thio)uracil	--	U	NO
5 (propynyl)uracil	--	U	NO
5 (trifluoromethyl)uracil	--	U	NO
5-(2-aminopropyl)uracil	--	U	NO
5-(alkyl)-2-(thio)pseudouracil	--	U	NO
5-(alkyl)-2,4 (dithio)pseudouracil	--	U	NO
5-(alkyl)-4 (thio)pseudouracil	--	U	NO
5-(alkyl)pseudouracil	--	U	NO
5-(alkyl)uracil	--	U	NO
5-(alkynyl)uracil	--	U	NO

5-(allylamino)uracil	--	U	NO
5-(cyanoalkyl)uracil	--	U	NO
5-(dialkylaminoalkyl)uracil	--	U	NO
5-(dimethylaminoalkyl)uracil	--	U	NO
5-(guanidiniumalkyl)uracil	--	U	NO
5-(halo)uracil	--	U	NO
5-(l,3-diazole-l-alkyl)uracil	--	U	NO
5-(methoxy)uracil	--	U	NO
5-(methoxycarbonylmethyl)-2-(thio)uracil	--	U	NO
5-(methoxycarbonyl-methyl)uracil	--	U	NO
5-(methyl) 2(thio)uracil	--	U	NO
5-(methyl) 2,4 (dithio)uracil	--	U	NO
5-(methyl) 4 (thio)uracil	--	U	NO
5-(methyl)-2-(thio)pseudouracil	--	U	NO
5-(methyl)-2,4 (dithio)pseudouracil	--	U	NO
5-(methyl)-4 (thio)pseudouracil	--	U	NO
5-(methyl)pseudouracil	--	U	NO
5-(methylaminomethyl)-2 (thio)uracil	--	U	NO
5-(methylaminomethyl)-2,4(dithio)uracil	--	U	NO
5-(methylaminomethyl)-4-(thio)uracil	--	U	NO
5-(propynyl)uracil	--	U	NO
5-(trifluoromethyl)uracil	--	U	NO
5-aminoallyl-uridine	--	U	NO
5-bromo-uridine	--	U	NO
5-iodo-uridine	--	U	NO
5-uracil	--	U	NO
6 (azo)uracil	--	U	NO
6-(azo)uracil	--	U	NO
6-aza-uridine	--	U	NO
allyamino-uracil	--	U	NO
aza uracil	--	U	NO
deaza uracil	--	U	NO
N3 (methyl)uracil	--	U	NO
P seudo-UTP-1-2-ethanoic acid	--	U	NO
Pseudouracil	--	U	NO
4-Thio-pseudo-UTP	--	U	NO
1-carboxymethyl-pseudouridine	--	U	NO

1-methyl-1-deaza-pseudouridine	--	U	NO
1-propynyl-uridine	--	U	NO
1-taurinomethyl-1-methyl-uridine	--	U	NO
1-taurinomethyl-4-thio-uridine	--	U	NO
1-taurinomethyl-pseudouridine	--	U	NO
2-methoxy-4-thio-pseudouridine	--	U	NO
2-thio-1-methyl-1-deaza-pseudouridine	--	U	NO
2-thio-1-methyl-pseudouridine	--	U	NO
2-thio-5-aza-uridine	--	U	NO
2-thio-dihydropseudouridine	--	U	NO
2-thio-dihydrouridine	--	U	NO
2-thio-pseudouridine	--	U	NO
4-methoxy-2-thio-pseudouridine	--	U	NO
4-methoxy-pseudouridine	--	U	NO
4-thio-1-methyl-pseudouridine	--	U	NO
4-thio-pseudouridine	--	U	NO
5-aza-uridine	--	U	NO
Dihydropseudouridine	--	U	NO
(±)1-(2-Hydroxypropyl)pseudouridine TP	--	U	NO
(2R)-1-(2-Hydroxypropyl)pseudouridine TP	--	U	NO
(2S)-1-(2-Hydroxypropyl)pseudouridine TP	--	U	NO
(E)-5-(2-Bromo-vinyl)ara-uridine TP	--	U	NO
(E)-5-(2-Bromo-vinyl)uridine TP	--	U	NO
(Z)-5-(2-Bromo-vinyl)ara-uridine TP	--	U	NO
(Z)-5-(2-Bromo-vinyl)uridine TP	--	U	NO
1-(2,2,2-Trifluoroethyl)-pseudo-UTP	--	U	NO
1-(2,2,3,3,3-Pentafluoropropyl)pseudouridine TP	--	U	NO
1-(2,2-Diethoxyethyl)pseudouridine TP	--	U	NO
1-(2,4,6-Trimethylbenzyl)pseudouridine TP	--	U	NO
1-(2,4,6-Trimethyl-benzyl)pseudo-UTP	--	U	NO
1-(2,4,6-Trimethyl-phenyl)pseudo-UTP	--	U	NO
1-(2-Amino-2-carboxyethyl)pseudo-UTP	--	U	NO
1-(2-Amino-ethyl)pseudo-UTP	--	U	NO
1-(2-Hydroxyethyl)pseudouridine TP	--	U	NO
1-(2-Methoxyethyl)pseudouridine TP	--	U	NO
1-(3,4-Bis-trifluoromethoxybenzyl)pseudouridine TP	--	U	NO
1-(3,4-Dimethoxybenzyl)pseudouridine TP	--	U	NO

1-(3-Amino-3-carboxypropyl)pseudo-UTP	--	U	NO
1-(3-Amino-propyl)pseudo-UTP	--	U	NO
1-(3-Cyclopropyl-prop-2-ynyl)pseudouridine TP	--	U	NO
1-(4-Amino-4-carboxybutyl)pseudo-UTP	--	U	NO
1-(4-Amino-benzyl)pseudo-UTP	--	U	NO
1-(4-Amino-butyl)pseudo-UTP	--	U	NO
1-(4-Amino-phenyl)pseudo-UTP	--	U	NO
1-(4-Azidobenzyl)pseudouridine TP	--	U	NO
1-(4-Bromobenzyl)pseudouridine TP	--	U	NO
1-(4-Chlorobenzyl)pseudouridine TP	--	U	NO
1-(4-Fluorobenzyl)pseudouridine TP	--	U	NO
1-(4-Iodobenzyl)pseudouridine TP	--	U	NO
1-(4-Methanesulfonylbenzyl)pseudouridine TP	--	U	NO
1-(4-Methoxybenzyl)pseudouridine TP	--	U	NO
1-(4-Methoxy-benzyl)pseudo-UTP	--	U	NO
1-(4-Methoxy-phenyl)pseudo-UTP	--	U	NO
1-(4-Methylbenzyl)pseudouridine TP	--	U	NO
1-(4-Methyl-benzyl)pseudo-UTP	--	U	NO
1-(4-Nitrobenzyl)pseudouridine TP	--	U	NO
1-(4-Nitro-benzyl)pseudo-UTP	--	U	NO
1(4-Nitro-phenyl)pseudo-UTP	--	U	NO
1-(4-Thiomethoxybenzyl)pseudouridine TP	--	U	NO
1-(4-Trifluoromethoxybenzyl)pseudouridine TP	--	U	NO
1-(4-Trifluoromethylbenzyl)pseudouridine TP	--	U	NO
1-(5-Amino-pentyl)pseudo-UTP	--	U	NO
1-(6-Amino-hexyl)pseudo-UTP	--	U	NO
1,6-Dimethyl-pseudo-UTP	--	U	NO
1-[3-(2-{2-[2-(2-Aminoethoxy)-ethoxy]-ethoxy}-ethoxy)-propionyl]pseudouridine TP	--	U	NO
1-{3-[2-(2-Aminoethoxy)-ethoxy]-propionyl } pseudouridine TP	--	U	NO
1-Acetylpseudouridine TP	--	U	NO
1-Alkyl-6-(1-propynyl)-pseudo-UTP	--	U	NO
1-Alkyl-6-(2-propynyl)-pseudo-UTP	--	U	NO
1-Alkyl-6-allyl-pseudo-UTP	--	U	NO
1-Alkyl-6-ethynyl-pseudo-UTP	--	U	NO
1-Alkyl-6-homoallyl-pseudo-UTP	--	U	NO
1-Alkyl-6-vinyl-pseudo-UTP	--	U	NO
1-Allylpseudouridine TP	--	U	NO

1-Aminomethyl-pseudo-UTP	--	U	NO
1-Benzoylpseudouridine TP	--	U	NO
1-Benzyloxymethylpseudouridine TP	--	U	NO
1-Benzyl-pseudo-UTP	--	U	NO
1-Biotinyl-PEG2-pseudouridine TP	--	U	NO
1-Biotinylpseudouridine TP	--	U	NO
1-Butyl-pseudo-UTP	--	U	NO
1-Cyanomethylpseudouridine TP	--	U	NO
1-Cyclobutylmethyl-pseudo-UTP	--	U	NO
1-Cyclobutyl-pseudo-UTP	--	U	NO
1-Cycloheptylmethyl-pseudo-UTP	--	U	NO
1-Cycloheptyl-pseudo-UTP	--	U	NO
1-Cyclohexylmethyl-pseudo-UTP	--	U	NO
1-Cyclohexyl-pseudo-UTP	--	U	NO
1-Cyclooctylmethyl-pseudo-UTP	--	U	NO
1-Cyclooctyl-pseudo-UTP	--	U	NO
1-Cyclopentylmethyl-pseudo-UTP	--	U	NO
1-Cyclopentyl-pseudo-UTP	--	U	NO
1-Cyclopropylmethyl-pseudo-UTP	--	U	NO
1-Cyclopropyl-pseudo-UTP	--	U	NO
1-Ethyl-pseudo-UTP	--	U	NO
1-Hexyl-pseudo-UTP	--	U	NO
1-Homoallylpseudouridine TP	--	U	NO
1-Hydroxymethylpseudouridine TP	--	U	NO
1-iso-propyl-pseudo-UTP	--	U	NO
1-Me-2-thio-pseudo-UTP	--	U	NO
1-Me-4-thio-pseudo-UTP	--	U	NO
1-Me-alpha-thio-pseudo-UTP	--	U	NO
1-Methanesulfonylmethylpseudouridine TP	--	U	NO
1-Methoxymethylpseudouridine TP	--	U	NO
1-Methyl-6-(2,2,2-Trifluoroethyl)pseudo-UTP	--	U	NO
1-Methyl-6-(4-morpholino)-pseudo-UTP	--	U	NO
1-Methyl-6-(4-thiomorpholino)-pseudo-UTP	--	U	NO
1-Methyl-6-(substituted phenyl)pseudo-UTP	--	U	NO
1-Methyl-6-amino-pseudo-UTP	--	U	NO
1-Methyl-6-azido-pseudo-UTP	--	U	NO
1-Methyl-6-bromo-pseudo-UTP	--	U	NO

1-Methyl-6-butyl-pseudo-UTP	--	U	NO
1-Methyl-6-chloro-pseudo-UTP	--	U	NO
1-Methyl-6-cyano-pseudo-UTP	--	U	NO
1-Methyl-6-dimethylamino-pseudo-UTP	--	U	NO
1-Methyl-6-ethoxy-pseudo-UTP	--	U	NO
1-Methyl-6-ethylcarboxylate-pseudo-UTP	--	U	NO
1-Methyl-6-ethyl-pseudo-UTP	--	U	NO
1-Methyl-6-fluoro-pseudo-UTP	--	U	NO
1-Methyl-6-formyl-pseudo-UTP	--	U	NO
1-Methyl-6-hydroxyamino-pseudo-UTP	--	U	NO
1-Methyl-6-hydroxy-pseudo-UTP	--	U	NO
1-Methyl-6-iodo-pseudo-UTP	--	U	NO
1-Methyl-6-iso-propyl-pseudo-UTP	--	U	NO
1-Methyl-6-methoxy-pseudo-UTP	--	U	NO
1-Methyl-6-methylamino-pseudo-UTP	--	U	NO
1-Methyl-6-phenyl-pseudo-UTP	--	U	NO
1-Methyl-6-propyl-pseudo-UTP	--	U	NO
1-Methyl-6-tert-butyl-pseudo-UTP	--	U	NO
1-Methyl-6-trifluoromethoxy-pseudo-UTP	--	U	NO
1-Methyl-6-trifluoromethyl-pseudo-UTP	--	U	NO
1-Morpholinomethylpseudouridine TP	--	U	NO
1-Pentyl-pseudo-UTP	--	U	NO
1-Phenyl-pseudo-UTP	--	U	NO
1-Pivaloylpseudouridine TP	--	U	NO
1-Propargylpseudouridine TP	--	U	NO
1-Propyl-pseudo-UTP	--	U	NO
1-propynyl-pseudouridine	--	U	NO
1-p-tolyl-pseudo-UTP	--	U	NO
1-tert-Butyl-pseudo-UTP	--	U	NO
1-Thiomethoxymethylpseudouridine TP	--	U	NO
1-Thiomorpholinomethylpseudouridine TP	--	U	NO
1-Trifluoroacetylpseudouridine TP	--	U	NO
1-Trifluoromethyl-pseudo-UTP	--	U	NO
1-Vinylpseudouridine TP	--	U	NO
2,2'-anhydro-uridine TP	--	U	NO
2'-bromo-deoxyuridine TP	--	U	NO
2'-F-5-Methyl-2'-deoxy-UTP	--	U	NO

2'-OMe-5-Me-UTP	--	U	NO
2'-OMe-pseudo-UTP	--	U	NO
2'-a-Ethynyluridine TP	--	U	NO
2'-a-Trifluoromethyluridine TP	--	U	NO
2'-b-Ethynyluridine TP	--	U	NO
2'-b-Trifluoromethyluridine TP	--	U	NO
2'-Deoxy-2',2'-difluorouridine TP	--	U	NO
2'-Deoxy-2'-a-mercaptouridine TP	--	U	NO
2'-Deoxy-2'-a-thiomethoxyuridine TP	--	U	NO
2'-Deoxy-2'-b-aminouridine TP	--	U	NO
2'-Deoxy-2'-b-azidouridine TP	--	U	NO
2'-Deoxy-2'-b-bromouridine TP	--	U	NO
2'-Deoxy-2'-b-chlorouridine TP	--	U	NO
2'-Deoxy-2'-b-fluorouridine TP	--	U	NO
2'-Deoxy-2'-b-iodouridine TP	--	U	NO
2'-Deoxy-2'-b-mercaptouridine TP	--	U	NO
2'-Deoxy-2'-b-thiomethoxyuridine TP	--	U	NO
2-methoxy-4-thio-uridine	--	U	NO
2-methoxyuridine	--	U	NO
2'-O-Methyl-5-(1-propynyl)uridine TP	--	U	NO
3-Alkyl-pseudo-UTP	--	U	NO
4'-Azidouridine TP	--	U	NO
4'-Carbocyclic uridine TP	--	U	NO
4'-Ethynyluridine TP	--	U	NO
5-(1-Propynyl)ara-uridine TP	--	U	NO
5-(2-Furanyl)uridine TP	--	U	NO
5-Cyanouridine TP	--	U	NO
5-Dimethylaminouridine TP	--	U	NO
5'-Homo-uridine TP	--	U	NO
5-iodo-2'-fluoro-deoxyuridine TP	--	U	NO
5-Phenylethynyluridine TP	--	U	NO
5-Trideuteromethyl-6-deuterouridine TP	--	U	NO
5-Trifluoromethyl-Uridine TP	--	U	NO
5-Vinylarauridine TP	--	U	NO
6-(2,2,2-Trifluoroethyl)-pseudo-UTP	--	U	NO
6-(4-Morpholino)-pseudo-UTP	--	U	NO
6-(4-Thiomorpholino)-pseudo-UTP	--	U	NO

6-(Substituted-Phenyl)-pseudo-UTP	--	U	NO
6-Amino-pseudo-UTP	--	U	NO
6-Azido-pseudo-UTP	--	U	NO
6-Bromo-pseudo-UTP	--	U	NO
6-Butyl-pseudo-UTP	--	U	NO
6-Chloro-pseudo-UTP	--	U	NO
6-Cyano-pseudo-UTP	--	U	NO
6-Dimethylamino-pseudo-UTP	--	U	NO
6-Ethoxy-pseudo-UTP	--	U	NO
6-Ethylcarboxylate-pseudo-UTP	--	U	NO
6-Ethyl-pseudo-UTP	--	U	NO
6-Fluoro-pseudo-UTP	--	U	NO
6-Formyl-pseudo-UTP	--	U	NO
6-Hydroxyamino-pseudo-UTP	--	U	NO
6-Hydroxy-pseudo-UTP	--	U	NO
6-Iodo-pseudo-UTP	--	U	NO
6-iso-Propyl-pseudo-UTP	--	U	NO
6-Methoxy-pseudo-UTP	--	U	NO
6-Methylamino-pseudo-UTP	--	U	NO
6-Methyl-pseudo-UTP	--	U	NO
6-Phenyl-pseudo-UTP	--	U	NO
6-Phenyl-pseudo-UTP	--	U	NO
6-Propyl-pseudo-UTP	--	U	NO
6-tert-Butyl-pseudo-UTP	--	U	NO
6-Trifluoromethoxy-pseudo-UTP	--	U	NO
6-Trifluoromethyl-pseudo-UTP	--	U	NO
Alpha-thio-pseudo-UTP	--	U	NO
Pseudouridine 1-(4-methylbenzenesulfonic acid) TP	--	U	NO
Pseudouridine 1-(4-methylbenzoic acid) TP	--	U	NO
Pseudouridine TP 1-[3-(2-ethoxy)]propionic acid	--	U	NO
Pseudouridine TP 1-[3-{2-(2-[2-(2-ethoxy)-ethoxy]-ethoxy)-ethoxy}]propionic acid	--	U	NO
Pseudouridine TP 1-[3-{2-(2-[2-{2(2-ethoxy)-ethoxy}-ethoxy]-ethoxy)-ethoxy}]propionic acid	--	U	NO
Pseudouridine TP 1-[3-{2-(2-[2-ethoxy]-ethoxy)-ethoxy}]propionic acid	--	U	NO
Pseudouridine TP 1-[3-{2-(2-ethoxy)-ethoxy}] propionic acid	--	U	NO
Pseudouridine TP 1-methylphosphonic acid	--	U	NO
Pseudouridine TP 1-methylphosphonic acid diethyl ester	--	U	NO

Pseudo-UTP-N1-3-propionic acid	--	U	NO
Pseudo-UTP-N1-4-butanoic acid	--	U	NO
Pseudo-UTP-N1-5-pentanoic acid	--	U	NO
Pseudo-UTP-N1-6-hexanoic acid	--	U	NO
Pseudo-UTP-N1-7-heptanoic acid	--	U	NO
Pseudo-UTP-N1-methyl-p-benzoic acid	--	U	NO
Pseudo-UTP-N1-p-benzoic acid	--	U	NO
Wybutosine	yW	W	YES
Hydroxywybutosine	OHyW	W	YES
Isowyosine	imG2	W	YES
Peroxywybutosine	o2yW	W	YES
undermodified hydroxywybutosine	OHyW*	W	YES
4-demethylwyosine	imG-14	W	YES

[0484] Other modifications which may be useful in the polynucleotides of the NAVs of the present invention are listed in Table 23.

Table 23. Additional Modification types

Name	Type
2,6-(diamino)purine	Other
1-(aza)-2-(thio)-3-(aza)-phenoxazin-1-yl	Other
1,3-(diaza)-2-(oxo)-phenthiazin-l-yl	Other
1,3-(diaza)-2-(oxo)-phenoxazin-1-yl	Other
1,3,5-(triaza)-2,6-(dioxa)-naphthalene	Other
2 (amino)purine	Other
2,4,5-(trimethyl)phenyl	Other
2' methyl, 2'amino, 2'azido, 2'fluro-cytidine	Other
2' methyl, 2'amino, 2'azido, 2'fluro-adenine	Other
2'methyl, 2'amino, 2'azido, 2'fluro-uridine	Other
2'-amino-2'-deoxyribose	Other
2-amino-6-Chloro-purine	Other
2-aza-inosinyl	Other
2'-azido-2'-deoxyribose	Other
2'fluoro-2'-deoxyribose	Other
2'-fluoro-modified bases	Other
2'-O-methyl-ribose	Other
2-oxo-7-aminopyridopyrimidin-3-yl	Other
2-oxo-pyridopyrimidine-3-yl	Other
2-pyridinone	Other
3 nitropyrrole	Other
3-(methyl)-7-(propynyl)isocarbostyrilyl	Other
3-(methyl)isocarbostyrilyl	Other
4-(fluoro)-6-(methyl)benzimidazole	Other
4-(methyl)benzimidazole	Other
4-(methyl)indolyl	Other

4,6-(dimethyl)indolyl	Other
5 nitroindole	Other
5 substituted pyrimidines	Other
5-(methyl)isocarbostyrilyl	Other
5-nitroindole	Other
6-(aza)pyrimidine	Other
6-(azo)thymine	Other
6-(methyl)-7-(aza)indolyl	Other
6-chloro-purine	Other
6-phenyl-pyrrolo-pyrimidin-2-on-3-yl	Other
7-(aminoalkylhydroxy)-1-(aza)-2-(thio)-3-(aza)-phenthiazin-l-yl	Other
7-(aminoalkylhydroxy)-1-(aza)-2-(thio)-3-(aza)-phenoxazin-1-yl	Other
7-(aminoalkylhydroxy)-1,3-(diaza)-2-(oxo)-phenoxazin-1-yl	Other
7-(aminoalkylhydroxy)-l,3-(diaza)-2-(oxo)-phenthiazin-l-yl	Other
7-(aminoalkylhydroxy)-l,3-(diaza)-2-(oxo)-phenoxazin-l-yl	Other
7-(aza)indolyl	Other
7-(guanidiniumalkylhydroxy)-1-(aza)-2-(thio)-3-(aza)-phenoxazinl-yl	Other
7-(guanidiniumalkylhydroxy)-1-(aza)-2-(thio)-3-(aza)-phenthiazin-l-yl	Other
7-(guanidiniumalkylhydroxy)-1-(aza)-2-(thio)-3-(aza)-phenoxazin-1-yl	Other
7-(guanidiniumalkylhydroxy)-1,3-(diaza)-2-(oxo)-phenoxazin-1-yl	Other
7-(guanidiniumalkyl-hydroxy)-l,3-(diaza)-2-(oxo)-phenthiazin-l-yl	Other
7-(guanidiniumalkylhydroxy)-l,3-(diaza)-2-(oxo)-phenoxazin-l-yl	Other
7-(propynyl)isocarbostyrilyl	Other
7-(propynyl)isocarbostyrilyl, propynyl-7-(aza)indolyl	Other
7-deaza-inosinyl	Other
7-substituted 1-(aza)-2-(thio)-3-(aza)-phenoxazin-1-yl	Other
7-substituted 1,3-(diaza)-2-(oxo)-phenoxazin-1-yl	Other
9-(methyl)-imidizopyridinyl	Other
Aminoindolyl	Other
Anthracenyl	Other
bis-ortho-(aminoalkylhydroxy)-6-phenyl-pyrrolo-pyrimidin-2-on-3-yl	Other
bis-ortho-substituted-6-phenyl-pyrrolo-pyrimidin-2-on-3-yl	Other
Difluorotolyl	Other
Hypoxanthine	Other
Imidizopyridinyl	Other
Inosinyl	Other
Isocarbostyrilyl	Other

Isoguanisine	Other
N2-substituted purines	Other
N6-methyl-2-amino-purine	Other
N6-substituted purines	Other
N-alkylated derivative	Other
Napthalenyl	Other
Nitrobenzimidazolyl	Other
Nitroimidazolyl	Other
Nitroindazolyl	Other
Nitropyrazolyl	Other
Nubularine	Other
O6-substituted purines	Other
O-alkylated derivative	Other
ortho-(aminoalkylhydroxy)-6-phenyl-pyrrolo-pyrimidin-2-on-3-yl	Other
ortho-substituted-6-phenyl-pyrrolo-pyrimidin-2-on-3-yl	Other
Oxoformycin TP	Other
para-(aminoalkylhydroxy)-6-phenyl-pyrrolo-pyrimidin-2-on-3-yl	Other
para-substituted-6-phenyl-pyrrolo-pyrimidin-2-on-3-yl	Other
Pentacenyl	Other
Phenanthracenyl	Other
Phenyl	Other
propynyl-7-(aza)indolyl	Other
Pyrenyl	Other
pyridopyrimidin-3-yl	Other
pyridopyrimidin-3-yl, 2-oxo-7-amino-pyridopyrimidin-3-yl	Other
pyrrolo-pyrimidin-2-on-3-yl	Other
Pyrrolopyrimidinyl	Other
Pyrrolopyrizinyl	Other
Stilbenzyl	Other
substituted 1,2,4-triazoles	Other
Tetracenyl	Other
Tubercidine	Other
Xanthine	Other
Xanthosine-5'-TP	Other
2-thio-zebularine	Other
5-aza-2-thio-zebularine	Other
7-deaza-2-amino-purine	Other

pyridin-4-one ribonucleoside	Other
2-Amino-riboside-TP	Other
Formycin A TP	Other
Formycin B TP	Other
Pyrrolosine TP	Other
2'-OH-ara-adenosine TP	Other
2'-OH-ara-cytidine TP	Other
2'-OH-ara-uridine TP	Other
2'-OH-ara-guanosine TP	Other
5-(2-carbomethoxyvinyl)uridine TP	Other
N6-(19-Amino-pentaoxanonadecyl)adenosine TP	Other

[0485] The polynucleotides of the NAVs can include any useful linker between the nucleosides. Such linkers, including backbone modifications are given in Table 24.

Table 24. Linker modifications

Name	TYPE
3'-alkylene phosphonates	Linker
3'-amino phosphoramidate	Linker
alkene containing backbones	Linker
Aminoalkylphosphoramidates	Linker
Aminoalkylphosphotriesters	Linker
Boranophosphates	Linker
-CH2-0-N(CH3)-CH2-	Linker
-CH2-N(CH3)-N(CH3)-CH2-	Linker
-CH2-NH-CH2-	Linker
chiral phosphonates	Linker
chiral phosphorothioates	Linker
formacetyl and thioformacetyl backbones	Linker
methylene (methylimino)	Linker
methylene formacetyl and thioformacetyl backbones	Linker
methyleneimino and methylenehydrazino backbones	Linker
morpholino linkages	Linker
-N(CH3)-CH2-CH2-	Linker
oligonucleosides with heteroatom internucleoside linkage	Linker
Phosphinates	Linker
phosphoramidates	Linker
Phosphorodithioates	Linker
phosphorothioate internucleoside linkages	Linker
Phosphorothioates	Linker
Phosphotriesters	Linker
PNA	Linker
siloxane backbones	Linker
sulfamate backbones	Linker
sulfide sulfoxide and sulfone backbones	Linker
sulfonate and sulfonamide backbones	Linker
Thionoalkylphosphonates	Linker
Thionoalkylphosphotriesters	Linker
Thionophosphoramidates	Linker

[0486] The polynucleotides of the NAVs of the invention can include any useful modification, such as to the sugar, the nucleobase, or the internucleoside linkage (*e.g.* to a linking phosphate / to a phosphodiester linkage / to the phosphodiester backbone). One or more atoms of a pyrimidine nucleobase may be replaced or substituted with optionally substituted amino, optionally substituted thiol, optionally substituted alkyl (e.g., methyl or ethyl), or halo (e.g., chloro or fluoro). In certain embodiments, modifications (e.g., one or more modifications) are present in each of the sugar and the internucleoside linkage. Modifications according to the present invention may be modifications of ribonucleic acids (RNAs) to deoxyribonucleic acids (DNAs), threose nucleic acids (TNAs), glycol nucleic acids (GNAs), peptide nucleic acids (PNAs),

locked nucleic acids (LNAs) or hybrids thereof). Additional modifications are described herein. The at least one chemical modification of the present invention is set out in the appended claims.

[0487] Non-natural modified nucleotides may be introduced to polynucleotides, i.e., of the NAVs of the invention, or nucleic acids during synthesis or post-synthesis of the chains to achieve desired functions or properties. The modifications may be on internucleotide lineage, the purine or pyrimidine bases, or sugar. The modification may be introduced at the terminal of a chain or anywhere else in the chain; with chemical synthesis or with a polymerase enzyme. For example, hexitol nucleic acids (HNAs) are nuclease resistant and provide strong hybridization to RNA Short messenger RNAs (mRNAs) with hexitol residues in two codons have been constructed (Lavrik et al., Biochemistry, 40, 11777-11784 (2001)). The antisense effects of a chimeric HNA gapmer oligonucleotide comprising a phosphorothioate central sequence flanked by 5' and 3' HNA sequences have also been studied (See e.g., Kang et al., Nucleic Acids Research, vol. 32(4), 4411-4419 (2004)). The preparation and uses of modified nucleotides comprising 6-member rings in RNA interference, antisense therapy or other applications are disclosed in US Pat. Application No. 2008/0261905, US Pat. Application No. 201010009865, and PCT Application No. WO97/30064 to Herdewijn et al.). Modified nucleic acids and their synthesis are disclosed in copending PCT applications No. PCT/US2012/058519. The synthesis and strategy of modified polynucleotides is reviewed by Verma and Eckstein in Annual Review of Biochemistry, vol. 76, 99-134 (1998).

[0488] Either enzymatic or chemical ligation methods can be used to conjugate polynucleotides or their regions with different functional blocks, such as fluorescent labels, liquids, nanoparticles, delivery agents, etc. The conjugates of polynucleotides and modified polynucleotides are reviewed by Goodchild in Bioconjugate Chemistry, vol. 1(3), 165-187 (1990). US Pat. No. 6,835,827 and US Pat. No. 6,525,183 to Vinayak et al. teach synthesis of labeled oligonucleotides using a labeled solid support.

[0489] In certain embodiments, it may desirable to intracellularly degrade a polynulcleotide introduced into the cell. For example, degradation of a polynulcleofide may be preferable if precise timing of protein production is desired. Thus, in some embodiments, the invention provides a polynulcleotide containing a degradation domain, which is capable of being acted on in a directed manner within a cell.

[0490] Any of the regions of the polynucleotides may be chemically modified as taught herein or as taught in International Application Number PCT/2012/058519 filed October 3, 2012 and U.S. Provisional Application Number 61/837297 filed June 20, 2013.

Modified Polynucleotide Molecules

[0491] The present invention also includes building blocks, e.g., modified ribonucleosides, and modified ribonucleotides, of polynucleotide molecules, e.g., of the NAVs of the invention. For example, these building blocks can be useful for preparing the polynucleotides of the invention. Such building blocks are taught in International Application Number PCT/2012/058519 filed October 3, 2012 and U.S. Provisional Application Number 61/837297 filed June 20, 2013.

Modifications on the Sugar

[0492] The modified nucleosides and nucleotides (e.g., building block molecules), which may be incorporated into a polynucleotide (e.g., RNA or mRNA, as described herein), can be modified on the sugar of the ribonucleic acid. For example, the 2' hydroxyl group (OH) can be modified or replaced with a number of different substituents. Exemplary substitutions at the 2'-position include, but are not limited to, H, halo, optionally substituted C_{1-6} alkyl; optionally substituted C_{1-6} alkoxy; optionally substituted C_{6-10} aryloxy; optionally substituted C_{3-8} cycloalkyl; optionally substituted C_{3-8} cycloalkoxy; optionally substituted C_{6-10} aryloxy; optionally substituted C_{6-10} aryl-C_{1-6} alkoxy, optionally substituted C_{1-12} (heterocyclyl)oxy; a sugar (e.g., ribose, pentose, or any described herein); a polyethyleneglycol (PEG), $-O(CH_2CH_2O)_nCH_2CH_2OR$, where R is H or optionally substituted alkyl, and n is an integer from 0 to 20 (e.g., from 0 to 4, from 0 to 8, from 0 to 10, from 0 to 16, from 1 to 4, from 1 to 8, from 1 to 10, from 1 to 16, from 1 to 20, from 2 to 4, from 2 to 8, from 2 to 10, from 2 to 16, from 2 to 20, from 4 to 8, from 4 to 10, from 4 to 16, and from 4 to 20), "locked" nucleic acids (LNA) in which the 2'-hydroxyl is connected by a C_{1-6} alkylene or C_{1-6} heteroalkylene bridge to the 4'-carbon of the same ribose sugar, where exemplary bridges included methylene, propylene, ether, or amino bridges; aminoalkyl, as defined herein; aminoalkoxy, as defined herein; amino as defined herein; and amino acid, as defined herein

[0493] Generally, RNA includes the sugar group ribose, which is a 5-membered ring having an oxygen. Exemplary, non-limiting modified nucleotides include replacement of the oxygen in ribose (e.g., with S, Se, or alkylene, such as methylene or ethylene); addition of a double bond (e.g., to replace ribose with cyclopentenyl or cyclohexenyl); ring contraction of ribose (e.g., to form a 4-membered ring of cyclobutane or oxetane); ring expansion of ribose (e.g., to form a 6- or 7-membered ring having an additional carbon or heteroatom, such as for anhydrohexitol, altritol, mannitol, cyclohexanyl, cyclohexenyl, and morpholino that also has a phosphoramidate backbone); multicyclic forms (e.g., tricyclo; and "unlocked" forms, such as glycol nucleic acid (GNA) (e.g., R-GNA or S-GNA, where ribose is replaced by glycol units attached to phosphodiester bonds), threose nucleic acid (TNA, where ribose is replace with α-L-threofurano-

syl-(3'→2')), and peptide nucleic acid (PNA, where 2-amino-ethyl-glycine linkages replace the ribose and phosphodiester backbone). The sugar group can also contain one or more carbons that possess the opposite stereochemical configuration than that of the corresponding carbon in ribose. Thus, a polynucleotide molecule can include nucleotides containing, e.g., arabinose, as the sugar. Such sugar modifications are taught International Application Number PCT/2012/058519 filed October 3, 2012 and U.S. Provisional Application Number 61/837297 filed June 20, 2013.

Modifications on the Nucleobase

[0494] The present disclosure provides for modified nucleosides and nucleotides. As described herein "nucleoside" is defined as a compound containing a sugar molecule (e.g., a pentose or ribose) or a derivative thereof in combination with an organic base (e.g., a purine or pyrimidine) or a derivative thereof (also referred to herein as "nucleobase"). As described herein, "nucleotide" is defined as a nucleoside including a phosphate group. The modified nucleotides may by synthesized by any useful method, as described herein (e.g., chemically, enzymatically, or recombinantly to include one or more modified or non-natural nucleosides). The polynucleotides may comprise a region or regions of linked nucleosides. Such regions may have variable backbone linkages. The linkages may be standard phosphoester linkages, in which case the polynucleotides would comprise regions of nucleotides.

[0495] The modified nucleotide base pairing encompasses not only the standard adenosine-thymine, adenosine-uracil, or guanosine-cytosine base pairs, but also base pairs formed between nucleotides and/or modified nucleotides comprising non-standard or modified bases, wherein the arrangement of hydrogen bond donors and hydrogen bond acceptors permits hydrogen bonding between a non-standard base and a standard base or between two complementary non-standard base structures. One example of such non-standard base pairing is the base pairing between the modified nucleotide inosine and adenine, cytosine or uracil.

[0496] The modified nucleosides and nucleotides can include a modified nucleobase. Examples of nucleobases found in RNA include, but are not limited to, adenine, guanine, cytosine, and uracil. Examples of nucleobase found in DNA include, but are not limited to, adenine, guanine, cytosine, and thymine. Such modified nucleobases (including the distinctions between naturally occurring and non-naturally occurring) are taught in International Application Number PCT/2012/058519 filed October 3, 2012 and U.S. Provisional Application Number 61/837297 filed June 20, 2013.

Combinations of Modified Sugars, Nucleobases, and Internucleoside Linkages

[0497] The polynucleotides of the invention, i.e., the NAVs of the invention, can include a combination of modifications to the sugar, the nucleobase, and/or the internucleoside linkage. These combinations can include any one or more modifications described herein. The at least one chemical modification of the present invention is set out in the appended claims.

[0498] Examples of modified nucleotides and modified nucleotide combinations are provided below in Table 25. These combinations of modified nucleotides can be used to form the polynucleotides of the invention. Unless otherwise noted, the modified nucleotides may be completely substituted for the natural nucleotides of the polynucleotides of the invention. As a non-limiting example, the natural nucleotide uridine may be substituted with a modified nucleoside described herein. In another non-limiting example, the natural nucleotide uridine may be partially substituted (e.g., about 0.1%, 1%, 5%, 10%, 15%, 20%, 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95% or 99.9%) with at least one of the modified nucleoside disclosed herein. Any combination of base/sugar or linker may be incorporated into the polynucleotides of the invention and such modifcations are taught in International Application Number PCT/2012/058519 filed October 3, 2012 and U.S. Provisional Application Number 61/837297 filed June 20,.

Table 25. Combinations

Modified Nucleotide	Modified Nucleotide Combination
α-thio-cytidine	α-thio-cytidine/5-iodo-uridine
	α-thio-cytidine/N1-methyl-pseudouridine
	α-thio-cytidine/α-thio-uridine
	α-thio-cytidine/5-methyl-uridine
	α-thio-cytidine/pseudo-uridine
	about 50% of the cytosines are α-thio-cytidine
Pseudoisocytidine	pseudoisocytidine/5-iodo-uridine
	pseudoisocytidine/N1-methyl-pseudouridine
	pseudoisocytidine/α-thio-uridine
	pseudoisocytidine/5-methyl-uridine
	pseudoisocytidine/pseudouridine
	about 25% of cytosines are pseudoisocytidine
	pseudoisocytidine/about 50% of uridines are N1-methyl-pseudouridine and about 50% of uridines are pseudouridine
	pseudoisocytidine/about 25% of uridines are N1-methyl-pseudouridine and about 25% of uridines are pseudouridine
pyrrolo-cytidine	pyrrolo-cytidine/5-iodo-uridine
	pyrrolo-cytidine/N1-methyl-pseudouridine
	pyrrolo-cytidine/α-thio-uridine
	pyrrolo-cytidine/5-methyl-uridine
	pyrrolo-cytidine/pseudouridine
	about 50% of the cytosines are pyrrolo-cytidine
5-methyl-cytidine	5-methyl-cytidine/5-iodo-uridine
	5-methyl-cytidine/N1-methyl-pseudouridine
	5-methyl-cytidine/α-thio-uridine
	5-methyl-cytidine/5-methyl-uridine
	5-methyl-cytidine/pseudouridine
	about 25% of cytosines are 5-methyl-cytidine
	about 50% of cytosines are 5-methyl-cytidine
	5-methyl-cytidine/5-methoxy-uridine
	5-methyl-cytidine/5-bromo-uridine
	5-methyl-cytidine/2-thio-uridine
	5-methyl-cytidine/about 50% of uridines are 2-thio-uridine
	about 50% of uridines are 5-methyl-cytidine/ about 50% of uridines are 2-thio-uridine
N4-acetyl-cytidine	N4-acetyl-cytidine /5-iodo-uridine
	N4-acetyl-cytidine /N1-methyl-pseudouridine
	N4-acetyl-cytidine /α-thio-uridine
	N4-acetyl-cytidine /5-methyl-uridine
	N4-acetyl-cytidine /pseudouridine
	about 50% of cytosines are N4-acetyl-cytidine
	about 25% of cytosines are N4-acetyl-cytidine
	N4-acetyl-cytidine /5-methoxy-uridine
	N4-acetyl-cytidine /5-bromo-uridine
	N4-acetyl-cytidine /2-thio-uridine
	about 50% of cytosines are N4-acetyl-cytidine/ about 50% of uridines are 2-thio-uridine

V. Pharmaceutical Vaccine Compositions

Formulation, Administration, Delivery and Dosing

[0499] The present disclosure provides pharmaceutical compositions including NAVs and NAV compositions and/or complexes optionally in combination with one or more pharmaceutically acceptable excipients.

[0500] The present inventors have discovered that NAVs are superior to current vaccines in several ways. First, subcutaneous and/or intradermal injection is better than intramuscular administration as a route of delivery. Second, the lipid nanoparticle delivery is superior to other formulations including the protamine approach in the literature by a factor of between 10-100 fold and no additional adjuvants were found to be necessary. Third modified and formulated NAVs were superior to unmodified formulated NAVs by a factor of 50 fold.

[0501] The present disclosure provides NAVs and NAV pharmaceutical compositions and complexes optionally in combination with one or more pharmaceutically acceptable excipients. Pharmaceutical compositions may optionally comprise one or more additional active substances, e.g. therapeutically and/or prophylactically active substances. Pharmaceutical compositions of the present disclosure may be sterile and/or pyrogen-free. General considerations in the formulation and/or manufacture of pharmaceutical agents may be found, for example, in Remington: The Science and Practice of Pharmacy 21st ed, Lippincott Williams & Wilkins, 2005.

[0502] In some instances, compositions are administered to humans, human patients or subjects. For the purposes of the present disclosure, the phrase "active ingredient" generally refers to the NAVs or the polynucleotides contained therein, e.g., antigen-encoding polynucleotides, for example, RNA polynucleotides, to be delivered as described herein.

[0503] Although the descriptions of pharmaceutical compositions provided herein are principally directed to pharmaceutical compositions which are suitable for administration to humans, it will be understood by the skilled artisan that such compositions are generally suitable for administration to any other animal, *e.g.*, to non-human animals, e.g. non-human mammals. Modification of pharmaceutical compositions suitable for administration to humans in order to render the compositions suitable for administration to various animals is well understood, and the ordinarily skilled veterinary pharmacologist can design and/or perform such modification with merely ordinary, if any, experimentation. Subjects to which administration of the pharmaceutical compositions is contemplated include, but are not limited to, humans and/or other primates; mammals, including commercially relevant mammals such as cattle, pigs, horses, sheep, cats, dogs, mice, and/or rats; and/or birds, including commercially relevant birds such as poultry, chickens, ducks, geese, and/or turkeys.

[0504] Formulations of the pharmaceutical compositions described herein may be prepared by any method known or hereafter developed in the art of pharmacology. In general, such preparatory methods include the step of bringing the active ingredient into association with an excipient and/or one or more other accessory ingredients, and then, if necessary and/or desirable, dividing, shaping and/or packaging the product into a desired single- or multi-dose unit.

[0505] Relative amounts of the active ingredient, the pharmaceutically acceptable excipient, and/or any additional ingredients in a pharmaceutical composition in accordance with the disclosure will vary, depending upon the identity, size, and/or condition of the subject treated and further depending upon the route by which the composition is to be administered. By way of example, the composition may comprise between 0.1% and 100%, e.g., between 0.5 and 50%, between 1-30%, between 5-80%, at least 80% (w/w) active ingredient.

Formulations

[0506] The NAVs of the disclosure can be formulated using one or more excipients to (1) increase stability; (2) increase cell transfection; (3) permit the sustained or delayed release (e.g., from a depot formulation); (4) alter the biodistribution (e.g., target to specific tissues or cell types); (5) increase the translation of encoded protein *in vivo;* and/or (6) alter the release profile of encoded protein (antigen) in vivo. In addition to traditional excipients such as any and all solvents, dispersion media, diluents, or other liquid vehicles, dispersion or suspension aids, surface active agents, isotonic agents, thickening or emulsifying agents, preservatives, excipients of the present invention can include, without limitation, lipidoids, liposomes, lipid nanoparticles, polymers, lipoplexes, core-shell nanoparticles, peptides, proteins, cells transfected with NAVs (e.g., for transplantation into a subject), hyaluronidase, nanoparticle mimics and combinations thereof.

[0507] Accordingly, the formulations of the disclosure can include one or more excipients, each in an amount that may increases the stability of the NAV, increases cell transfection by the NAV, increases the expression of polynucleotides encoded protein, and/or alters the release profile of polynucleotide encoded proteins. Further, the polynucleotides of the present disclosure may be formulated using self-assembled nucleic acid nanoparticles.

[0508] Formulations of the pharmaceutical compositions described herein may be prepared by any method known or hereafter developed in the art of pharmacology. In general, such preparatory methods include the step of associating the active ingredient with an excipient and/or one or more other accessory ingredients.

[0509] A pharmaceutical composition in accordance with the present disclosure may be prepared, packaged, and/or

sold in bulk, as a single unit dose, and/or as a plurality of single unit doses. As used herein, a "unit dose" refers to a discrete amount of the pharmaceutical composition comprising a predetermined amount of the active ingredient. The amount of the active ingredient is generally equal to the dosage of the active ingredient which would be administered to a subject and/or a convenient fraction of such a dosage such as, for example, one-half or one-third of such a dosage.

[0510] Relative amounts of the active ingredient, the pharmaceutically acceptable excipient, and/or any additional ingredients in a pharmaceutical composition in accordance with the present disclosure may vary, depending upon the identity, size, and/or condition of the subject being treated and further depending upon the route by which the composition is to be administered. For example, the composition may comprise between 0.1% and 99% (w/w) of the active ingredient. By way of example, the composition may comprise between 0.1% and 100%, e.g., between .5 and 50%, between 1-30%, between 5-80%, at least 80% (w/w) active ingredient.

[0511] In some instances, the formulations described herein may contain at least one polynucleotide, *e.g.*, antigen-encoding polynucleotide. As a non-limiting example, the formulations may contain 1, 2, 3, 4 or 5 polynucleotides.

[0512] In one instance, the formulations described herein may comprise more than one type of polynucleotide, *e.g.*, antigen-encoding polynucleotide. In one instance, the formulation may comprise a chimeric polynucleotide in linear and circular form. In another instance, the formulation may comprise a circular polynucleotide and an IVT polynucleotide. In yet another instance, the formulation may comprise an IVT polynucleotide, a chimeric polynucleotide and a circular polynucleotide.

[0513] In one instancethe formulation may contain polynucleotide encoding proteins selected from categories such as, but not limited to, human proteins, veterinary proteins, bacterial proteins, biological proteins, antibodies, immunogenic proteins, therapeutic peptides and proteins, secreted proteins, plasma membrane proteins, cytoplasmic and cytoskeletal proteins, intracellular membrane bound proteins, nuclear proteins, proteins associated with human disease and/or proteins associated with non-human diseases. In one instance, the formulation contains at least three polynucleotides encoding proteins. In one instance, the formulation contains at least five polynucleotide encoding proteins.

[0514] Pharmaceutical formulations may additionally comprise a pharmaceutically acceptable excipient, which, as used herein, includes, but is not limited to, any and all solvents, dispersion media, diluents, or other liquid vehicles, dispersion or suspension aids, surface active agents, isotonic agents, thickening or emulsifying agents, preservatives, and the like, as suited to the particular dosage form desired. Various excipients for formulating pharmaceutical compositions and techniques for preparing the composition are known in the art (see Remington: The Science and Practice of Pharmacy, 21st Edition, A. R. Gennaro, Lippincott, Williams & Wilkins, Baltimore, MD, 2006). The use of a conventional excipient medium may be contemplated within the scope of the present disclosure, except insofar as any conventional excipient medium may be incompatible with a substance or its derivatives, such as by producing any undesirable biological effect or otherwise interacting in a deleterious manner with any other component(s) of the pharmaceutical composition.

[0515] In some instances, the particle size of the lipid nanoparticle may be increased and/or decreased. The change in particle size may be able to help counter biological reaction such as, but not limited to, inflammation or may increase the biological effect of the modified mRNA delivered to mammals.

[0516] Pharmaceutically acceptable excipients used in the manufacture of pharmaceutical compositions include, but are not limited to, inert diluents, surface active agents and/or emulsifiers, preservatives, buffering agents, lubricating agents, and/or oils. Such excipients may optionally be included in the pharmaceutical formulations of the disclosure.

Lipid Nanoparticles

[0517] The NAVs of the invention are formulated using one or more lipid nanoparticles.

[0518] In some embodiments, the ratio of PEG in the lipid nanoparticle (LNP) formulations may be increased or decreased and/or the carbon chain length of the PEG lipid may be modified from C14 to C18 to alter the pharmacokinetics and/or biodistribution of the LNP formulations. As a non-limiting example, LNP formulations may contain from about 0.5% to about 3.0%, from about 1.0% to about 3.5%, from about 1.5% to about 40%, from about 2.0% to about 45%, from about 2.5% to about 5.0% and/or from about 3.0% to about 6.0% of the lipid molar ratio of PEG-c-DOMG (R-3-[(w-methoxy-poly(ethyleneglycol)2000)carbamoyl)]-1,2-dimyristyloxypropyl-3-amine) (also referred to herein as PEG-DOMG) as compared to the cationic lipid, DSPC and cholesterol. In another embodiment the PEG-c-DOMG may be replaced with a PEG lipid such as, but not limited to, PEG- DSG (1,2-Distearoyl-sn-glycerol, methoxypolyethylene glycol), PEG-DMG (1,2-Dimyristoyl-sn-glycerol) and/or PEG-DPG (1,2-Dipalmitoyl-sn-glycerol, methoxypolyethylene glycol). The cationic lipid may be selected from any lipid known in the art such as, but not limited to, DLin-MC3-DMA, DLin-DMA, C12-200 and DLin-KC2-DMA.

[0519] In one embodiment, the NAVs may be formulated in a lipid nanoparticle such as those described in International Publication No. WO2012170930.

[0520] In one instance, the NAV formulation comprising the polynucleotide is a nanoparticle which may comprise at least one lipid. The lipid may be selected from, but is not limited to, DLin-DMA, DLin-K-DMA, 98N12-5, C12-200, DLin-MC3-DMA, DLin-KC2-DMA, DODMA, PLGA, PEG, PEG-DMG, PEGylated lipids and amino alcohol lipids. In another

aspect, the lipid may be a cationic lipid such as, but not limited to, DLin-DMA, DLin-D-DMA, DLin-MC3-DMA, DLin-KC2-DMA, DODMA and amino alcohol lipids. The amino alcohol cationic lipid may be the lipids described in and/or made by the methods described in US Patent Publication No. US20130150625. As a non-limiting example, the cationic lipid may be 2-amino-3-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]-2-{[(9Z,2Z)-ocfadeca-9,12-dien-1-yloxy]mefhyl}propan-1-ol (Compound 1 in US20130150625); 2-amino-3-[(9Z)-octadec-9-en-1-yloxy]-2-{[(9Z)-octadec-9-en-1-yloxy]methyl}propan-1-ol (Compound 2 in US20130150625); 2-amino-3-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]-2-[(octyloxy)methyl]propan-1-ol (Compound 3 in US20130150625); and 2-(dimethylamino)-3-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]-2-{[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]mefhyl}propan-1-ol (Compound 4 in US20130150625); or any pharmaceutically acceptable salt or stereoisomer thereof.

[0521] Lipid nanoparticle formulations typically comprise a lipid, in particular, an ionizable cationic lipid, for example, 2,2-dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), or di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319), and further comprise a neutral lipid, a sterol and a molecule capable of reducing particle aggregation, for example a PEG or PEG-modified lipid.

[0522] In one embodiment, the lipid nanoparticle formulation consists essentially of (i) at least one lipid selected from the group consisting of 2,2-dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), and di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319); (ii) a neutral lipid selected from DSPC, DPPC, POPC, DOPE and SM; (iii) a sterol, e.g., cholesterol; and (iv) a PEG-lipid, e.g., PEG-DMG or PEG-cDMA, in a molar ratio of about 20-60% cationic lipid: 5-25% neutral lipid: 25-55% sterol; 0.5-15% PEG-lipid.

[0523] In one instance, the formulation includes from about 25% to about 75% on a molar basis of a cationic lipid selected from 2,2-dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), and di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319), e.g., from about 35 to about 65%, from about 45 to about 65%, about 60%, about 57.5%, about 50% or about 40% on a molar basis. In the invention, the molar ration of the lipid nanoparticle is set out in the appended claims.

[0524] In one instance, the formulation includes from about 0.5% to about 15% on a molar basis of the neutral lipid e.g., from about 3 to about 12%, from about 5 to about 10% or about 15%, about 10%, or about 7.5% on a molar basis. Exemplary neutral lipids include, but are not limited to, DSPC, POPC, DPPC, DOPE and SM. In one instance, the formulation includes from about 5% to about 50% on a molar basis of the sterol (*e.g.*, about 15 to about 45%, about 20 to about 40%, about 40%, about 38.5%, about 35%, or about 31% on a molar basis. An exemplary sterol is cholesterol. In one instance, the formulation includes from about 0.5% to about 20% on a molar basis of the PEG or PEG-modified lipid (e.g., about 0.5 to about 10%, about 0.5 to about 5%, about 1.5%, about 0.5%, about 1.5%, about 3.5%, or about 5% on a molar basis. In one embodiment, the PEG or PEG modified lipid comprises a PEG molecule of an average molecular weight of 2,000 Da. In other embodiments, the PEG or PEG modified lipid comprises a PEG molecule of an average molecular weight of less than 2,000, for example around 1,500 Da, around 1,000 Da, or around 500 Da. Exemplary PEG-modified lipids include, but are not limited to, PEG-disfearoyl glycerol (PEG-DMG) (also referred herein as PEG-C14 or C14-PEG), PEG-cDMA (further discussed in Reyes et al. J. Controlled Release, 107, 276-287 (2005). In the invention, the molar ration of the lipid nanoparticle is set out in the appended claims.

[0525] In one instance, the formulations of the disclosure include 25-75% of a cationic lipid selected from 2,2-dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), and di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319), 0.5-15% of the neutral lipid, 5-50% of the sterol, and 0.5-20% of the PEG or PEG-modified lipid on a molar basis.

[0526] In one instance, the formulations of the disclosure include 35-65% of a cationic lipid selected from 2,2-dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), and di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319), 3-12% of the neutral lipid, 15-45% of the sterol, and 0.5-10% of the PEG or PEG-modified lipid on a molar basis.

[0527] In one embodiment, the formulations of the inventions include 45-65% of a cationic lipid selected from 2,2-dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), and di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319), 5-10% of the neutral lipid, 25-40% of the sterol, and 0.5-10% of the PEG or PEG-modified lipid on a molar basis.

[0528] In one embodiment, the formulations of the inventions include about 60% of a cationic lipid selected from 2,2-dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), and di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319), about 7.5% of the neutral lipid, about 31 % of the sterol, and about 1.5% of the PEG or PEG-modified lipid on a molar basis.

[0529] In one embodiment, the formulations of the inventions include about 50% of a cationic lipid selected from 2,2-dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), and di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319), about 10% of the neutral lipid, about 38.5 % of the sterol, and about 1.5% of the PEG or PEG-modified lipid on a molar basis.

[0530] In one embodiment, the formulations of the inventions include about 50% of a cationic lipid selected from 2,2-

dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), and di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319), about 10% of the neutral lipid, about 35 % of the sterol, about 4.5% or about 5% of the PEG or PEG-modified lipid, and about 0.5% of the targeting lipid on a molar basis.

[0531] In one embodiment, the formulations of the inventions include about 40% of a cationic lipid selected from 2,2-dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), and di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319), about 15% of the neutral lipid, about 40% of the sterol, and about 5% of the PEG or PEG-modified lipid on a molar basis.

[0532] In one embodiment, the formulations of the inventions include about 57.2% of a cationic lipid selected from 2,2-dilinoleyl-4-dimethylaminoethyl-[1,3]-dioxolane (DLin-KC2-DMA), dilinoleyl-methyl-4-dimethylaminobutyrate (DLin-MC3-DMA), and di((Z)-non-2-en-1-yl) 9-((4-(dimethylamino)butanoyl)oxy)heptadecanedioate (L319), about 7.1% of the neutral lipid, about 34.3% of the sterol, and about 1.4% of the PEG or PEG-modified lipid on a molar basis.

[0533] In one embodiment, the formulations of the inventions include about 57.5% of a cationic lipid selected from the PEG lipid is PEG-cDMA (PEG-cDMA is further discussed in Reyes et al. (J. Controlled Release, 107, 276-287 (2005)), about 7.5% of the neutral lipid, about 31.5 % of the sterol, and about 3.5% of the PEG or PEG-modified lipid on a molar basis.

[0534] In particular embodiments, the molar lipid ratio is approximately 50110138.511.5 (mol% cationic lipid/neutral lipid, *e.g.*, DSPC/Chol/PEG-modified lipid, *e.g.*, PEG-DMG, PEG-DSG or PEG-DPG), 57.217.1134.311.4 (mol% cationic lipid/ neutral lipid, *e.g.*, DPPC/Chol/ PEG-modified lipid, *e.g.*, PEG-cDMA), 40/15/40/5 (mol% cationic lipid/ neutral lipid, *e.g.*, DSPC/Chol/ PEG-modified lipid, *e.g.*, PEG-DMG), 50/10/35/4.5/0.5 (mol% cationic lipid/ neutral lipid, *e.g.*, DSPC/Chol/ PEG-modified lipid, *e.g.*, PEG-DSG), 50/10/35/5 (cationic lipid/ neutral lipid, *e.g.*, DSPC/Chol/ PEG-modified lipid, *e.g.*, PEG-DMG), 40/10/40/10 (mol% cationic lipid/ neutral lipid, *e.g.,* DSPC/Chol/ PEG-modified lipid, *e.g.,* PEG-DMG or PEG-cDMA), 35/15/40/10 (mol% cationic lipid/ neutral lipid, *e.g.,* DSPC/Chol/ PEG-modified lipid, *e.g.,* PEG-DMG or PEG-cDMA) or 52/13/30/5 (mol% cationic lipid/ neutral lipid, *e.g.,* DSPC/Chol/ PEG-modified lipid, *e.g.,* PEG-DMG or PEG-cDMA).

[0535] Exemplary lipid nanoparticle compositions and methods of making same are described, for example, in Semple et al. (2010) Nat. Biotechnol. 28:172-176; Jayarama et al. (2012), Angew. Chem. Int. Ed., 51: 8529-8533, and Maier et al. (2013) Molecular Therapy 21, 1570-1578

[0536] The lipid nanoparticle formulations may be 4 component lipid nanoparticles. The lipid nanoparticle may comprise a cationic lipid, a non-cationic lipid, a PEG lipid and a structural lipid. As a non-limiting example, the lipid nanoparticle may comprise about 40-60% of cationic lipid, about 5-15% of a non-cationic lipid, about 1-2% of a PEG lipid and about 30-50% of a structural lipid. As another non-limiting example, the lipid nanoparticle may comprise about 50% cationic lipid, about 10% non-cationic lipid, about 1.5% PEG lipid and about 38.5% structural lipid. As yet another non-limiting example, the lipid nanoparticle may comprise about 55% cationic lipid, about 10% non-cationic lipid, about 2.5% PEG lipid and about 32.5% structural lipid. In one embodiment, the cationic lipid may be any cationic lipid described herein such as, but not limited to, DLin-KC2-DMA, DLin-MC3-DMA and L319.

[0537] The lipid nanoparticle formulations comprise a cationic lipid, a non-cationic lipid, a PEG lipid and a structural lipid. As a non-limiting example, the lipid nanoparticle comprise about 50% of the cationic lipid DLin-KC2-DMA, about 10% of the non-cationic lipid DSPC, about 1.5% of the PEG lipid PEG-DOMG and about 38.5% of the structural lipid cholesterol. As a non-limiting example, the lipid nanoparticle comprise about 50% of the cationic lipid DLin-MC3-DMA, about 10% of the non-cationic lipid DSPC, about 1.5% of the PEG lipid PEG-DOMG and about 38.5% of the structural lipid cholesterol. As a non-limiting example, the lipid nanoparticle comprise about 50% of the cationic lipid DLin-MC3-DMA, about 10% of the non-cationic lipid DSPC, about 1.5% of the PEG lipid PEG-DMG and about 38.5% of the structural lipid cholesterol. As yet another non-limiting example, the lipid nanoparticle comprise about 55% of the cationic lipid L319, about 10% of the non-cationic lipid DSPC, about 2.5% of the PEG lipid PEG-DMG and about 32.5% of the structural lipid cholesterol.

[0538] In one embodiment, the cationic lipid may be selected from, but not limited to, a cationic lipid described in International Publication Nos. WO2012040184, WO2011153120, WO2011149733, WO2011090965, WO2011043913, WO2011022460, WO2012061259, WO2012054365, WO2012044638, WO2010080724, WO201021865, WO2008103276, WO2013086373 and WO2013086354, US Patent Nos. 7,893,302, 7,404,969, 8,283,333, and 8,466,122 and US Patent Publication No. US20100036115, US20120202871, US20130064894, US20130129785, US20130150625, US20130178541 and US20130225836. In another embodiment, the cationic lipid may be selected from, but not limited to, formula A described in International Publication Nos. WO2012040184, WO2011153120, WO2011149733, WO2011090965, WO2011043913, WO2011022460, WO2012061259, WO2012054365, WO2012044638 and WO2013116126 or US Patent Publication No. US20130178541 and US20130225836. In yet another embodiment, the cationic lipid may be selected from, but not limited to, formula CLI-CLXXIX of International Publication No. WO2008103276, formula CLI-CLXXIX of US Patent No. 7,893,302, formula CLI-CLXXXXII of US Patent No. 7,404,969 and formula I-VI of US Patent Publication No. US20100036115, formula I of US Patent Publication No

US20130123338. As a non-limiting example, the cationic lipid may be selected from (20Z,23Z)-N,N-dimethylnonacosa-20,23-dien-10-amine, (17Z,20Z)-N,N-dimemylhexacosa-17,20-dien-9-amine, (1Z,19Z)-N5N-dimethylpentacosa-l6, 19-dien-8-amine, (13Z,16Z)-N,N-dimethyldocosa-13,16-dien-5-amine, (12Z, 15Z)-N,N-dimethylhenicosa-12, 15-dien-4-amine, (14Z, 17Z)-N,N-dimethyltricosa-14, 17-dien-6-amine, (15Z,18Z)-N,N-dimethyltetracosa-15,18-dien-7-amine, (18Z,21Z)-N,N-dimethylheptacosa-18,21-dien-10-amine, (15Z,18Z)-N,N-dimethyltetracosa-15,18-dien-5-amine, (14Z, 17Z)-N,N-dimethyltricosa-14, 17-dien-4-amine, (19Z,22Z)-N,N-dimeihyloctacosa-19,22-dien-9-amine, (18Z,21 Z)-N,N-dimethylheptacosa- 18 21 -dien-8 -amine, (17Z,20Z)-N,N-dimethylhexacosa- 17,20-dien-7-amine, (16Z,19Z)-N,N-dimethylpentacosa-16,19-dien-6-amine, (22Z,25Z)-N,N-dimethylhentriaconta-22,25-dien-10-amine, (21 Z ,24Z)-N,N-dimethyltriaconta-21,24-dien-9-amine, (18Z)-N,N-dimetylheptacos-18-en-10-amine, (17Z)-N,N-dimethylhexacos-17-en-9-amine, (19Z,22Z)-N,N-dimethyloctacosa-19,22-dien-7-amine, N,N-dimethylheptacosan-10-amine, (20Z,23Z)-N-ethyl-N-methylnonacosa-20,23-dien-l0-amine, 1-[(11Z,14Z)-l-nonylicosa-11,14-dien-l-yl] pyrrolidine, (20Z)-N,N-dimethylheptacos-20-en-l0-amine, (15Z)-N,N-dimethyl eptacos-15-en-l0-amine, (14Z)-N,N-dimethylnonacos-14-en-l0-amine, (17Z)-N,N-dimethylnonacos-17-en-10-amine, (24Z)-N,N-dimethyltritriacont-24-en-l0-amine, (20Z)-N,N-dimethylnonacos-20-en-l0-amine, (22Z)-N,N-dimethylhentriacont-22-en-l0-amine, (16Z)-N,N-dimethylpentacos-16-en-8-amine, (12Z, 15Z)-N,N-dimethyl-2-nonylhenicosa-12, 15-dien-1-amine, (13Z,16Z)-N,N-dimethyl-3-nonyldocosa-l3,16-dien-l-amine, N,N-dimefhyl-l-[(IS,2R)-2-ocfylcyclopropyl] eptadecan-8-amine, 1-[(1S,2R)-2-hexylcyclopropyl]-N,N-dimethylnonadecan-10-amine, N,N-dimethyl-1-[(1S ,2R)-2-octylcyclopropyl]nonadecan-10-amine, N,N-dimethyl-21-[(IS,2R)-2-octylcyclopropyl]henicosan-l0-amine,N,N-dimethyl-1-[(1S,2S)-2-{[(IR,2R)-2-pentylcyclopropyl]methyl}cyclopropyl]nonadecan-10-amine,N,N-dimethyl-1-[(1S,2R)-2-octylcyclopropyl]hexadecan-8-amine, N,N-dimethyl-[(IR,2S)-2-undecylcyclopropyl]tetradecan-5-amine, N,N-dimethyl-3-{7-[(1S,2R)-2-octylcyclopropyl]heptyl} dodecan-1-amine, 1-[(1R,2S)-2-hepty lcyclopropyl]-N,N-dimethyloctadecan-9-amine, 1-[(1S,2R)-2-decylcyclopropyl]-N,N-dimethylpentadecan-6-amine, N,N-dimethyl-l-[(IS,2R)-2-octylcyclopropyl]pentadecan-8-amine, R-N,N-dimethyl-1-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]-3-(octyloxy)propan-2-amine, S-N,N-dimethyl-1-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]-3-(octyloxy)propan-2-amine, 1-{2-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]-1-[(octyloxy)methyl]ethyl}pyrrolidine, (2S)-N,N-dimethyl-1-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]-3-[(5Z)-oct-5-en-1-yloxy]propan-2-amine, 1-{2-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]-1-[(octyloxy)methyl]ethyl}azetidine, (2S)-1-(hexyloxy)-N,N-dimethyl-3-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]propan-2-amine, (2S)-1-(heptyloxy)-N,N-dimethyl-3-[(9Z, 12Z)-octadeca-9, 12-dien-1-yloxy]propan-2-amine, N,N-dimethyl-1-(nonyloxy)-3-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]propan-2-amine, N,N-dimethyl-1-[(9Z)-octadec-9-en-1-yloxy]-3-(octyloxy)propan-2-amine; (2S)-N,N-dimethyl-1-[(6Z,9Z,12Z)-octadeca-6,9,12-trien-1-yloxy]-3-(octyloxy)propan-2-amine, (2S)-1-[(11Z,14Z)-icosa-11,14-dien-1-yloxy]-N,N-dimethyl-3-(pentyloxy)propan-2-amine, (2S)-1-(hexyloxy)-3-[(11Z,14Z)-icosa-11,14-dien-1-yloxy]-N,N-dimethylpropan-2-amine, 1-[(11Z,14Z)-icosa-11,14-dien-1-yloxy]-N,N-dimethy1-3-(octyloxy)propan-2-amine, 1-[(13Z,16Z)-docosa-l3,16-dien-l-yloxy]-N,N-dimethyl-3-(octyloxy)propan-2-amine, (2S)-1-[(13Z,16Z)-docosa-13,16-dien-1-yloxy]-3-(hexyloxy)-N,N-dimethylpropan-2-amine, (2S)-1-[(13Z)-docos-13-en-1-yloxy]-3-(hexyloxy)-N,N-dimethylpropan-2-amine, 1-[(13Z)-docos-13-en-1-yloxy]-N,N-dimethyl-3-(octyloxy)propan-2-amine, 1-[(9Z)-hexadec-9-en-1-yloxy]-N,N-dimethyl-3-(octyloxy)propan-2-amine, (2R)-N,N-dimethyl-H(1-metoylo ctyl)oxy]-3-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]propan-2-amine, (2R)-1-[(3,7-dimethyloctyl)oxy]-N,N-dimethyl-3-[(9Z,12Z)-octadeca-9,12-dien-1-yloxy]propan-2-amine, N,N-dimethyl-1-(octyloxy)-3-({8-[(1S,2S)-2-{[(1R,2R)-2-pentylcyclopropyl]methyl}cyclopropyl]octyl}oxy)propan-2-amine, N,N-dimethyl-1-{[8-(2-oc1ylcyclopropyl)octyl]oxy}-3-(octyloxy)propan-2-amine and (1lE,20Z,23Z)-N,N-dimethylnonacosa-l1,20,2-trien-10-amine or a pharmaceutically acceptable salt or stereoisomer thereof.

[0539] In one embodiment, the lipid may be a cleavable lipid such as those described in International Publication No. WO2012170889.

[0540] In another embodiment, the lipid may be a cationic lipid such as, but not limited to, Formula (I) of U.S. Patent Application No. US20130064894.

[0541] In one embodiment, the cationic lipid may be synthesized by methods known in the art and/or as described in International Publication Nos. WO2012040184, WO2011153120, WO2011149733, WO2011090965, WO2011043913, WO2011022460, WO2012061259, WO2012054365, WO2012044638, WO2010080724, WO201021865, WO2013086373 and WO2013086354.

[0542] In another embodiment, the cationic lipid may be a trialkyl cationic lipid. Non-limiting examples of trialkyl cationic lipids and methods of making and using the trialkyl cationic lipids are described in International Patent Publication No. WO2013126803.

[0543] In one embodiment, the LNP formulations of the NAVs may contain PEG-c-DOMG at 3% lipid molar ratio. In another embodiment, the LNP formulations RNAVs may contain PEG-c-DOMG at 1.5% lipid molar ratio.

[0544] In one embodiment, the pharmaceutical compositions of the NAVs may include at least one of the PEGylated lipids described in International Publication No. WO2012099755.

[0545] In one embodiment, the LNP formulation may contain PEG-DMG 2000 (1,2-dimyristoyl-sn-glycero-3-phophoethanolamine-N-[methoxy(polyethylene glycol)-2000). In one embodiment, the LNP formulation may contain PEG-DMG 2000, a cationic lipid known in the art and at least one other component. In another embodiment, the LNP formulation

may contain PEG-DMG 2000, a cationic lipid known in the art, DSPC and cholesterol. As a non-limiting example, the LNP formulation may contain PEG-DMG 2000, DLin-DMA, DSPC and cholesterol. As another non-limiting example the LNP formulation may contain PEG-DMG 2000, DLin-DMA, DSPC and cholesterol in a molar ratio of 2:40:10:48 (see e.g., Geall et al., Nonviral delivery of self-amplifying RNA vaccines, PNAS 2012; PMID: 22908294).

[0546] The nanoparticle formulations may comprise a phosphate conjugate. The phosphate conjugate may increase *in vivo* circulation times and/or increase the targeted delivery of the nanoparticle. Phosphate conjugates for use with the present invention may be made by the methods described in International Application No. WO2013033438 or US Patent Publication No. US20130196948. As a non-limiting example, the phosphate conjugates may include a compound of any one of the formulas described in International Application No. WO2013033438.

[0547] The nanoparticle formulation may comprise a polymer conjugate. The polymer conjugate may be a water soluble conjugate. The polymer conjugate may have a structure as described in U.S. Patent Application No. 20130059360. In one aspect, polymer conjugates with the polynucleotides of the present invention may be made using the methods and/or segmented polymeric reagents described in U.S. Patent Application No. 20130072709. In another aspect, the polymer conjugate may have pendant side groups comprising ring moieties such as, but not limited to, the polymer conjugates described in US Patent Publication No. US20130196948.

[0548] The nanoparticle formulations may comprise a conjugate to enhance the delivery of nanoparticles of the present invention in a subject. Further, the conjugate may inhibit phagocytic clearance of the nanoparticles in a subject. In one aspect, the conjugate may be a "self' peptide designed from the human membrane protein CD47 (e.g., the "self' particles described by Rodriguez et al (Science 2013 339, 971-975)). As shown by Rodriguez et al. the self peptides delayed macrophage-mediated clearance of nanoparticles which enhanced delivery of the nanoparticles In another aspect, the conjugate may be the membrane protein CD47 (e.g., see Rodriguez et al. Science 2013 339, 971-975). Rodriguez et al. showed that, similarly to "self' peptides, CD47 can increase the circulating particle ratio in a subject as compared to scrambled peptides and PEG coated nanoparticles

[0549] In one embodiment, the NAVs of the present invention are formulated in nanoparticles which comprise a conjugate to enhance the delivery of the nanoparticles of the present invention in a subject. The conjugate may be the CD47 membrane or the conjugate may be derived from the CD47 membrane protein, such as the "self" peptide described previously. In another aspect the nanoparticle may comprise PEG and a conjugate of CD47 or a derivative thereof. In yet another aspect, the nanoparticle may comprise both the "self" peptide described above and the membrane protein CD47.

[0550] In another aspect, a "self" peptide and/or CD47 protein may be conjugated to a virus-like particle or pseudovirion, as described herein for delivery of the NAVs of the present invention.

[0551] In another embodiment, NAV pharmaceutical compositions comprising the polynucleotides of the present invention and a conjugate which may have a degradable linkage. Non-limiting examples of conjugates include an aromatic moiety comprising an ionizable hydrogen atom, a spacer moiety, and a water-soluble polymer. As a non-limiting example, pharmaceutical compositions comprising a conjugate with a degradable linkage and methods for delivering such pharmaceutical compositions are described in US Patent Publication No. US20130184443.

[0552] The nanoparticle formulations may be a carbohydrate nanoparticle comprising a carbohydrate carrier and a NAV. As a non-limiting example, the carbohydrate carrier may include, but is not limited to, an anhydride-modified phytoglycogen or glycogen-type material, phtoglycogen octenyl succinate, phytoglycogen beta-dextrin, anhydride-modified phytoglycogen beta-dextrin. (See e.g., International Publication No. WO2012109121).

[0553] Nanoparticle formulations of the present invention may be coated with a surfactant or polymer in order to improve the delivery of the particle. In one embodiment, the nanoparticle may be coated with a hydrophilic coating such as, but not limited to, PEG coatings and/or coatings that have a neutral surface charge. The hydrophilic coatings may help to deliver nanoparticles with larger payloads such as, but not limited to, NAVs within the central nervous system. As a non-limiting example nanoparticles comprising a hydrophilic coating and methods of making such nanoparticles are described in US Patent Publication No. US20130183244.

[0554] Lipid nanoparticle formulations may be improved by replacing the cationic lipid with a biodegradable cationic lipid which is known as a rapidly eliminated lipid nanoparticle (reLNP). Ionizable cationic lipids, such as, but not limited to, DLinDMA, DLin-KC2-DMA, and DLin-MC3-DMA, have been shown to accumulate in plasma and tissues over time and may be a potential source of toxicity. The rapid metabolism of the rapidly eliminated lipids can improve the tolerability and therapeutic index of the lipid nanoparticles by an order of magnitude from a 1 mg/kg dose to a 10 mg/kg dose in rat. Inclusion of an enzymatically degraded ester linkage can improve the degradation and metabolism profile of the cationic component, while still maintaining the activity of the reLNP formulation. The ester linkage can be internally located within the lipid chain or it may be terminally located at the terminal end of the lipid chain. The internal ester linkage may replace any carbon in the lipid chain.

[0555] In one embodiment, the internal ester linkage may be located on either side of the saturated carbon.

[0556] In one embodiment, an immune response may be elicited by delivering a lipid nanoparticle which may include a nanospecies, a polymer and an immunogen. (U.S. Publication No. 20120189700 and International Publication No.

WO2012099805). The polymer may encapsulate the nanospecies or partially encapsulate the nanospecies. The immunogen may be a recombinant protein, a modified RNA and/or a polynucleotide described herein. In one embodiment, the lipid nanoparticle may be formulated for use in a vaccine such as, but not limited to, against a pathogen.

[0557] Lipid nanoparticles may be engineered to alter the surface properties of particles so the lipid nanoparticles may penetrate the mucosal barrier Mucus is located on mucosal tissue such as, but not limted to, oral (e.g., the buccal and esophageal membranes and tonsil tissue), ophthalmic, gastrointestinal (e.g., stomach, small intestine, large intestine, colon, rectum), nasal, respiratory (e.g., nasal, pharyngeal, tracheal and bronchial membranes), genital (e.g., vaginal, cervical and urethral membranes). Nanoparticles larger than 10-200 nm which are preferred for higher drug encapsulation efficiency and the ability to provide the sustained delivery of a wide array of drugs have been thought to be too large to rapidly diffuse through mucosal barriers. Mucus is continuously secreted, shed, discarded or digested and recycled so most of the trapped particles may be removed from the mucosla tissue within seconds or within a few hours. Large polymeric nanoparticles (200nm -500nm in diameter) which have been coated densely with a low molecular weight polyethylene glycol (PEG) diffused through mucus only 4 to 6-fold lower than the same particles diffusing in water (Lai et al. PNAS 2007 104(5):1482-487; Lai et al. Adv Drug Deliv Rev. 2009 61(2): 158-171). The transport of nanoparticles may be determined using rates of permeation and/or fluorescent microscopy techniques including, but not limited to, fluorescence recovery after photobleaching (FRAP) and high resolution multiple particle tracking (MPT). As a non-limiting example, compositions which can penetrate a mucosal barrier may be made as described in U.S. Pat. No. 8,241,670 or International Patent Publication No. WO2013110028.

[0558] The lipid nanoparticle engineered to penetrate mucus may comprise a polymeric material (i.e. a polymeric core) and/or a polymer-vitamin conjugate and/or a tri-block co-polymer. The polymeric material may include, but is not limited to, polyamines, polyethers, polyamides, polyesters, polycarbamates, polyureas, polycarbonates, poly(styrenes), polyimides, polysulfones, polyurethanes, polyacetylenes, polyethylenes, polyethyeneimines, polyisocyanates, polyacrylates, polymethacrylates, polyacrylonitriles, and polyarylates. The polymeric material may be biodegradable and/or biocompatible Non-limiting examples of biocompatible polymers are described in International Patent Publication No. WO2013116804. The polymeric material may additionally be irradiated. As a non-limiting example, the polymeric material may be gamma irradiated (See e.g., International App. No. WO201282165). Non-limiting examples of specific polymers include poly(caprolactone) (PCL), ethylene vinyl acetate polymer (EVA), poly (lactic acid) (PLA), poly(L-lactic acid) (PLLA), poly(glycolic acid) (PGA), poly(lactic acid-co-glycolic acid) (PLGA), poly(L-lactic acid-co-glycolic acid) (PLLGA), poly(D,L-lactide) (PDLA), poly(L-lactide) (PLLA), poly(D,L-lactide-co-caprolactone), poly(D,L-lactide-co-caprolactone-co-glycolide), poly(D,L-lactide-co-PEO-co-D,L-lactide), poly(D,L-lactide-co-PPO-co-D,L-lactide), polyalkyl cyanoacralate, polyurethane, poly-L-lysine (PLL), hydroxypropyl methacrylate (HPMA), polyethyleneglycol, poly-L-glutamic acid, poly(hydroxy acids), polyanhydrides, polyorthoesters, poly(ester amides), polyamides, poly(ester ethers), polycarbonates, polyalkylenes such as polyethylene and polypropylene, polyalkylene glycols such as polyethylene glycol) (PEG), polyalkylene oxides (PEG), polyalkylene terephthalates such as polyethylene terephthalate), polyvinyl alcohols (PVA), polyvinyl ethers, polyvinyl esters such as poly(vinyl acetate), polyvinyl halides such as poly(vinyl chloride) (PVC), polyvinylpyrrolidone, polysiloxanes, polystyrene (PS), polyurethanes, derivatized celluloses such as alkyl celluloses, hydroxyalkyl celluloses, cellulose ethers, cellulose esters, nitro celluloses, hydroxypropylcellulose, carboxymethylcellulose, polymers of acrylic acids, such as poly(methyl(meth)acrylate) (PMMA), poly(ethyl(meth)acrylate), poly(butyl(meth)acrylate), poly(isobutyl(meth)acrylate), poly(hexyl(meth)acrylate), poly(isodecyl(meth)acrylate), poly(lauryl(meth)acrylate), poly(phenyl(meth)acrylate), poly(methyl acrylate), poly(isopropyl acrylate), poly(isobutyl acrylate), poly(octadecyl acrylate) and copolymers and mixtures thereof, polydioxanone and its copolymers, polyhydroxyalkanoates, polypropylene fumarate, polyoxymethylene, poloxamers, poly(ortho)esters, poly(butyric acid), poly(valeric acid), poly(lactide-co-caprolactone), PEG-PLGA-PEG and trimethylene carbonate, polyvinylpyrrolidone.The lipid nanoparticle may be coated or associated with a co-polymer such as, but not limited to, a block co-polymer (such as a branched polyether-polyamide block copolymer described in International Publication No. WO2013012476), and (poly(ethylene glycol))-(poly(propylene oxide))-(poly(ethylene glycol)) triblock copolymer (see e.g., US Publication 20120121718 and US Publication 20100003337 and U.S. Pat. No. 8,263,665). The co-polymer may be a polymer that is generally regarded as safe (GRAS) and the formation of the lipid nanoparticle may be in such a way that no new chemical entities are created For example, the lipid nanoparticle may comprise poloxamers coating PLGA nanoparticles without forming new chemical entities which are still able to rapidly penetrate human mucus (Yang et al. Angew. Chem. Int. Ed. 2011 50:2597-2600). A non-limiting scalable method to produce nanoparticles which can penetrate human mucus is described by Xu et al. (See e.g., J Control Release 2013,170(2):279-86).

[0559] The vitamin of the polymer-vitamin conjugate may be vitamin E. The vitamin portion of the conjugate may be substituted with other suitable components such as, but not limited to, vitamin A, vitamin E, other vitamins, cholesterol, a hydrophobic moiety, or a hydrophobic component of other surfactants (e.g., sterol chains, fatty acids, hydrocarbon chains and alkylene oxide chains).

[0560] The lipid nanoparticle engineered to penetrate mucus may include surface altering agents such as, but not limited to, polynucleotides, anionic proteins (e.g., bovine serum albumin), surfactants (e.g., cationic surfactants such as

for example dimethyldioctadecyl-ammonium bromide), sugars or sugar derivatives (e.g., cyclodextrin), nucleic acids, polymers (e.g., heparin, polyethylene glycol and poloxamer), mucolytic agents (e.g., N-acetylcysteine, mugwort, bromelain, papain, clerodendrum, acetylcysteine, bromhexine, carbocisteine, eprazinone, mesna, ambroxol, sobrerol, domiodol, letosteine, stepronin, tiopronin, gelsolin, thymosin β4 dornase alfa, neltenexine, erdosteine) and various DNases including rhDNase. The surface altering agent may be embedded or enmeshed in the particle's surface or disposed (e.g., by coating, adsorption, covalent linkage, or other process) on the surface of the lipid nanoparticle (see e.g., US Publication 20100215580 and US Publication 20080166414 and US20130164343).

[0561] In one embodiment, the mucus penetrating lipid nanoparticles may comprise at least one polynucleotide described herein. The polynucleotide may be encapsulated in the lipid nanoparticle and/or disposed on the surface of the paricle. The polynucleotide may be covalently coupled to the lipid nanoparticle. Formulations of mucus penetrating lipid nanoparticles may comprise a plurality of nanoparticles. Further, the formulations may contain particles which may interact with the mucus and alter the structural and/or adhesive properties of the surrounding mucus to decrease mucoadhesion which may increase the delivery of the mucus penetrating lipid nanoparticles to the mucosal tissue.

[0562] In another embodiment, the mucus penetrating lipid nanoparticles may be a hypotonic formulation comprising a mucosal penetration enhancing coating. The formulation may be hypotonice for the epithelium to which it is being delivered. Non-limiting examples of hypotonic formulations may be found in International Patent Publication No. WO2013110028.

[0563] In one embodiment, in order to enhance the delivery through the mucosal barrier the NAV formulation may comprise or be a hypotonic solution. Hypotonic solutions were found to increase the rate at which mucoinert particles such as, but not limited to, mucus-penetrating particles, were able to reach the vaginal epithelial surface (See e.g., Ensign et al. Biomaterials 2013 34(28):6922-9).

[0564] In one embodiment such formulations may also be constructed or compositions altered such that they passively or actively are directed to different cell types *in vivo,* including but not limited to hepatocytes, immune cells, tumor cells, endothelial cells, antigen presenting cells, and leukocytes (Akinc et al. Mol Ther. 2010 18:1357-1364; Song et al., Nat Biotechnol. 2005 23:709-717; Judge et al., J Clin Invest. 2009 119:661-673; Kaufmann et al., Microvasc Res 2010 80:286-293; Santel et al., Gene Ther 2006 13:1222-1234; Santel et al., Gene Ther 2006 13:1360-1370; Gutbier et al., Pulm Pharmacol. Ther. 2010 23:334-344; Basha et al., Mol. Ther. 2011 19:2186-2200; Fenske and Cullis, Expert Opin Drug Deliv. 2008 5:25-44; Peer et al., Science. 2008 319:627-630; Peer and Lieberman, Gene Ther. 2011 18:1127-1133). One example of passive targeting of formulations to liver cells includes the DLin-DMA, DLin-KC2-DMA and DLin-MC3-DMA-based lipid nanoparticle formulations which have been shown to bind to apolipoprotein E and promote binding and uptake of these formulations into hepatocytes in vivo (Akinc et al. Mol Ther. 2010 18:1357-1364Formulations can also be selectively targeted through expression of different ligands on their surface as exemplified by, but not limited by, folate, transferrin, N-acetylgalactosamine (GalNAc), and antibody targeted approaches (Kolhatkar et al., Curr Drug Discov Technol. 2011 8:197-206; Musacchio and Torchilin, Front Biosci. 2011 16:1388-1412; Yu et al., Mol Membr Biol. 2010 27:286-298; Patil et al., Crit Rev Ther Drug Carrier Syst. 2008 25:1-61; Benoit et al., Biomacromolecules. 2011 12:2708-2714; Zhao et al., Expert Opin Drug Deliv. 2008 5:309-319; Akinc et al., Mol Ther. 2010 18:1357-1364; Srinivasan et al., Methods Mol Biol. 2012 820:105-116; Ben-Arie et al., Methods Mol Biol. 2012757:497-507; Peer 2010 J Control Release. 20:63-68; Peer et al., Proc Natl Acad Sci USA. 2007 104:4095-4100; Kim et al., Methods Mol Biol. 2011 721:339-353; Subramanya et al., Mol Ther. 2010 18:2028-2037; Song et al., Nat Biotechnol. 2005 23:709-717; Peer et al., Science. 2008 319:627-630; Peer and Lieberman, Gene Ther. 2011 18:1127-1133).

[0565] In one embodiment, the NAVs of the present invention can be formulated for controlled release and/or targeted delivery. As used herein, "controlled release" refers to a pharmaceutical composition or compound release profile that conforms to a particular pattern of release to effect a therapeutic outcome. In one embodiment, the RNAVs may be encapsulated into a delivery agent described herein and/or known in the art for controlled release and/or targeted delivery. As used herein, the term "encapsulate" means to enclose, surround or encase. As it relates to the formulation of the compounds of the invention, encapsulation may be substantial, complete or partial. The term "substantially encapsulated" means that at least greater than 50, 60, 70, 80, 85, 90, 95, 96, 97, 98, 99, 99.9, 99.9 or greater than 99.999% of the pharmaceutical composition or compound of the invention may be enclosed, surrounded or encased within the delivery agent. "Partially encapsulation" means that less than 10, 10, 20, 30, 40 50 or less of the pharmaceutical composition or compound of the invention may be enclosed, surrounded or encased within the delivery agent. Advantageously, encapsulation may be determined by measuring the escape or the activity of the pharmaceutical composition or compound of the invention using fluorescence and/or electron micrograph. For example, at least 1, 5, 10, 20, 30, 40, 50, 60, 70, 80, 85, 90, 95, 96, 97, 98, 99, 99.9, 99.99 or greater than 99.99% of the pharmaceutical composition or compound of the invention are encapsulated in the delivery agent.

[0566] In one embodiment, the controlled release formulation may include, but is not limited to, tri-block co-polymers. As a non-limiting example, the formulation may include two different types of tri-block co-polymers (International Pub. No. WO2012131104 and WO2012131106).

[0567] In another embodiment, the NAVs may be encapsulated into a lipid nanoparticle or a rapidly eliminated lipid

nanoparticle and the lipid nanoparticles or a rapidly eliminated lipid nanoparticle may then be encapsulated into a polymer, hydrogel and/or surgical sealant described herein and/or known in the art. As a non-limiting example, the polymer, hydrogel or surgical sealant may be PLGA, ethylene vinyl acetate (EVAc), poloxamer, GELSITE® (Nanotherapeutics, Inc. Alachua, FL), HYLENEX® (Halozyme Therapeutics, San Diego CA), surgical sealants such as fibrinogen polymers (Ethicon Inc. Cornelia, GA), TISSELLO (Baxter International, Inc Deerfield, IL), PEG-based sealants, and COSEAL® (Baxter International, Inc Deerfield, IL).

[0568] In another embodiment, the lipid nanoparticle may be encapsulated into any polymer known in the art which may form a gel when injected into a subject. As another non-limiting example, the lipid nanoparticle may be encapsulated into a polymer matrix which may be biodegradable.

[0569] In one embodiment, the the NAV formulation for controlled release and/or targeted delivery may also include at least one controlled release coating. Controlled release coatings include, but are not limited to, OPADRY®, polyvinylpyrrolidone/vinyl acetate copolymer, polyvinylpyrrolidone, hydroxypropyl methylcellulose, hydroxypropyl cellulose, hydroxyethyl cellulose, EUDRAGIT RL®, EUDRAGIT RSO and cellulose derivatives such as ethylcellulose aqueous dispersions (AQUACOAT® and SURELEASE®).

[0570] In one embodiment, the NAV controlled release and/or targeted delivery formulation may comprise at least one degradable polyester which may contain polycationic side chains. Degradeable polyesters include, but are not limited to, poly(serine ester), poly(L-lactide-co-L-lysine), poly(4-hydroxy-L-proline ester), and combinations thereof. In another embodiment, the degradable polyesters may include a PEG conjugation to form a PEGylated polymer.

[0571] In one embodiment, the NAV controlled release and/or targeted delivery formulation comprising at least one polynucleotide may comprise at least one PEG and/or PEG related polymer derivatives as described in US Patent No. 8,404,222.

[0572] In another embodiment, the NAV controlled release delivery formulation comprising at least one polynucleotide may be the controlled release polymer system described in US20130130348.

[0573] In one embodiment, the the NAVs of the present invention may be encapsulated in a therapeutic nanoparticle, referred to herein as "therapeutic nanoparticle RNAVs." Therapeutic nanoparticles may be formulated by methods described herein and known in the art such as, but not limited to, International Pub Nos. WO2010005740, WO2010030763, WO2010005721, WO2010005723, WO2012054923, US Pub. Nos. US20110262491, US20100104645, US20100087337, US20100068285, US20110274759, US20100068286, US20120288541, US20130123351 and US20130230567 and US Pat No. 8,206,747, 8,293,276, 8,318,208 and 8,318,211. In another embodiment, therapeutic polymer nanoparticles may be identified by the methods described in US Pub No. US20120140790.

[0574] In one embodiment, the therapeutic nanoparticle NAV may be formulated for sustained release. As used herein, "sustained release" refers to a pharmaceutical composition or compound that conforms to a release rate over a specific period of time. The period of time may include, but is not limited to, hours, days, weeks, months and years. As a non-limiting example, the sustained release nanoparticle may comprise a polymer and a therapeutic agent such as, but not limited to, the the polynucleotides of the present invention (see International Pub No. 2010075072 and US Pub No. US20100216804, US20110217377 and US20120201859). In another non-limiting example, the sustained release formulation may comprise agents which permit persistent bioavailability such as, but not limited to, crystals, macromolecular gels and/or particulate suspensions (see US Patent Publication No US20130150295).

[0575] In one embodiment, the therapeutic nanoparticle NAVs may be formulated to be target specific. As a non-limiting example, the thereapeutic nanoparticles may include a corticosteroid (see International Pub. No. WO2011084518). As a non-limiting example, the therapeutic nanoparticles may be formulated in nanoparticles described in International Pub No. WO2008121949, WO2010005726, WO2010005725, WO2011084521 and US Pub No. US20100069426, US20120004293 and US20100104655.

[0576] In one embodiment, the nanoparticles of the present invention may comprise a polymeric matrix. As a non-limiting example, the nanoparticle may comprise two or more polymers such as, but not limited to, polyethylenes, polycarbonates, polyanhydrides, polyhydroxyacids, polypropylfumerates, polycaprolactones, polyamides, polyacetals, polyethers, polyesters, poly(orthoesters), polycyanoacrylates, polyvinyl alcohols, polyurethanes, polyphosphazenes, polyacrylates, polymethacrylates, polycyanoacrylates, polyureas, polystyrenes, polyamines, polylysine, polyethylene imine), poly(serine ester), poly(L-lactide-co-L-lysine), poly(4-hydroxy-L-proline ester) or combinations thereof.

[0577] In one embodiment, the therapeutic nanoparticle comprises a diblock copolymer. In one embodiment, the diblock copolymer may include PEG in combination with a polymer such as, but not limited to, polyethylenes, polycarbonates, polyanhydrides, polyhydroxyacids, polypropylfumerates, polycaprolactones, polyamides, polyacetals, polyethers, polyesters, poly(orthoesters), polycyanoacrylates, polyvinyl alcohols, polyurethanes, polyphosphazenes, polyacrylates, polymethacrylates, polycyanoacrylates, polyureas, polystyrenes, polyamines, polylysine, polyethylene imine), poly(serine ester), poly(L-lactide-co-L-lysine), poly(4-hydroxy-L-proline ester) or combinations thereof. In yet another embodiment, the diblock copolymer may be a high-X diblock copolymer such as those described in International Patent Publication No. WO2013120052.

[0578] As a non-limiting example the therapeutic nanoparticle comprises a PLGA-PEG block copolymer (see US Pub. No. US20120004293 and US Pat No. 8,236,330). In another non-limiting example, the therapeutic nanoparticle is a stealth nanoparticle comprising a diblock copolymer of PEG and PLA or PEG and PLGA (see US Pat No 8,246,968 and International Publication No. WO2012166923). In yet another non-limiting example, the therapeutic nanoparticle is a stealth nanoparticle or a target-specific stealth nanoparticle as described in US Patent Publication No. US20130172406.

[0579] In one embodiment, the therapeutic nanoparticle may comprise a multiblock copolymer (See e.g., U.S. Pat. No. 8,263,665 and 8,287,910 and US Patent Pub. No. US20130195987).

[0580] In yet another non-limiting example, the lipid nanoparticle comprises the block copolymer PEG-PLGA-PEG (see e.g., the thermosensitive hydrogel (PEG-PLGA-PEG) was used as a TGF-beta1 gene delivery vehicle in Lee et al. Thermosensitive Hydrogel as a Tgf-β1 Gene Delivery Vehicle Enhances Diabetic Wound Healing. Pharmaceutical Research, 2003 20(12): 1995-2000; as a controlled gene delivery system in Li et al. Controlled Gene Delivery System Based on Thermosensitive Biodegradable Hydrogel. Pharmaceutical Research 2003 20(6):884-888; and Chang et al., Non-ionic amphiphilic biodegradable PEG-PLGA-PEG copolymer enhances gene delivery efficiency in rat skeletal muscle. J Controlled Release. 2007 118:245-253). The NAVs of the present invention may be formulated in lipid nanoparticles comprising the PEG-PLGA-PEG block copolymer.

[0581] In one embodiment, the therapeutic nanoparticle may comprise a multiblock copolymer (See e.g., U.S. Pat. No. 8,263,665 and 8,287,910 and US Patent Pub. No. US20130195987).

[0582] In one embodiment, the block copolymers described herein may be included in a polyion complex comprising a non-polymeric micelle and the block copolymer. (See e.g., U.S. Pub. No. 20120076836).

[0583] In one embodiment, the therapeutic nanoparticle may comprise at least one acrylic polymer. Acrylic polymers include but are not limited to, acrylic acid, methacrylic acid, acrylic acid and methacrylic acid copolymers, methyl methacrylate copolymers, ethoxyethyl methacrylates, cyanoethyl methacrylate, amino alkyl methacrylate copolymer, poly(acrylic acid), poly(methacrylic acid), polycyanoacrylates and combinations thereof.

[0584] In one embodiment, the therapeutic nanoparticles may comprise at least one poly(vinyl ester) polymer. The poly(vinyl ester) polymer may be a copolymer such as a random copolymer. As a non-limiting example, the random copolymer may have a structure such as those described in International Application No. WO2013032829 or US Patent Publication No US20130121954. In one aspect, the poly(vinyl ester) polymers may be conjugated to the polynucleotides described herein. In another aspect, the poly(vinyl ester) polymer which may be used in the present invention may be those described in

[0585] In one embodiment, the therapeutic nanoparticle may comprise at least one diblock copolymer. The diblock copolymer may be, but it not limited to, a poly(lactic) acid-poly(ethylene)glycol copolymer (see e.g., International Patent Publication No. WO2013044219). As a non-limiting example, the therapeutic nanoparticle may be used to treat cancer (see International publication No. WO2013044219).

[0586] In one embodiment, the therapeutic nanoparticles may comprise at least one cationic polymer described herein and/or known in the art.

[0587] In one embodiment, the therapeutic nanoparticles may comprise at least one amine-containing polymer such as, but not limited to polylysine, polyethylene imine, poly(amidoamine) dendrimers, poly(beta-amino esters) (See e.g., U.S. Pat. No. 8,287,849) and combinations thereof.

[0588] In another embodiment, the nanoparticles described herein may comprise an amine cationic lipid such as those described in International Patent Application No. WO2013059496. In one aspect the cationic lipids may have an amino-amine or an amino-amide moiety.

[0589] In one embodiment, the therapeutic nanoparticles may comprise at least one degradable polyester which may contain polycationic side chains. Degradeable polyesters include, but are not limited to, poly(serine ester), poly(L-lactide-co-L-lysine), poly(4-hydroxy-L-proline ester), and combinations thereof. In another embodiment, the degradable polyesters may include a PEG conjugation to form a PEGylated polymer.

[0590] In another embodiment, the therapeutic nanoparticle may include a conjugation of at least one targeting ligand. The targeting ligand may be any ligand known in the art such as, but not limited to, a monoclonal antibody. (Kirpotin et al, Cancer Res. 2006 66:6732-6740).

[0591] In one embodiment, the therapeutic nanoparticle may be formulated in an aqueous solution which may be used to target cancer (see International Pub No. WO2011084513 and US Pub No. US20110294717).

[0592] In one embodiment, the therapeutic nanoparticle NAVs, e.g., therapeutic nanoparticles comprising at least one NAV may be formulated using the methods described by Podobinski et al in US Patent No. 8,404,799

[0593] In one embodiment, the nanoparticle may be optimized for oral administration. The nanoparticle may comprise at least one cationic biopolymer such as, but not limited to, chitosan or a derivative thereof. As a non-limiting example, the nanoparticle may be formulated by the methods described in U.S. Pub. No. 20120282343.

[0594] In some embodiments, LNPs comprise the lipid KL52 (an amino-lipid disclosed in U.S. Application Publication No. 2012/0295832). Activity and/or safety (as measured by examining one or more of ALTIAST, white blood cell count and cytokine induction) of LNP administration may be improved by incorporation of such lipids. LNPs comprising KL52

may be administered intravenously and/or in one or more doses. In some embodiments, administration of LNPs comprising KL52 results in equal or improved mRNA and/or protein expression as compared to LNPs comprising MC3.

[0595] In some embodiments, NAV may be delivered using smaller LNPs. Such particles may comprise a diameter from below 0.1 um up to 100 nm such as, but not limited to, less than 0.1 um, less than 1.0 um, less than 5 um, less than 10 um, less than 15 um, less than 20 um, less than 25 um, less than 30 um, less than 35 um, less than 40 um, less than 50 um, less than 55 um, less than 60 um, less than 65 um, less than 70 um, less than 75 um, less than 80 um, less than 85 um, less than 90 um, less than 95 um, less than 100 um, less than 125 um, less than 150 um, less than 175 um, less than 200 um, less than 225 um, less than 250 um, less than 275 um, less than 300 um, less than 325 um, less than 350 um, less than 375 um, less than 400 um, less than 425 um, less than 450 um, less than 475 um, less than 500 um, less than 525 um, less than 550 um, less than 575 um, less than 600 um, less than 625 um, less than 650 um, less than 675 um, less than 700 um, less than 725 um, less than 750 um, less than 775 um, less than 800 um, less than 825 um, less than 850 um, less than 875 um, less than 900 um, less than 925 um, less than 950 um, less than 975 um,

[0596] In another embodiment, NAVs may be delivered using smaller LNPs which may comprise a diameter from about 1 nm to about 100 nm, from about 1 nm to about 10 nm, about 1 nm to about 20 nm, from about 1 nm to about 30 nm, from about 1 nm to about 40 nm, from about 1 nm to about 50 nm, from about 1 nm to about 60 nm, from about 1 nm to about 70 nm, from about 1 nm to about 80 nm, from about 1 nm to about 90 nm, from about 5 nm to about from 100 nm, from about 5 nm to about 10 nm, about 5 nm to about 20 nm, from about 5 nm to about 30 nm, from about 5 nm to about 40 nm, from about 5 nm to about 50 nm, from about 5 nm to about 60 nm, from about 5 nm to about 70 nm, from about 5 nm to about 80 nm, from about 5 nm to about 90 nm, about 10 to about 50 nM, from about 20 to about 50 nm, from about 30 to about 50 nm, from about 40 to about 50 nm, from about 20 to about 60 nm, from about 30 to about 60 nm, from about 40 to about 60 nm, from about 20 to about 70 nm, from about 30 to about 70 nm, from about 40 to about 70 nm, from about 50 to about 70 nm, from about 60 to about 70 nm, from about 20 to about 80 nm, from about 30 to about 80 nm, from about 40 to about 80 nm, from about 50 to about 80 nm, from about 60 to about 80 nm, from about 20 to about 90 nm, from about 30 to about 90 nm, from about 40 to about 90 nm, from about 50 to about 90 nm, from about 60 to about 90 nm and/or from about 70 to about 90 nm.

[0597] In some embodiments, such LNPs are synthesized using methods comprising microfluidic mixers. Exemplary microfluidic mixers may include, but are not limited to a slit interdigitial micromixer including, but not limited to those manufactured by Microinnova (Allerheiligen bei Wildon, Austria) and/or a staggered herringbone micromixer (SHM) (Zhigaltsev, I.V. et al., Bottom-up design and synthesis of limit size lipid nanoparticle systems with aqueous and triglyceride cores using millisecond microfluidic mixing have been published (Langmuir. 2012. 28:3633-40; Belliveau, N.M. et al., Microfluidic synthesis of highly potent limit-size lipid nanoparticles for in vivo delivery of siRNA. Molecular Therapy-Nucleic Acids. 2012. 1:e37; Chen, D. et al., Rapid discovery of potent siRNA-containing lipid nanoparticles enabled by controlled microfluidic formulation. J Am Chem Soc. 2012.134(16):6948-51). In some instances, methods of LNP generation comprising SHM, further comprise the mixing of at least two input streams wherein mixing occurs by microstructure-induced chaotic advection (MICA). According to this method, fluid streams flow through channels present in a herringbone pattern causing rotational flow and folding the fluids around each other. This method may also comprise a surface for fluid mixing wherein the surface changes orientations during fluid cycling. Methods of generating LNPs using SHM include those disclosed in U.S. Application Publication Nos. 2004/0262223 and 2012/0276209.

[0598] In one embodiment, the NAV of the present invention may be formulated in lipid nanoparticles created using a micromixer such as, but not limited to, a Slit Interdigital Microstructured Mixer (SIMM-V2) or a Standard Slit Interdigital Micro Mixer (SSIMM) or Caterpillar (CPMM) or Impinging-jet (IJMM)from the Institut für Mikrotechnik Mainz GmbH, Mainz Germany).

[0599] In one embodiment, the NAVs of the present invention may be formulated in lipid nanoparticles created using microfluidic technology (see Whitesides, George M. The Origins and the Future of Microfluidics. Nature, 2006 442: 368-373; and Abraham et al. Chaotic Mixer for Microchannels. Science, 2002 295: 647-651). As a non-limiting example, controlled microfluidic formulation includes a passive method for mixing streams of steady pressure-driven flows in micro channels at a low Reynolds number (See e.g., Abraham et al. Chaotic Mixer for Microchannels. Science, 2002 295: 647-651).

[0600] In one embodiment, the NAVs of the present invention may be formulated in lipid nanoparticles created using a micromixer chip such as, but not limited to, those from Harvard Apparatus (Holliston, MA) or Dolomite Microfluidics (Royston, UK). A micromixer chip can be used for rapid mixing of two or more fluid streams with a split and recombine mechanism.

[0601] In one embodiment, the NAVs of the invention may be formulated for delivery using the drug encapsulating microspheres described in International Patent Publication No. WO2013063468 or U.S. Patent No. 8,440,614. The microspheres may comprise a compound of the formula (I), (II), (III), (IV), (V) or (VI) as described in International Patent Publication No. WO2013063468. In another aspect, the amino acid, peptide, polypeptide, lipids (APPL) are useful in delivering the NAVs of the invention to cells (see International Patent Publication No. WO2013063468).

[0602] In one embodiment, the NAVs of the invention may be formulated in lipid nanoparticles having a diameter from

about 10 to about 100 nm such as, but not limited to, about 10 to about 20 nm, about 10 to about 30 nm, about 10 to about 40 nm, about 10 to about 50 nm, about 10 to about 60 nm, about 10 to about 70 nm, about 10 to about 80 nm, about 10 to about 90 nm, about 20 to about 30 nm, about 20 to about 40 nm, about 20 to about 50 nm, about 20 to about 60 nm, about 20 to about 70 nm, about 20 to about 80 nm, about 20 to about 90 nm, about 20 to about 100 nm, about 30 to about 40 nm, about 30 to about 50 nm, about 30 to about 60 nm, about 30 to about 70 nm, about 30 to about 80 nm, about 30 to about 90 nm, about 30 to about 100 nm, about 40 to about 50 nm, about 40 to about 60 nm, about 40 to about 70 nm, about 40 to about 80 nm, about 40 to about 90 nm, about 40 to about 100 nm, about 50 to about 60 nm, about 50 to about 70 nm about 50 to about 80 nm, about 50 to about 90 nm, about 50 to about 100 nm, about 60 to about 70 nm, about 60 to about 80 nm, about 60 to about 90 nm, about 60 to about 100 nm, about 70 to about 80 nm, about 70 to about 90 nm, about 70 to about 100 nm, about 80 to about 90 nm, about 80 to about 100 nm and/or about 90 to about 100 nm.

[0603] In one embodiment, the lipid nanoparticles may have a diameter from about 10 to 500 nm.

[0604] In one embodiment, the lipid nanoparticle may have a diameter greater than 100 nm, greater than 150 nm, greater than 200 nm, greater than 250 nm, greater than 300 nm, greater than 350 nm, greater than 400 nm, greater than 450 nm, greater than 500 nm, greater than 550 nm, greater than 600 nm, greater than 650 nm, greater than 700 nm, greater than 750 nm, greater than 800 nm, greater than 850 nm, greater than 900 nm, greater than 950 nm or greater than 1000 nm.

[0605] The nanoparticles of the present invention may further include nutrients such as, but not limited to, those which deficiencies can lead to health hazards from anemia to neural tube defects (see e.g, the nanoparticles described in International Patent Publication No WO2013072929). As a non-limiting example, the nutrient may be iron in the form of ferrous, ferric salts or elemental iron, iodine, folic acid, vitamins or micronutrients.

Introduction Into Cells

[0606] A variety of methods are known in the art and suitable for introduction of nucleic acid into a cell, including viral and non-viral mediated techniques and any of these may be used to introduce the NAVs of the present disclosure. Examples of typical non-viral mediated techniques include, but are not limited to, electroporation, calcium phosphate mediated transfer, nucleofection, sonoporation, heat shock, magnetofection, liposome mediated transfer, microinjection, microprojectile mediated transfer (nanoparticles), cationic polymer mediated transfer (DEAE-dextran, polyethylenimine, polyethylene glycol (PEG) and the like) or cell fusion.

[0607] The technique of sonoporation, or cellular sonication, is the use of sound (e.g., ultrasonic frequencies) for modifying the permeability of the cell plasma membrane. Sonoporation methods are known to those in the art and are used to deliver nucleic acids *in vivo* (Yoon and Park, Expert Opin Drug Deliv. 2010 7:321-330; Postema and Gilja, Curr Pharm Biotechnol. 2007 8:355-361; Newman and Bettinger, Gene Ther. 2007 14:465-475). Sonoporation methods are known in the art and are also taught for example as it relates to bacteria in US Patent Publication 20100196983 and as it relates to other cell types in, for example, US Patent Publication 20100009424.

[0608] Electroporation techniques are also well known in the art and are used to deliver nucleic acids *in vivo* and clinically (Andre et al., Curr Gene Ther. 2010 10:267-280; Chiarella et al., Curr Gene Ther. 2010 10:281-286; Hojman, Curr Gene Ther. 2010 10:128-138). Electroporation devices are sold by many companies worldwide including, but not limited to BTX® Instruments (Holliston, MA) (e.g., the AgilePulse In Vivo System) and Inovio (Blue Bell, PA) (e.g., Inovio SP-5P intramuscular delivery device or the CELLECTRA® 3000 intradermal delivery device). In one instance, NAVs may be delivered by electroporation as described in Example 9.

Excipients

[0609] NAV pharmaceutical formulations may additionally comprise a pharmaceutically acceptable excipient, which, as used herein, includes, but are not limited to, any and all solvents, dispersion media, diluents, or other liquid vehicles, dispersion or suspension aids, surface active agents, isotonic agents, thickening or emulsifying agents, preservatives, solid binders, lubricants, flavoring agents, stabilizers, antioxidants, osmolality adjusting agents, pH adjusting agents and the like, as suited to the particular dosage form desired. Various excipients for formulating pharmaceutical compositions and techniques for preparing the composition are known in the art (see Remington: The Science and Practice of Pharmacy, 21st Edition, A. R. Gennaro (Lippincott, Williams & Wilkins, Baltimore, MD, 2006). The use of a conventional excipient medium may be contemplated within the scope of the present disclosure, except insofar as any conventional excipient medium is incompatible with a substance or its derivatives, such as by producing any undesirable biological effect or otherwise interacting in a deleterious manner with any other component(s) of the pharmaceutical composition, its use is contemplated to be within the scope of this disclosure.

[0610] In some instances, a pharmaceutically acceptable excipient may be at least 95%, at least 96%, at least 97%, at least 98%, at least 99%, or 100% pure. In some instances, an excipient is approved for use for humans and for

veterinary use. In some instances, an excipient may be approved by United States Food and Drug Administration. In some instances, an excipient may be of pharmaceutical grade. In some instances, an excipient may meet the standards of the United States Pharmacopoeia (USP), the European Pharmacopoeia (EP), the British Pharmacopoeia, and/or the International Pharmacopoeia.

[0611] Pharmaceutically acceptable excipients used in the manufacture of pharmaceutical compositions include, but are not limited to, inert diluents, dispersing and/or granulating agents, surface active agents and/or emulsifiers, disintegrating agents, binding agents, preservatives, buffering agents, lubricating agents, and/or oils. Such excipients may optionally be included in pharmaceutical compositions The composition may also include excipients such as cocoa buffer and suppository waxes, coloring agents, coating agents, sweetening, flavoring, and/or perfuming agents.

[0612] Exemplary diluents include, but are not limited to, calcium carbonate, sodium carbonate, calcium phosphate, dicalcium phosphate, calcium sulfate, calcium hydrogen phosphate, sodium phosphate lactose, sucrose, cellulose, microcrystalline cellulose, kaolin, mannitol, sorbitol, inositol, sodium chloride, dry starch, cornstarch, powdered sugar, etc., and/or combinations thereof.

[0613] Exemplary granulating and/or dispersing agents include, but are not limited to, potato starch, corn starch, tapioca starch, sodium starch glycolate, clays, alginic acid, guar gum, citrus pulp, agar, bentonite, cellulose and wood products, natural sponge, cation-exchange resins, calcium carbonate, silicates, sodium carbonate, cross-linked poly(vinyl-pyrrolidone) (crospovidone), sodium carboxymethyl starch (sodium starch glycolate), carboxymethyl cellulose, cross-linked sodium carboxymethyl cellulose (croscarmellose), methylcellulose, pregelatinized starch (starch 1500), microcrystalline starch, water insoluble starch, calcium carboxymethyl cellulose, magnesium aluminum silicate (VEEGUM®), sodium lauryl sulfate, quaternary ammonium compounds, etc., and/or combinations thereof.

[0614] Exemplary surface active agents and/or emulsifiers include, but are not limited to, natural emulsifiers (e.g. acacia, agar, alginic acid, sodium alginate, tragacanth, chondrux, cholesterol, xanthan, pectin, gelatin, egg yolk, casein, wool fat, cholesterol, wax, and lecithin), colloidal clays (e.g. bentonite [aluminum silicate] and VEEGUM® [magnesium aluminum silicate]), long chain amino acid derivatives, high molecular weight alcohols (e.g. stearyl alcohol, cetyl alcohol, oleyl alcohol, triacetin monostearate, ethylene glycol distearate, glyceryl monostearate, and propylene glycol monostearate, polyvinyl alcohol), carbomers (e.g. carboxy polymethylene, polyacrylic acid, acrylic acid polymer, and carboxyvinyl polymer), carrageenan, cellulosic derivatives (e.g. carboxymethylcellulose sodium, powdered cellulose, hydroxymethyl cellulose, hydroxypropyl cellulose, hydroxypropyl methylcellulose, methylcellulose), sorbitan fatty acid esters (e.g. polyoxyethylene sorbitan monolaurate [TWEEN®20], polyoxyethylene sorbitan [TWEEN®60], polyoxyethylene sorbitan monooleate [TWEEN®80], sorbitan monopalmitate [SPAN®40], sorbitan monostearate [SPAN®60], sorbitan tristearate [SPAN®65], glyceryl monooleate, sorbitan monooleate [SPAN®80]), polyoxyethylene esters (e.g. polyoxyethylene monostearate [MYRJ®45], polyoxyethylene hydrogenated castor oil, polyethoxylated castor oil, polyoxymethylene stearate, and SOLUTOL®), sucrose fatty acid esters, polyethylene glycol fatty acid esters (e.g. CREMOPHOR®), polyoxyethylene ethers, (e.g. polyoxyethylene lauryl ether [BRIJ®30]), poly(vinyl-pyrrolidone), diethylene glycol monolaurate, triethanolamine oleate, sodium oleate, potassium oleate, ethyl oleate, oleic acid, ethyl laurate, sodium lauryl sulfate, PLUORINC®F 68, POLOXAMER®188, cetrimonium bromide, cetylpyridinium chloride, benzalkonium chloride, docusate sodium, etc. and/or combinations thereof.

[0615] Exemplary binding agents include, but are not limited to, starch (e.g. cornstarch and starch paste); gelatin; sugars (e.g. sucrose, glucose, dextrose, dextrin, molasses, lactose, lactitol, mannitol); amino acids (e.g., glycine); natural and synthetic gums (e.g. acacia, sodium alginate, extract of Irish moss, panwar gum, ghatti gum, mucilage of isapol husks, carboxymethylcellulose, methylcellulose, ethylcellulose, hydroxyethylcellulose, hydroxypropyl cellulose, hydroxypropyl methylcellulose, microcrystalline cellulose, cellulose acetate, poly(vinyl-pyrrolidone), magnesium aluminum silicate (VEEGUM®), and larch arabogalactan); alginates; polyethylene oxide; polyethylene glycol; inorganic calcium salts; silicic acid; polymethacrylates; waxes, water, alcohol; etc.; and combinations thereof.

[0616] Exemplary preservatives may include, but are not limited to, antioxidants, chelating agents, antimicrobial preservatives, antifungal preservatives, alcohol preservatives, acidic preservatives, and/or other preservatives. Oxidation is a potential degradation pathway for mRNA, especially for liquid mRNA formulations. In order to prevent oxidation, antioxidants can be added to the formulation. Exemplary antioxidants include, but are not limited to, alpha tocopherol, ascorbic acid, acorbyl palmitate, benzyl alcohol, butylated hydroxyanisole, EDTA, m-cresol, methionine, butylated hydroxytoluene, monothioglycerol, potassium metabisulfite, propionic acid, propyl gallate, sodium ascorbate, sodium bisulfite, sodium metabisulfite, thioglycerol and/or sodium sulfite. Exemplary chelating agents include ethylenediaminetetraacetic acid (EDTA), citric acid monohydrate, disodium edetate, dipotassium edetate, edetic acid, fumaric acid, malic acid, phosphoric acid, sodium edetate, tartaric acid, and/or trisodium edetate. Exemplary antimicrobial preservatives include, but are not limited to, benzalkonium chloride, benzethonium chloride, benzyl alcohol, bronopol, cetrimide, cetylpyridinium chloride, chlorhexidine, chlorobutanol, chlorocresol, chloroxylenol, cresol, ethyl alcohol, glycerin, hexetidine, imidurea, phenol, phenoxyethanol, phenylethyl alcohol, phenylmercuric nitrate, propylene glycol, and/or thimerosal. Exemplary antifungal preservatives include, but are not limited to, butyl paraben, methyl paraben, ethyl paraben, propyl paraben, benzoic acid, hydroxybenzoic acid, potassium benzoate, potassium sorbate, sodium benzoate, sodium propi-

onate, and/or sorbic acid. Exemplary alcohol preservatives include, but are not limited to, ethanol, polyethylene glycol, phenol, phenolic compounds, bisphenol, chlorobutanol, hydroxybenzoate, and/or phenylethyl alcohol. Exemplary acidic preservatives include, but are not limited to, vitamin A, vitamin C, vitamin E, beta-carotene, citric acid, acetic acid, dehydroacetic acid, ascorbic acid, sorbic acid, and/or phytic acid. Other preservatives include, but are not limited to, tocopherol, tocopherol acetate, deteroxime mesylate, cetrimide, butylated hydroxyanisol (BHA), butylated hydroxytoluened (BHT), ethylenediamine, sodium lauryl sulfate (SLS), sodium lauryl ether sulfate (SLES), sodium bisulfite, sodium metabisulfite, potassium sulfite, potassium metabisulfite, GLYDANT PLUS®, PHENONIP®, methylparaben, GERMALL®115, GERMABEN®II, NEOLONE™, KATHON™, and/or EUXYL®.

[0617] In some embodiments, the pH of the NAV solutions are maintained between pH 5 and pH 8 to improve stability. Exemplary buffers to control pH may include, but are not limited to sodium phosphate, sodium citrate, sodium succinate, histidine (or histidine-HCl), sodium carbonate, and/or sodium malate. In another embodiment, the exemplary buffers listed above may be used with additional monovalent counterions (including, but not limited to potassium). Divalent cations may also be used as buffer counterions; however, these are not preferred due to complex formation and/or mRNA degradation.

[0618] Exemplary buffering agents may also include, but are not limited to, citrate buffer solutions, acetate buffer solutions, phosphate buffer solutions, ammonium chloride, calcium carbonate, calcium chloride, calcium citrate, calcium glubionate, calcium gluceptate, calcium gluconate, D-gluconic acid, calcium glycerophosphate, calcium lactate, propanoic acid, calcium levulinate, pentanoic acid, dibasic calcium phosphate, phosphoric acid, tribasic calcium phosphate, calcium hydroxide phosphate, potassium acetate, potassium chloride, potassium gluconate, potassium mixtures, dibasic potassium phosphate, monobasic potassium phosphate, potassium phosphate mixtures, sodium acetate, sodium bicarbonate, sodium chloride, sodium citrate, sodium lactate, dibasic sodium phosphate, monobasic sodium phosphate, sodium phosphate mixtures, tromethamine, magnesium hydroxide, aluminum hydroxide, alginic acid, pyrogen-free water, isotonic saline, Ringer's solution, ethyl alcohol, etc., and/or combinations thereof.

[0619] Exemplary lubricating agents include, but are not limited to, magnesium stearate, calcium stearate, stearic acid, silica, talc, malt, glyceryl behanate, hydrogenated vegetable oils, polyethylene glycol, sodium benzoate, sodium acetate, sodium chloride, leucine, magnesium lauryl sulfate, sodium lauryl sulfate, etc., and combinations thereof.

[0620] Exemplary oils include, but are not limited to, almond, apricot kernel, avocado, babassu, bergamot, black current seed, borage, cade, camomile, canola, caraway, carnauba, castor, cinnamon, cocoa buffer, coconut, cod liver, coffee, corn, cotton seed, emu, eucalyptus, evening primrose, fish, flaxseed, geraniol, gourd, grape seed, hazel nut, hyssop, isopropyl myristate, jojoba, kukui nut, lavandin, lavender, lemon, litsea cubeba, macademia nut, mallow, mango seed, meadowfoam seed, mink, nutmeg, olive, orange, orange roughy, palm, palm kernel, peach kernel, peanut, poppy seed, pumpkin seed, rapeseed, rice bran, rosemary, safflower, sandalwood, sasquana, savoury, sea buckthorn, sesame, shea buffer, silicone, soybean, sunflower, tea tree, thistle, tsubaki, vetiver, walnut, and wheat germ oils. Exemplary oils include, but are not limited to, butyl stearate, caprylic triglyceride, capric triglyceride, cyclomethicone, diethyl sebacate, dimethicone 360, isopropyl myristate, mineral oil, octyldodecanol, oleyl alcohol, silicone oil, and/or combinations thereof.

[0621] Excipients such as cocoa buffer and suppository waxes, coloring agents, coating agents, sweetening, flavoring, and/or perfuming agents can be present in the composition, according to the judgment of the formulator.

[0622] Exemplary additives include physiologically biocompatible buffers (e.g., trimethylamine hydrochloride), addition of chelants (such as, for example, DTPA or DTPA-bisamide) or calcium chelate complexes (as for example calcium DTPA, CaNaDTPA-bisamide), or, optionally, additions of calcium or sodium salts (for example, calcium chloride, calcium ascorbate, calcium gluconate or calcium lactate). In addition, antioxidants and suspending agents can be used.

Cryoprotectants

[0623] In some embodiments, NAV formulations may comprise cyroprotectants. As used herein, there term "cryoprotectant" refers to one or more agent that when combined with a given substance, helps to reduce or eliminate damage to that substance that occurs upon freezing. In some embodiments, cryoprotectants are combined with NAVs in order to stabilize them during freezing. Frozen storage of NAVs between -20°C and -80°C may be advantageous for long term (e.g. 36 months) stability of polynucleotide. In some embodiments, cryoprotectants are included in NAV formulations to stabilize polynucleotide through freeze/thaw cycles and under frozen storage conditions. Cryoprotectants of the present invention may include, but are not limited to sucrose, trehalose, lactose, glycerol, dextrose, raffinose and/or mannitol. Trehalose is listed by the Food and Drug Administration as being generally regarded as safe (GRAS) and is commonly used in commercial pharmaceutical formulations.

Bulking agents

[0624] In some embodiments, NAV formulations may comprise bulking agents. As used herein, ther term "bulking agent" refers to one or more agents included in formulations to impart a desired consistency to the formulation and/or

stabilization of formulation components. In some embodiments, bulking agents are included in lyophilized NAV formulations to yield a "pharmaceutically elegant" cake, stabilizing the lyophilized NAVs during long term (e.g. 36 month) storage. Bulking agents of the present invention may include, but are not limited to sucrose, trehalose, mannitol, glycine, lactose and/or raffinose. In some embodiments, combinations of cryoprotectants and bulking agents (for example, sucroselglycine or trehalose/mannitol) may be included to both stabilize NAVs during freezing and provide a bulking agent for lyophilization.

[0625] Non-limiting examples of formulations and methods for formulating the NAVs of the present invention are also provided in International Publication No WO2013090648 filed December 14, 2012.

Inactive Ingredients

[0626] In some embodiments, NAV formulations may comprise at least one excipient which is an inactive ingredient. As used herein, ther term "inactive ingredient" refers to one or more inactive agents included in formulations. In some embodiments, all, none or some of the inactive ingredients which may be used in the formulations of the present invention may be approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients and the routes of administration the inactive ingredients may be formulated in are described in Table 26. In Table 26, "AN" means anesthetic, "CNBLK" means cervical nerve block, "NBLK" means nerve block, "IV" means intravenous, "IM" means intramuscular and "SC" means subcutaneous

Table 26. Inactive Ingredients

Inactive Ingredient	Route of Administration
Alpha-Terpineol	Topical
Alpha-Tocopherol	Intravenous; Topical
Alpha-Tocopherol Acetate, Dl-	Topical
Alpha-Tocopherol, Dl-	Intravenous; Topical
1,2,6-Hexanetriol	Topical
1,2-Dimyristoyl-Sn-Glycero-3-(Phospho-S-(1-Glycerol))	Intravenous; Infusion (IV)
1,2-Dimyristoyl-Sn-Glycero-3-Phosphocholine	Intravenous; Infusion (IV)
1,2-Dioleoyl-Sn-Glycero-3-Phosphocholine	Epidural
1,2-Dipalmitoyl-Sn-Glycero-3-(Phospho-Rac-(1-Glycerol))	Epidural
1,2-Distearoyl-Sn-Glycero-3-(Phospho-Rac-(1-Glycerol))	Intravenous
1,2-Distearoyl-Sn-Glycero-3-Phosphocholine	Intravenous
1-O-Tolylbiguanide	Topical
2-Ethyl-1,6-Hexanediol	Topical
Acetic Acid	Infiltration; Auricular (Otic); Extracorporeal; Intramuscular; Intravenous; Subcutaneous; Intra-articualr; Intralesional; Intramuscular; Intrasynovial; Intratracheal; Intravenous; Irrigation; Infusion (IV); Nasal; Nerve block; Ophthalmic; Photopheresis; Soft Tissue; Submucosal; Topical
Acetic Acid, Glacial	Intravenous; Infusion (IV); Subcutaneous
Acetic Anhydride	Intravenous
Acetone	Implantation; Topical
Acetone Sodium Bisulfite	Intrathecal (AN,CNBLK); Infiltration (AN); Dental; Inhalation; Nerve Block
Acetylated Lanolin Alcohols	Topical
Acetylated Monoglycerides	Intravenous
Acetylcysteine	Inhalation
Acetyltryptophan, DL-	Intravenous
Acrylates Copolymer	Topical; Transdermal
Acrylic Acid-Isooctyl Acrylate Copolymer	Transdermal
Acrylic Adhesive 788	Transdermal

Activated Charcoal	Intramuscular; Intravenous; Irrigation; Infusion (IV)
Adcote 72A103	Transdermal
Adhesive Tape	Topical
Adipic Acid	Intramuscular; Vaginal
Aerotex Resin 3730	Transdermal
Alanine	Infusion (IV)
Albumin Aggregated	Intravenous
Albumin Colloidal	Intravenous
Albumin Human	Intravenous; Infusion (IV); Subcutaneous
Alcohol	Dental; Intramuscular; Intravenous; Subcutaneous; Inhalation; Intravascular; Infusion (IV); Ophthalmic; Rectal; Respiratory (Inhalation); Topical; Transdermal
Alcohol, Dehydrated	Dental; Extracorporeal; Intramuscular; Intravenous; Subcutaneous; Inhalation; Intracavitary; Intravascular; Intravesical; Nasal, Ophthalmic; Photopheresis, Rectal; Respiratory (Inhalation); Sublingual; Topical; Transdermal
Alcohol, Denatured	Denatal; Intravenous; Topical; Vaginal
Alcohol, Diluted	Intramuscular; Intravenous; Topical
Alfadex	Intracavitary
Alginic Acid	Ophthalmic
Alkyl Ammonium Sulfonic Acid Betaine	Topical
Alkyl Aryl Sodium Sulfonate	Topical
Allantoin	Topical; Vaginal
Allyl .Alpha.-Ionone	Nasal
Almond Oil	Topical
Aluminum Acetate	Auricular (Otic); Topical
Aluminum Chlorhydroxy Allantoinate	Topical
Aluminum Hydroxide	Topical
Aluminum Hydroxide - Sucrose, Hydrated	Topical
Aluminum Hydroxide Gel	Topical
Aluminum Hydroxide Gel F 500	Topical
Aluminum Hydroxide Gel F 5000	Topical
Aluminum Monostearate	Topical
Aluminum Oxide	Topical
Aluminum Polyester	Transdermal
Aluminum Silicate	Topical
Aluminum Starch Octenylsuccinate	Topical
Aluminum Stearate	Topical
Aluminum Subacetate	Rectal
Aluminum Sulfate Anhydrous	Auricular (Otic); Topical
Amerchol C	Topical
Amerchol-Cab	Ophthalmic; Topical
Aminomethylpropanol	Topical
Ammonia	Inhalation
Ammonia Solution	Topical
Ammonia Solution, Strong	Topical
Ammonium Acetate	Intramuscular; Intravenous; Infusion (IV)
Ammonium Hydroxide	Intravenous; Ophthalmic; Subcutaneous; Topical
Ammonium Lauryl Sulfate	Topical
Ammonium Nonoxynol-4 Sulfate	Topical
Ammonium Salt Of C-12-C-15 Linear Primary Alcohol Ethoxylate	Topical
Ammonium Sulfate	Intravenous

Ammonyx	Topical
Amphoteric-2	Topical
Amphoteric-9	Topical
Anethole	Dental
Anhydrous Citric Acid	Intravenous; Infusion (IV); Rectal; Topical
Anhydrous Dextrose	Intramuscular; Intravenous; Subcutaneous; Infusion (IV); Nasal; Spinal
Anhydrous Lactose	Intramuscular; Intravenous; Intracavitary; Intravenous; Infusion (IV); Vaginal
Anhydrous Trisodium Citrate	Intramuscular; Intravenous; Intra-arterial; Intra-articular; Intrabursal; Infusion (IV); Nasal; Ophthalmic; Soft Tissue; Topical
Aniseed Oil	Rectal
Anoxid Sbn	Topical
Antifoam	Topical
Antipyrine	Ophthalmic
Apaflurane	Respiratory (Inhalation)
Apricot Kernel Oil Peg-6 Esters	Topical; Vaginal
Aquaphor	Topical
Arginine	Intramuscular; Intravenous; Infusion (IV)
Arlacel	Topical
Ascorbic Acid	Infiltration (AN); Caudal Block; Epidural; Intramuscular; Intravenous; Inhalation; Infusion (IV); Nerve Block; Rectal; Subcutaneous; Topical
Ascorbyl Palmitate	Rectal; Topical
Aspartic Acid	Infusion (IV)
Balsam Peru	Rectal
Barium Sulfate	Intrauterine; Vaginal
Beeswax	Topical; Vaginal
Beeswax, Synthetic	Topical
Beheneth-10	Topical
Bentonite	Topical; Transdermal; Vaginal
Benzalkonium Chloride	Auricular (Otic); Inhalation; Intra-Articular; Intrabursal; Intradermal; Intralesional; Intramuscular; Intraocular; Nasal; Ophthalmic; Respiratory (Inhalation); Topical
Benzenesulfonic Acid	Intravenous; Infusion (IV)
Benzethonium Chloride	Auricular (Otic); Intramuscular; Intravenous; Infusion (IV); Nasal; Ophthalmic
Benzododecinium Bromide	Ophthalmic
Benzoic Acid	Intramuscular; Intravenous; Irrigation; Infusion (IV); Rectal; Topical; Vaginal
Benzyl Alcohol	Infiltration (AN); Auricular (Otic); Dental; Epidural; Extracorporeal; Interstitial; Intra-Arterial; Intra-Articular; Intrabursal; Intracavitary; Intradermal; Intralesional; Intramuscular; Intraperitoneal; Intrapleural; Intrasynovial; Intrathecal; Intratracheal; Intratumor; Intravenous; Infusion(IV); Nasal; Nerve Block; Rectal; Soft Tissue; Subconjunctival; Subcutaneous; Topical; Ureteral; Vaginal
Benzyl Benzoate	Intramuscular
Benzyl Chloride	Intravenous
Betadex	Topical
Bibapcitide	Intravenous
Bismuth Subgallate	Rectal

Boric Acid	Auricular (Otic); Intravenous; Ophthalmic; Topical
Brocrinat	Infusion (IV)
Butane	Topical
Butyl Alcohol	Topical
Butyl Ester Of Vinyl Methyl Ether/Maleic Anhydride Copolymer (125000 Mw)	Topical
Butyl Stearate	Topical
Butylated Hydroxyanisole	Intramuscular; Infusion (IV); Nasal; Rectal; Topical; Vaginal
Butylated Hydroxytoluene	Intramuscular; Intravenous; Infusion (IV); Nasal; Rectal; Topical; Transdermal; Vaginal
Butylene Glycol	Topical; Transdermal
Butylparaben	Intramuscular; Rectal; Topical
Butyric Acid	Transdermal
C20-40 Pareth-24	Topical
Caffeine	Nasal; Ophthalmic
Calcium	Intramuscular
Calcium Carbonate	Auricular (Otic); Respiratory (Inhalation)
Calcium Chloride	Infiltration (AN); Caudal Block; Epidural; Intramuscular; Intravenous; Intraocular; Intraperitoneal; Intravascular; Intravitreal; Nerve Block; Ophthalmic; Subctaneous; Topical
Calcium Gluceptate	Intravenous
Calcium Hydroxide	Intravenous; Subcutaneous; Topical
Calcium Lactate	Vaginal
Calcobutrol	Intravenous
Caldiamide Sodium	Intravenous
Caloxetate Trisodium	Intravenous
Calteridol Calcium	Intravenous
Canada Balsam	Topical
Caprylic/Capric Triglyceride	Topical; Transdermal
Caprylic/Capric/Stearic Triglyceride	Topical
Captan	Topical
Captisol	Intravenous
Caramel	Rectal; Topical
Carbomer 1342	Ophthalmic; Topical; Transdermal
Carbomer 1382	Topical
Carbomer 934	Rectal; Topical; Vaginal
Carbomer 934p	Ophthalmic; Rectal; Topical; Vaginal
Carbomer 940	Ophthalmic; Topical; Transdermal
Carbomer 941	Topical
Carbomer 980	Topical; Transdermal
Carbomer 981	Topical
Carbomer Homopolymer Type B (Allyl Pentaerythritol Crosslinked)	Ophthalmic; Topical
Carbomer Homopolymer Type C (Allyl Pentaerythritol Crosslinked)	Topical
Carbon Dioxide	Infiltration (AN); Intramuscular (IM); Infusion (IV); Inhalation; Intra-arterial; Intracardiac; Intrathecal; Intravascular; Intravenous
Carboxy Vinyl Copolymer	Topical
Carboxymethylcellulose	Intra-articular; Intrabursal; Intralesional; Intramuscular; Soft tissue; Topical

Carboxymethylcellulose Sodium	Dental; Intra-articular; Intrabursal; Intradermal; Intramuscular; Intrasynovial; Intratracheal; Nasal; Ophthalmic; Soft tissue; Subcutaneous; Topical
Carboxypolymethylene	Rectal; Topical
Carrageenan	Dental; Topical; Transdermal
Carrageenan Salt	Topical
Castor Oil	Intramuscular; Ophthalmic; Topical
Cedar Leaf Oil	Topical
Cellulose	Topical
Cellulose, Microcrystalline	Intra-articular; Intramuscular; Intravenous; Intravitreal; Nasal; Vaginal
Cerasynt-Se	Rectal; Topical
Ceresin	Topical
Ceteareth-12	Topical
Ceteareth-15	Topical
Ceteareth-30	Topical
Cetearyl Alcohol/Ceteareth-20	Topical
Cetearyl Ethylhexanoate	Topical
Ceteth-10	Topical
Ceteth-2	Topical
Ceteth-20	Topical; Vaginal
Ceteth-23	Topical
Cetostearyl Alcohol	Topical; Vaginal
Cetrimonium Chloride	Topical
Cetyl Alcohol	Auricular (Otic); Ophthalmic; Rectal; Topical; Vaginal
Cetyl Esters Wax	Topical; Vaginal
Cetyl Palmitate	Topical; Vaginal
Cetylpyridinium Chloride	Inhalation; Iontophoresis; Transdermal
Chlorobutanol	Infiltration (AN); Auricular (Otic); Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inhalation; Intravenous; Nasal; Nerve Block; Ophthalmic; Topical
Chlorobutanol Hemihydrate	Intramuscular; Intravenous
Chlorobutanol, Anhydrous	Intramuscular; Intravenous; Ophthalmic
Chlorocresol	Topical
Chloroxylenol	Auricular (Otic); Topical
Cholesterol	Epidural; Infiltration; Intravecous; Ophthalmic; Topical; Vaginal
Choleth	Vaginal
Choleth-24	Topical
Citrate	Intravenous
Citric Acid	Intratheca! (AN,CNBLK); Infiltration (AN); Auricular (Otic); Caudal Block; Epidural; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Infiltration; Inhalation; Intra-amniotic; Intra-arterial; Intra-articular; Intrabursal; Intracardiac; Intralesional; Iintrapleural; Intrasynovial; Intratheca!; Intravascular; Intravenous; Iontophoresis; Nasal; Nerve Block; Ophthalmic; Peridural; Soft tissue; Topical; Transdermal; Vaginal
Citric Acid Monohydrate	Infiltration (AN); Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intracardiac; Intraocular; Intravenous; Nasal; Nerve Block; Ophthalmic; Topical; Vaginal
Citric Acid, Hydrous	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intravenous

Cocamide Ether Sulfate	Topical
Cocamine Oxide	Topical
Coco Betaine	Topical
Coco Diethanolamide	Topical
Coco Monoethanolamide	Topical
Cocoa Butter	Rectal; Topical
Coco-Glycerides	Topical
Coconut Oil	Topical
Coconut Oil, Hydrogenated	Rectal
Coconut Oil/Palm Kernel Oil Glycerides, Hydrogenated	Rectal; Vaginal
Cocoyl Caprylocaprate	Topical
Cola Nitida Seed Extract	Rectal
Collagen	Topical
Coloring Suspension	Topical
Corn Oil	Intramuscular
Cottonseed Oil	Intramuscular
Cream Base	Topical
Creatine	Intra-articular; Intralesional; Intramuscular
Creatinine	Auricular (Otic); Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intra-articular; Intrabursal; Intradermal; Intralesional; Intrasynovial; Ophthalmic; Soft tissue; Topical
Cresol	Subcutaneous
Croscarmellose Sodium	Intramuscular
Crospovidone	Implantation; Intra-articluar; Intramuscular; Intrauterine; Topical; Transdermal; Vagiinal
Cupric Sulfate	Auricular (Otic)
Cupric Sulfate Anhydrous	Auricular (Otic)
Cyclomethicone	Topical
Cyclomethicone/Dimethicone Copolyol	Topical
Cysteine	Intramuscular (IM); Subcutaneous (SC); Intravenous; Infusion (IV)
Cysteine Hydrochloride	Intravenous; Infusion (IV)
Cysteine Hydrochloride Anhydrous	Intradiscal
Cysteine, Dl-	Intradiscal
D&C Red No. 28	Topical
D&C Red No. 33	Topical
D&C Red No. 36	Topical
D&C Red No. 39	Topical
D&C Yellow No. 10	Dental; Inhalation; Rectal; Topical
Dalfampridine	Intravenous
Daubert 1-5 Pestr (Matte) 164z	Transdermal
Decyl Methyl Sulfoxide	Topical
Dehydag Wax Sx	Topical
Dehydroacetic Acid	Topical
Dehymuls E	Topical
Denatonium Benzoate	Topical
Deoxycholic Acid	Infusion (IV)
Dextran	Intravenous
Dextran 40	Intravenous
Dextrin	Topical
Dextrose	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Interstitial; Intracavitary; Intraperitoneal; Intrapleural; Intraspinal; Intravenous; Nasal; Spinal

Dextrose Monohydrate	Intravenous
Dextrose Solution	Intravenous; Infusion (IV)
Diatrizoic Acid	Intra-arterial; Intra-articular; Intracardiac; Intradiscal; Intramuscular; Intrauterine; Intravascular; Intravenous; Infusion (IV); Periarticular; Subcutaneous; Ureteral; Urethral
Diazolidinyl Urea	Topical
Dichlorobenzyl Alcohol	Topical
Dichlorodifluoromethane	Inhalation; Intrapleural; Nasal; Rectal; Topical
Dichlorotetrafluoroethane	Inhalation; Nasal; Rectal; Topical
Diethanolamine	Infusion (IV); Ophthalmic; Topical
Diethyl Pyrocarbonate	Inflitration
Diethyl Sebacate	Topical
Diethylene Glycol Monoethyl Ether	Topical; Transdermal
Diethylhexyl Phthalate	Ophthalmic; Transdermal
Dihydroxyaluminum Aminoacetate	Topical
Diisopropanolamine	Topical
Diisopropyl Adipate	Topical
Diisopropyl Dilinoleate	Topical
Dimethicone 350	Topical
Dimethicone Copolyol	Topical; Transermal
Dimethicone Mdx4-4210	Transdermal
Dimethicone Medical Fluid 360	Dental; Intravenous; Topical; Transdermal
Dimethyl Isosorbide	Topical
Dimethyl Sulfoxide	Infusion (IV); Subcutanous; Topical
Dimethylaminoethyl Methacrylate - Butyl Methacrylate - Methyl Methacrylate Copolymer	Transdermal
Dimethyldioctadecylammonium Bentonite	Rectal
Dimethylsiloxane/Methylvinylsiloxane Copolymer	Implantation; Intrauterine
Dinoseb Ammonium Salt	Topical
Dipalmitoylphosphatidylglycerol, Dl-	Inflitration
Dipropylene Glycol	Transdermal
Disodium Cocoamphodiacetate	Topical
Disodium Laureth Sulfosuccinate	Topical
Disodium Lauryl Sulfosuccinate	Topical
Disodium Sulfosalicylate	Topical
Disofenin	Topical
Divinylbenzene Styrene Copolymer	Ophthalmic
Dmdm Hydantoin	Topical
Docosanol	Topical
Docusate Sodium	Intramuscular; Topical
Duro-Tak 280-2516	Transdermal
Duro-Tak 387-2516	Transdermal
Duro-Tak 80-1196	Transdermal
Duro-Tak 87-2070	Transdermal
Duro-Tak 87-2194	Transdermal
Duro-Tak 87-2287	Percutaneous; Transdermal
Duro-Tak 87-2296	Transdermal
Duro-Tak 87-2888	Transdermal
Duro-Tak 87-2979	Transdermal

Edetate Calcium Disodium	Infiltration (AN); Caudal Block; Epidural; Intramuscular (IM); Infusion (IV); Intra-articular; Intra-arterial; Intracardiac; Intradiscal; Intraperitoneal; Intrathecal; Intrauterine; Intravascular; Intravenous; Intravesical; Nerve Block; Periarticular; Rectal; Subcutaneous; Ureteral; Urethral
Edetate Disodium	Infiltration (AN), Auricular (Otic); Caudal Block; Epidural; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inhalation; Intra-arterial; Intra-articular; Intrabursal; Intracardiac; Intradermal; Intradiscal; Intralesional; Intrasynovial; Intrauterine; Intravascular; Intravenous; Iontophoresis; Nasal; Nerve Block; Ophthalmic; Rectal; Respiratory (Inhalation); Soft tissue; Topical; Transdermal; Ureteral; Urethral; Vaginal
Edetate Disodium Anhydrous	Intra-amniotic; Intramuscular; Intravenous; Infusion (IV); Ophthalmic
Edetate Sodium	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inhalation; Ophthalmic; Topical
Edetic Acid	Auricular (Otic); Rectal; Submucosal; Topical
Egg Phospholipids	Intravenous; Infusion (IV)
Entsufon	Topical
Entsufon Sodium	Topical
Epilactose	Rectal
Epitetracycline Hydrochloride	Topical
Essence Bouquet 9200	Topical
Ethanolamine Hydrochloride	Intravenous
Ethyl Acetate	Intramuscular; Topical; Transdermal
Ethyl Oleate	Transdermal
Ethylcelluloses	Topical; Transdermal; Vaginal
Ethylene Glycol	Topical
Ethylene Vinyl Acetate Copolymer	Implantation; Intrauerine; Ophthalmic; Periodontal; Subcutaneous; Transdermal
Ethylenediamine	Intravenous; Infusion (IV); Rectal; Topical
Ethylenediamine Dihydrochloride	Topical
Ethylene-Propylene Copolymer	Transdermal
Ethylene-Vinyl Acetate Copolymer (28% Vinyl Acetate)	Vaginal
Ethylene-Vinyl Acetate Copolymer (9% Vinylacetate)	Vaginal
Ethylhexyl Hydroxystearate	Topical
Ethylparaben	Topical
Eucalyptol	Dental
Exametazime	Intravenous
Fat, Edible	Rectal
Fat, Hard	Rectal
Fatty Acid Esters	Transdermal
Fatty Acid Pentaerythriol Ester	Topical
Fatty Acids	Topical
Fatty Alcohol Citrate	Topical
Fatty Alcohols	Vaginal
Fd&C Blue No. 1	Dental; Rectal; Topical
Fd&C Green No. 3	Dental; Rectal
Fd&C Red No. 4	Topical
Fd&C Red No. 40	Topical
Fd&C Yellow No. 10 (Delisted)	Topical
Fd&C Yellow No. 5	Topical; Vaginal

Fd&C Yellow No. 6	Inhalation; Rectal; Topical
Ferric Chloride	Intravenous
Ferric Oxide	Topical
Flavor 89-186	Dental
Flavor 89-259	Dental
Flavor Df-119	Dental
Flavor Df-1530	Dental
Flavor Enhancer	Dental
Flavor Fig 827118	Rectal
Flavor Raspberry Pfc-8407	Rectal
Flavor Rhodia Pharmaceutical No. Rf 451	Topical
Fluorochlorohydrocarbons	Inhalation
Formaldehyde	Topical
Formaldehyde Solution	Topical
Fractionated Coconut Oil	Topical
Fragrance 3949-5	Topical
Fragrance 520a	Topical
Fragrance 6.007	Topical
Fragrance 91-122	Topical
Fragrance 9128-Y	Topical
Fragrance 93498g	Topical
Fragrance Balsam Pine No. 5124	Topical
Fragrance Bouquet 10328	Topical
Fragrance Chemoderm 6401-B	Topical
Fragrance Chemoderm 6411	Topical
Fragrance Cream No. 73457	Topical
Fragrance Cs-28197	Topical
Fragrance Felton 066m	Topical
Fragrance Firmenich 47373	Topical
Fragrance Givaudan Ess 9090/1c	Topical
Fragrance H-6540	Topical
Fragrance Herbal 10396	Topical
Fragrance Nj-1085	Topical
Fragrance P O Fl-147	Topical
Fragrance Pa 52805	Topical
Fragrance Pera Derm D	Topical
Fragrance Rbd-9819	Topical
Fragrance Shaw Mudge U-7776	Topical
Fragrance Tf 044078	Topical
Fragrance Ungerer Honeysuckle K 2771	Topical
Fragrance Ungerer N5195	Topical
Fructose	Infusion (IV); Rectal
Gadolinium Oxide	Intravenous
Galactose	Rectal
Gamma Cyclodextrin	Intravenous
Gelatin	Dental; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intravenous; Respiratory (Inhalation); Topical; Vaginal
Gelatin, Crosslinked	Dental
Gelfoam Sponge	N/A
Gellan Gum (Low Acyl)	Ophthalmic
Gelva 737	Transdermal
Gentisic Acid	Intravenous
Gentisic Acid Ethanolamide	Infusion (IV)

Gluceptate Sodium	Intravenous
Gluceptate Sodium Dihydrate	Intravenous
Gluconolactone	Intramuscular (IM); Infusion (IV); Intravesou; Topical
Glucuronic Acid	Intravenous
Glutamic Acid, Dl-	Vaginal
Glutathione	Intramuscular
Glycerin	Auricular (Otic); Dental; Intramuscular; Infusion (IV); Subcutaneous (SC); Inhalation; Intradermal; Intravenous; Iontophoresis; Nasal; Ophthalmic; Perfusion; Biliary; Rectal; Topical; Transdermal; Vaginal
Glycerol Ester Of Hydrogenated Rosin	Nasal
Glyceryl Citrate	Topical
Glyceryl Isostearate	Topical; Vaginal
Glyceryl Laurate	Transdermal
Glyceryl Monostearate	Topical; Vaginal
Glyceryl Oleate	Topical; Transdermal
Glyceryl Oleate/Propylene Glycol	Topical
Glyceryl Palmitate	Rectal; Topical
Glyceryl Ricinoleate	Topical
Glyceryl Stearate	Auricular (Otic); Dental; Ophthalmic; Rectal; Topical; Vaginal
Glyceryl Stearate - Laureth-23	Topical
Glyceryl Stearate/Peg Stearate	Rectal
Glyceryl Stearate/Peg-100 Stearate	Topical
Glyceryl Stearate/Peg-40 Stearate	Rectal
Glyceryl Stearate-Stearamidoethyl Diethylamine	Topical
Glyceryl Trioleate	Epidural
Glycine	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intravenous; Rectal; Respiratory (Inhalation)
Glycine Hydrochloride	Subcutaneous
Glycol Distearate	Topical
Glycol Stearate	Topical
Guanidine Hydrochloride	Intravenous
Guar Gum	Topical; Vaginal
Hair Conditioner (18n195-1m)	Topical
Heptane	Transdermal
Hetastarch	Intravenous
Hexylene Glycol	Topical
High Density Polyethylene	Dental; Intrauterine; Ophthalmic; Topical; Transdermal; Vaginal
Histidine	Intravenous; Infusion (IV); Subcutaneous
Human Albumin Microspheres	Intravenous
Hyaluronate Sodium	Intra-articular; Intramuscular; Intravitreal; Topical
Hydrocarbon	Rectal
Hydrocarbon Gel, Plasticized	Dental; Ophthalmic; Topical

Hydrochloric Acid	Intrathecal (AN, CNBLK); Infiltration (AN); Sympathetic (AN, NBLK); Auricular (Otic); Caudal Block; Dental; Diagnostic; Epidural; Extracorporeal; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Infiltration; Inhalationi; Interstitial; Intra-amniotic; Intra-arterial; Intra-articular; Intrabursal; Intracardiac; Intracaudal; Intracavitary; Intradermal; Intralesional; Intraocular; Intraperitoneal; Intrapleural; Intraspinal; Intrasynovial; Intrathecal; Intratracheal; Intratumor; Intravascular; Intravenous; Intravesical; Intravitreal; Iontophoresis; Irrigation; Nasal; Nerve Block; Ophthalmic; Parenteral; Perfusion, Cardiac; Peridural; Perineural; Periodontal; Pectal; Respiratory (Inhalation); Retrobulbar; Soft tissue; Spinal; Subarachnoid; Subconjunctival; Subcutaneous; Topical; Transdermal; Ureteral; Urethral
Hydrochloric Acid, Diluted	Infiltration (AN); Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inhalation; Intra-arterial; Intravascular; Intravenous; Nerve Block; Ophthalmic; Topical
Hydrocortisone	Auricular (Otic)
Hydrogel Polymer	Vaginal
Hydrogen Peroxide	Topical
Hydrogenated Castor Oil	Topical
Hydrogenated Palm Oil	Rectal; Vaginal
Hydrogenated Palm/Palm Kernel Oil Peg-6 Esters	Topical
Hydrogenated Polybutene 635-690	Transdermal
Hydroxide Ion	Intramuscular; Infusion (IV)
Hydroxyethyl Cellulose	Auricular (Otic); Ophthalmic; Topical; Transdermal
Hydroxyethylpiperazine Ethane Sulfonic Acid	Intravenous
Hydroxymethyl Cellulose	Topical
Hydroxyoctacosanyl Hydroxystearate	Topical
Hydroxypropyl Cellulose	Topical
Hydroxypropyl Methylcellulose 2906	Ophthalmic
Hydroxypropyl-Bcyclodextrin	Intravenous; Infusion (IV)
Hypromellose 2208 (15000 Mpa.S)	Vaginal
Hypromellose 2910 (15000 Mpa.S)	Nasal; Ophthalmic
Hypromelloses	Irrigation; Ophthalmic; Rectal; Topical; Vaginal
Imidurea	Topical
Iodine	Intra-arterial; Intra-articular; Intracardiac; Intradiscal; Intravascular; Intravenous; Periarticular
Iodoxamic Acid	Intravenous
Iofetamine Hydrochloride	Intravenous
Irish Moss Extract	Topical
Isobutane	Topical
Isoceteth-20	Topical
Isoleucine	Infusion (IV)
Isooctyl Acrylate	Topical
Isopropyl Alcohol	Intravenous; Topical
Isopropyl Isostearate	Topical
Isopropyl Myristate	Auricular (Otic); Topical; Transdermal; Vaginal
Isopropyl Myristate - Myristyl Alcohol	Topical
Isopropyl Palmitate	Topical; Transdermal
Isopropyl Stearate	Topical
Isostearic Acid	Topical
Isostearyl Alcohol	Topical

Isotonic Sodium Chloride Solution	Epidural; Intratracheal; Intravenous; Infusion (IV)
Jelene	Ophthalmic; Topical
Kaolin	Topical
Kathon Cg	Topical
Kathon Cg II	Topical
Lactate	Topical
Lactic Acid	Infiltration (AN); Auricular (Otic); Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intracardiac; Intravenous; Nerve Block; Topical; Vaginal
Lactic Acid, Dl-	Intramuscular (IM); Infusion (IV); Intravesou; Topical; Vaginal
Lactic Acid, L-	Intravenous; Subcutanous
Lactobionic Acid	Intravenous; Infusion (IV)
Lactose	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inhalation; Intracavitary; Intravenous; Rectal; Transdermal; Vaginal
Lactose Monohydrate	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intracavitary; Intravenous; Respiratory (Inhalation); Vaginal
Lactose, Hydrous	Intramuscular (IM); Infusion (IV); Intravenous; Vaginal
Laneth	Topical
Lanolin	Ophthalmic; Rectal; Topical; Vaginal
Lanolin Alcohol - Mineral Oil	Topical
Lanolin Alcohols	Ophthalmic; Topical
Lanolin Anhydrous	Ophthalmic; Topical; Transdermal; Vaginal
Lanolin Cholesterols	Topical
Lanolin Nonionic Derivatives	Ophthalmic
Lanolin, Ethoxylated	Topical
Lanolin, Hydrogenated	Topical
Lauralkonium Chloride	Ophthalmic
Lauramine Oxide	Topical
Laurdimonium Hydrolyzed Animal Collagen	Topical
Laureth Sulfate	Topical
Laureth-2	Topical
Laureth-23	Topical
Laureth-4	Topical
Lauric Diethanolamide	Topical
Lauric Myristic Diethanolamide	Topical
Lauroyl Sarcosine	Ophthalmic
Lauryl Lactate	Transdermal
Lauryl Sulfate	Topical
Lavandula Angustifolia Flowering Top	Topical
Lecithin	Inhalation; Intramuscular; Rectal; Topical; Transdermal; Vaginal
Lecithin Unbleached	Topical
Lecithin, Egg	Intravenous
Lecithin, Hydrogenated	Auricular (Otic)
Lecithin, Hydrogenated Soy	Inhalation; Intravenous
Lecithin, Soybean	Inhalation; Vaginal
Lemon Oil	Topical
Leucine	Infusion (IV)
Levulinic Acid	Transdermal
Lidofenin	Intravenous
Light Mineral Oil	Ophthalmic; Rectal; Topical; Vaginal; Transdermal
Light Mineral Oil (85 Ssu)	Topical

Limonene, (+/-)-	Topical
Lipocol Sc-15	Topical
Lysine	Intramuscular (IM); Infusion (IV)
Lysine Acetate	Infusion (IV)
Lysine Monohydrate	Respiratory (Inhalation)
Magnesium Aluminum Silicate	Rectal; Topical; Vaginal
Magnesium Aluminum Silicate Hydrate	Rectal; Topical; Vaginal
Magnesium Chloride	Intramuscular; Intraocular; Intraperitoneal; Intravitreal; Infusion (IV); Ophthalmic; Subcutaneous
Magnesium Nitrate	Topical
Magnesium Stearate	Implantation; Intravitreal; Subcutaneous; Topical; Transmucosal; Vaginal
Maleic Acid	Intramuscular; Infusion (IV)
Mannitol	Intramuscular (IM); Infusion (IV); Subcutanous (SC); Intravenous; Ophthalmic; Parenteral; Respiratory (Inhalation); Submucosal; Topical; Transdermal
Maprofix	Topical
Mebrofenin	Intravenous
Medical Adhesive Modified S-15	Transdermal
Medical Antiform A-F Emulsion	Topical
Medronate Disodium	Intravenous
Medronic Acid	Intravenous
Meglumine	Intra-arterial; Intra-articular; Intracardiac; Intradiscal; Intramuscular; Intrauterine; Intravascular; Intravenous; Infusion (IV); Periarticular; Ureteral; Urethral
Menthol	Detanl; Inhalation; Topical
Metacresol	Intramuscular (IM); Infusion (IV); Subcutanous (SC); Intradermal
Metaphosphoric Acid	Infusion (IV)
Methanesulfonic Acid	Intramuscular (IM); Infusion (IV); Subcutaneous (SC)
Methionine	Intramuscular; Intrathecal; Intravenous; Infusion (IV); Subcutaneous
Methyl Alcohol	Transdermal
Methyl Gluceth-10	Topical
Methyl Gluceth-20	Topical
Methyl Gluceth-20 Sesquistearate	Topical
Methyl Glucose Sesquistearate	Topical
Methyl Laurate	Transdermal
Methyl Pyrrolidone	Periodontal; Subcutaneous
Methyl Salicylate	Topical
Methyl Stearate	Topical; Vaginal
Methylboronic Acid	Intravenous
Methylcellulose (4000 Mpa.S)	Ophthalmic
Methylcelluloses	Intra-articular; Intralesional; Intramuscular; Intrasynovial; Nasal; Ophthalmic; Soft tissue; Topical
Methylchloroisothiazolinone	Topical
Methylene Blue	Intravenous
Methylisothiazolinone	Topical

Methylparaben	Infiltration (AN); Auricular (Otic); Caudal Block; Epidural; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inhalation; Intra-arterial; Intra-articular; Intrabursal; Intradermal; Intralesional; Intrasynovial; Intravenous; Iontophoresis; Irrigation; Nasal; Nerve Block; Ophthalmic; Peridural; Rectal; Soft tissue; Topical; Ureteral; Urethral; Vaginal
Microcrystalline Wax	Topical; Vaginal
Mineral Oil	Auricular (Otic); Dental; Ophthalmic; Topical; Transdermal; Vaginal
Mono And Diglyceride	Topical
Monostearyl Citrate	Topical
Monothioglycerol	Infiltration (AN); Caudal Block; Epidural; Intramuscular (IM); Infusion (IV); Subcutanous (SC); Intravenous; Nerve Block
Multisterol Extract	Topical
Myristyl Alcohol	Topical
Myristyl Lactate	Topical
Myristyl-.Gamma.-Picolinium Chloride	Intra-articular; Intralesional; Intramuscular; Intrasynovial; Soft tissue
N-(Carbamoyl-Methoxy Peg-40)-1,2-Distearoyl-Cephalin Sodium	Intravenous
N,N-Dimethylacetamide	Intramuscular; Intravenous; Infusion (IV)
Niacinamide	Intramuscular; Infusion (IV); Intra-articular; Intralesional; Intrasynovial; Topical
Nioxime	Intravenous
Nitric Acid	Inhalation; Infusion (IV); Ophthalmic; Topical; Vaginal
Nitrogen	Infiltration (AN); Caudal Block; Dental; Epidural; Intramuscular; Infusion (IV); Subcutanous (SC); Inhalation; Intra-arterial; Intracavitary; Intramuscular (IM); Intrathecal; Intratumor; Intravascular; Intravenous; Intravesical; Irrigation; Nasal; Nerve Block; Ophthalmic; Parenteral; Submucosal; Topical; Transdermal
Nonoxynol Iodine	Topical
Nonoxynol-15	Topical
Nonoxynol-9	Ophthalmic; Topical
Norflurane	Inhalation; Nasal; Respiratory (Inhalation)
Oatmeal	Topical
Octadecene-1/Maleic Acid Copolymer	Topical
Octanoic Acid	Intravenous
Octisalate	Transdermal
Octoxynol-1	Topical
Octoxynol-40	Ophthalmic
Octoxynol-9	Topical
Octyldodecanol	Topical; Transdermal; Vaginal
Octylphenol Polymethylene	Ophthalmic
Oleic Acid	Inhalation; Nasal; Respiratory (Inhalation); Topical; Transdermal
Oleth-10/Oleth-5	Topical
Oleth-2	Topical
Oleth-20	Topical
Oleyl Alcohol	Topical; Transdermal
Oleyl Oleate	Topical; Transdermal
Olive Oil	Topical

Oxidronate Disodium	Intravenous
Oxyquinoline	Intravenous
Palm Kernel Oil	Rectal
Palmitamine Oxide	Topical
Parabens	Topical
Paraffin	Rectal; Topical
Paraffin, White Soft	Topical
Parfum Creme 45/3	Topical
Peanut Oil	Intramuscular; Intratracheal; Topical; Vaginal
Peanut Oil, Refined	Topical
Pectin	Dental; Topical
Peg 6-32 Stearate/Glycol Stearate	Topical; Vaginal
Peg Vegetable Oil	Intramuscular (IM); Infusion (IV); Subcutaneous (SC)
Peg-100 Stearate	Topical; Vaginal
Peg-12 Glyceryl Laurate	Topical
Peg-120 Glyceryl Stearate	Topical; Vaginal
Peg-120 Methyl Glucose Dioleate	Topical
Peg-15 Cocamine	Topical
Peg-150 Distearate	Topical
Peg-2 Stearate	Topical; Vaginal
Peg-20 Sorbitan Isostearate	Intramuscular
Peg-22 Methyl Ether/Dodecyl Glycol Copolymer	Topical
Peg-25 Propylene Glycol Stearate	Topical
Peg-4 Dilaurate	Topical
Peg-4 Laurate	Topical
Peg-40 Castor Oil	Intramuscular (IM); Subcutaneous (SC); Infusion (IV)
Peg-40 Sorbitan Diisostearate	Dental
Peg-45/Dodecyl Glycol Copolymer	Topical
Peg-5 Oleate	Topical; Vaginal
Peg-50 Stearate	Topical
Peg-54 Hydrogenated Castor Oil	Topical
Peg-6 Isostearate	Topical
Peg-60 Castor Oil	Infusion (IV)
Peg-60 Hydrogenated Castor Oil	Topical
Peg-7 Methyl Ether	Topical
Peg-75 Lanolin	Topical
Peg-8 Laurate	Topical
Peg-8 Stearate	Topical
Pegoxol 7 Stearate	Topical; Vaginal
Pentadecalactone	Transdermal
Pentaerythritol Cocoate	Topical
Pentasodium Pentetate	Intravenous
Pentetate Calcium Trisodium	Intrathecal; Intravenous; Infusion (IV)
Pentetic Acid	Intrathecal; Intravenous
Peppermint Oil	Dental; Topical
Perflutren	Intravenous
Perfume 25677	Topical
Perfume Bouquet	Topical
Perfume E-1991	Topical
Perfume Gd 5604	Topical
Perfume Tana 90/42 Scba	Topical
Perfume W-1952-1	Topical
Petrolatum	Auricular (Otic); Ophthalmic; Topical

Petrolatum, White	Auricular (Otic); Dental; Nasal; Ophthalmic; Rectal; Topical; Transdermal; Vaginal
Petroleum Distillates	Topical
Phenol	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intra-articular; Intradermal; Intralesional; Intrasynovial; Intravenous; Soft tissue
Phenol, Liquefied	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intravenous
Phenonip	Iontophoresis; Topical
Phenoxyethanol	Topical
Phenylalanine	Infusion (IV)
Phenylethyl Alcohol	Auricular (Otic); Nasal; Ophthalmic
Phenylmercuric Acetate	Ophthalmic; Topical; Vaginal
Phenylmercuric Nitrate	Intramuscular; Ophthalmic
Phosphatidyl Glycerol, Egg	Intravenous
Phospholipid	Infusion (IV)
Phospholipid, Egg	Intravenous; Infusion (IV)
Phospholipon 90g	Vaginal
Phosphoric Acid	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Infiltration; Intra-articular; Intralesional; Intravenous; Ophthalmic; Soft tissue; Topical; Vaginal
Pine Needle Oil (Pinus Sylvestris)	Topical
Piperazine Hexahydrate	Vaginal
Plastibase-50w	Dental; Topical
Polacrilin	Iontophoresis; Transdermal
Polidronium Chloride	Ophthalmic; Topical
Poloxamer 124	Topical
Poloxamer 181	Topical
Poloxamer 182	Topical
Poloxamer 188	Intravenous; Ophthalmic; Peridontal; Subcutaneous; Topical
Poloxamer 237	Topical
Poloxamer 407	Ophthalmic; Peridontal; Topical
Poly(Bis(P-Carboxyphenoxy)Propane Anhydride):Sebacic Acid	Implantation
Poly(Dimethylsiloxane/Methylvinylsiloxane/Methylhydrogen siloxane) Dimethylvinyl Or Dimethylhydroxy Or Trimethyl Endblocked	Vaginal
Poly(Dl-Lactic-Co-Glycolic Acid), (50:50	N/A
Poly(Dl-Lactic-Co-Glycolic Acid), Ethyl Ester Terminated, (50:50	N/A
Polyacrylic Acid (250000 Mw)	Transdermal
Polybutene (1400 Mw)	Transdermal
Polycarbophil	Ophthalmic; Topical; Vaginal
Polyester	Transdermal; Vaginal
Polyester Polyamine Copolymer	Transdermal
Polyester Rayon	Transdermal
Polyethylene Glycol 1000	Rectal; Respiratory (Inhalation); Topical; Vaginal
Polyethylene Glycol 1450	Topical; Urethral
Polyethylene Glycol 1500	Topical
Polyethylene Glycol 1540	Dental; Rectal; Topical
Polyethylene Glycol 200	Intramuscular; Topical
Polyethylene Glycol 300	Intramuscular (IM); Infusion (IV); Intravenous; Ophthalmic; Topical

Polyethylene Glycol 300-1600	Topical
Polyethylene Glycol 3350	Intra-articular; Intralesional; Intramuscular; Intrasynovial; Nasal; Rectal; Soft tissue; Subcutaneous; Topical; Vaginal
Polyethylene Glycol 400	Intramuscular (IM); Infusion (IV); Intravenous; Nasal; Ophthalmic; Rectal; Topical; Vaginal
Polyethylene Glycol 4000	Intra-articular; Intralesional; Intramuscular; Intrasynovial; Rectal; Soft tissue; Topical; Vaginal
Polyethylene Glycol 540	Topical
Polyethylene Glycol 600	Intravenous; Topical
Polyethylene Glycol 6000	Rectal; Topical; Vaginal
Polyethylene Glycol 8000	Ophthalmic; Rectal; Topical; Vaginal
Polyethylene Glycol 900	Topical
Polyethylene High Density Containing Ferric Oxide Black (<1%)	Intrauterine
Polyethylene Low Density Containing Barium Sulfate (20-24%)	Initrauterine
Polyethylene T	Initrauterine
Polyethylene Terephthalates	Transdermal
Polyglactin	Dental; Implantation; Intramuscular; Subcutaneous
Polyglyceryl-3 Oleate	Vaginl
Polyglyceryl-4 Oleate	Vaginl
Polyhydroxyethyl Methacrylate	Topical
Polyisobutylene	Topical; Transdermal
Polyisobutylene (1100000 Mw)	Topical; Transdermal
Polyisobutylene (35000 Mw)	Transdermal
Polyisobutylene 178-236	Transdermal
Polyisobutylene 241-294	Transdermal
Polyisobutylene 35-39	Transdermal
Polyisobutylene Low Molecular Weight	Transdermal
Polyisobutylene Medium Molecular Weight	Transdermal
Polyisobutylene/Polybutene Adhesive	Transdermal
Polylactide	Intramuscular; Peridontal
Polyols	Dental
Polyoxyethylene - Polyoxypropylene 1800	Ophthalmic; Topical
Polyoxyethylene Alcohols	Topical
Polyoxyethylene Fatty Acid Esters	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Topical
Polyoxyethylene Propylene	Topical
Polyoxyl 20 Cetostearyl Ether	Topical
Polyoxyl 35 Castor Oil	Intravesical; Infusion (IV); Ophthalmic
Polyoxyl 40 Hydrogenated Castor Oil	Dental; Ophthalmic; Topical
Polyoxyl 40 Stearate	Auricular (Otic); Dental; Ophthalmic; Topical
Polyoxyl 400 Stearate	Nasal; Topical
Polyoxyl 6 And Polyoxyl 32 Palmitostearate	Topical
Polyoxyl Distearate	Topical
Polyoxyl Glyceryl Stearate	Topical
Polyoxyl Lanolin	Topical
Polyoxyl Palmitate	Vaginl
Polyoxyl Stearate	Auricular (Otic); Topical
Polypropylene	Intrauterine; Topical; Transdermal
Polypropylene Glycol	Intramuscular (IM); Infusion (IV); Ophthalmic
Polyquaternium-10	Topical
Polyquaternium-7 (70/30 Acrylamide/Dadmac	N/A
Polysiloxane	Intravenous

Polysorbate 20	Auricular (Otic); Intramuscular (IM); Subcutaneous (SC); Intravenous; Infusion (IV); Nasal; Ophthalmic; Topical; Vaginal
Polysorbate 40	Intramuscular (IM); Infusion (IV); Topical
Polysorbate 60	Ophthalmic; Rectal; Topical; Vaginal
Polysorbate 65	Topical
Polysorbate 80	Auricular (Otic); Intra-articular; Intrabursal; Intradermal; Intralesional; Intramuscular; Intrasynovial; Intravenous; Infusion (IV); Nasal; Ophthalmic; Rectal; Soft tissue; Subcutaneous; Topical; Vaginal
Polyurethane	Vagianl
Polyvinyl Acetate	Transdermal
Polyvinyl Alcohol	Auricular (Otic); Intramuscular; Intraocular; Intravitreal; Iontophoresis; Ophthalmic; Topical; Transdermal
Polyvinyl Chloride	Transdermal
Polyvinyl Chloride-Polyvinyl Acetate Copolymer	Transdermal
Polyvinylpyridine	Transdermal
Poppy Seed Oil	Intralymphatic; Intrauterine
Potash	Topical
Potassium Acetate	Ophthalmic; Rectal
Potassium Alum	Vagianl
Potassium Bicarbonate	Transmucosal
Potassium Bisulfite	Intravenous
Potassium Chloride	Infiltration (AN); Caudal Block; Epidural; Intraocular; Intravenous; Intravitreal; Infusion (IV); Nerve Block; Ophthalmic
Potassium Citrate	Topical
Potassium Hydroxide	Intravascular; Intravenous; Infusion (IV); Topical; Vaginal
Potassium Metabisulfite	Infiltration (AN); Auricular (Otic); Intramuscular (IM); Infusion (IV); Nerve Block; Rectal
Potassium Phosphate, Dibasic	Intra-articular; Intramuscular; Intravenous; Infusion (IV); Subcutaneous
Potassium Phosphate, Monobasic	Infiltration (AN); Auricular (Otic); Intramuscular (IM); Infusion (IV); Intra-articular; Intramuterine; Intravenous; Intravesical; Nasal; Nerve Block; Ophthalmic; Subcutaneous
Potassium Soap	Topical
Potassium Sorbate	Nasal; Ophthalmic; Topical
Povidone Acrylate Copolymer	Topical
Povidone Hydrogel	Iontophoresis; Topical
Povidone K17	Subcutaneous
Povidone K25	Respiratory (Inhalation)
Povidone K29/32	Ophthalmic; Transdermal; Vaginal
Povidone K30	Ophthalmic
Povidone K90	Ophthalmic; Topical
Povidone K90f	Auricular (Otic)
Povidone/Eicosene Copolymer	Topical
Povidones	Auricular (Otic); Intramuscular; Intravenous; Infusion (IV); Ophthalmic; Subcutaneous; Topical; Transdermal; Vaginal
Ppg-12/Smdi Copolymer	Topical
Ppg-15 Stearyl Ether	Topical
Ppg-20 Methyl Glucose Ether Distearate	Topical
Ppg-26 Oleate	Topical
Product Wat	Topical

Proline	Infusion (IV)
Promulgen D	Topical; Vaginal
Promulgen G	Topical
Propane	Topical
Propellant A-46	Topical
Propyl Gallate	Topical; Intramuscular
Propylene Carbonate	Topical
Propylene Glycol	Auricular (Otic); Dental; Extracorporeal; Intramuscular (IM); Infusion (IV); Inhalation; Intravenous; Nasal; Ophthalmic; Photopheresis; Rectal; Subcutaneous; Topical; Transdermal; Vaginal
Propylene Glycol Diacetate	Auricular (Otic); Topical
Propylene Glycol Dicaprylate	Topical
Propylene Glycol Monolaurate	Transdermal
Propylene Glycol Monopalmitostearate	Topical; Vaginal
Propylene Glycol Palmitostearate	Topical
Propylene Glycol Ricinoleate	Topical
Propylene Glycol/Diazolidinyl Urea/Methylparaben/Propylparben	Topical
Propylparaben	Infiltration (AN); Auricular (Otic); Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inhalation; Intra-arterial; Intra-articular; Intrabursal; Intralesional; Intrasynovial; Intravenous; Nasal; Nerve Block; Ophthalmic; Rectal; Soft tissue; Topical; Ureteral; Urethral; Vaginal
Protamine Sulfate	Intramuscular (IM); Subcutaneous (SC); Intradermal
Protein Hydrolysate	Topical
Pvm/Ma Copolymer	Dental
Quaternium-15	Topical
Quaternium-15 Cis-Form	Topical; Vaginal
Quaternium-52	Topical
Ra-2397	Transdermal
Ra-3011	Transdermal
Saccharin	Inhalation; Topical
Saccharin Sodium	Dental; Intramuscular (IM); Infusion (IV); Inhalation; Intravenous; Rectal; Topical
Saccharin Sodium Anhydrous	Intramuscular (IM); Infusion (IV); Rectal
Safflower Oil	Topical
Sd Alcohol 3a	Topical
Sd Alcohol 40	Topical
Sd Alcohol 40-2	Topical
Sd Alcohol 40b	Topical
Sepineo P 600	Topical
Serine	Infusion (IV)
Sesame Oil	Intramuscular (IM); Subcutaneous (SC)
Shea Butter	Topical
Silastic Brand Medical Grade Tubing	Implantation
Silastic Medical Adhesive,Silicone Type A	Implantation
Silica, Dental	Dental
Silicon	Topical; Transdermal
Silicon Dioxide	Dental; Topical; Vaginal
Silicon Dioxide, Colloidal	Endocervical; Rectal; Respiratory (Inhalation); Transdermal; Vaginal

Silicone	Intramuscular (IM); Infusion (IV); Intrauterine; Topical; Transdermal; Vaginal
Silicone Adhesive 4102	Percutaneous; Transdermal
Silicone Adhesive 4502	Transdermal
Silicone Adhesive Bio-Psa Q7-4201	Transdermal; Topical
Silicone Adhesive Bio-Psa Q7-4301	Transdermal; Topical
Silicone Emulsion	Topical
Silicone/Polyester Film Strip	Transdermal
Simethicone	Intramuscular (IM); Infusion (IV); Rectal; Topical
Simethicone Emulsion	Topical
Sipon Ls 20np	Topical
Soda Ash	Ophthalmic
Sodium Acetate	Auricular (Otic); Extracorporeal; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Interstitial; Intra-articular; Intracavitary; Intradermal; Intralesional; Intraocular; Intraperitoneal; Intrapleural; Intrasynovial; Intravenous; Intravitreal; Nasal; Ophthalmic; Parenteral; Phtotpheresis; Soft tissue; Submucosal; Topical
Sodium Acetate Anhydrous	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intravenous; Topical
Sodium Alkyl Sulfate	Topical
Sodium Ascorbate	Intravenous
Sodium Benzoate	Dental; Intramuscular (IM); Infusion (IV); Intravenous; Rectal; Topical
Sodium Bicarbonate	Intramuscular (IM); Infusion (IV); Intraperitoneal; Intrathecal; Intratracheal; Intravenous; Intravitreal; Subcutaneous; Vaginal
Sodium Bisulfate	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inhalation; Ophthalmic
Sodium Bisulfite	Infiltration (AN); Auricular (Otic); Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Epidural; Inhalation; Intra-arterial; Intra-articular; Intrabursal; Intracardiac; Intradermal; Intradiscal; Intralesional; Intraperitoneal; Intrasynovial; Iontophoresis; Irrigation; Intravenous; Nerve Block; Ophthalmic; soft tissue; Topical
Sodium Borate	Auricular (Otic); Ophthalmic; Topical
Sodium Borate Decahydrate	Ophthalmic
Sodium Carbonate	Infiltration (AN); Intramuscular (IM); Infusion (IV); Intra-arterial; Intraperitoneal; Intrapleural; Intratumor; Intravascular; Intravenous; Intravitreal; Nerve Block; Ophthalmic; Rectal
Sodium Carbonate Decahydrate	Intravenous
Sodium Carbonate Monohydrate	Intra-arterial; Intracardiac; Intravenous; Ophthalmic
Sodium Cetostearyl Sulfate	Topical
Sodium Chlorate	Infiltration (AN); Intramuscular; Infusion (IV); Nerve Block
Sodium Chloride	Infiltration; Inhalation; Intra-arterial; Intra-articular; Intrabursal; Intracardiac; Intracaudal; Intracavitary; Intradermal; Intralesional; Intramuscular; Intraocular; Intraperitoneal; Intrapleural; Intrasynovial; Intrathecal; Intratracheal; Intratumor; Intravascular; Intravenous; Intravenous bolus; Intravesical; Intravitreal; Iontophoresis; Infusion (IV); Intramuscular (IM); Subcutaneous (SC); Nasal; Nerve Block; Ophthalmic; Parenteral; Peridural; Photopheresis; Rectal; Respiratory (Inhalation); Soft tissue; Subarachnoid; Submucosal; Topical; Transermal
Sodium Chloride Injection	Intramuscular

Sodium Chloride Injection, Bacteriostatic	Intraveous
Sodium Cholesteryl Sulfate	Infusion (IV)
Sodium Citrate	Infiltration (AN); Auricular (Otic); Epidural; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inhalation; Intra-arterial; Intra-articular; Intracardiac; Intravacitary; Intralesional; Intraocular; Iintraperitoneal; Intrapleural; Intrasynovial; Intrathecal; Intratracheal; Intrauterine; Intravasular; Intravenous; Iontophoresis; Irrigation; Nasal; Nerve Block; Ophthalmic; Rectal; Respiratory (Inhalation); Soft tissue; Topical; Transdermal; Ureteral; Vaginal
Sodium Cocoyl Sarcosinate	Topical
Sodium Desoxycholate	Infusion (IV)
Sodium Dithionite	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intravenous
Sodium Dodecylbenzenesulfonate	Topical
Sodium Formaldehyde Sulfoxylate	Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Topical
Sodium Gluconate	Intravenous; Infusion (IV)
Sodium Hydroxide	Intrathecal (AN, CNBLK); Inflitration (AN); Sympathetic (AN, NBLK); Auricular (Otic); Caudal Block; Dental; Epidural; Extracorporeal; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inflitration; Inhalationi; Interstitial; Intra-amniotic; Intra-arterial; Intra-articular; Intrabursal; Intracardiac; Intracaudl; Intracavitary; Intradermal; Intradiscal; Intralesional; Intraocular; Intraperioneal; Intrapleural; Intraspinal; Intrasynovial; Intrathecal; Intratracheal; Intratumor; Intrauterine; Intravascular; Intravenous; Intravitreal; Iontophoresis; Irrigation; Nasal; Nerve Block; Ophthalmic; Parenteral; Perfusion, cardiac; Peridural; Perineural; Photopheresis; Rectal; Respiratory (Inhalation); Retrobular; Soft tissue; Spinal; Subarachnoid; Subconjunctival; Submucosal; Topical; Transdermal; Ureteral; Urethral; Vaginal
Sodium Hypochlorite	Infusion (IV)
Sodium Iodide	Intravenous; Topical
Sodium Lactate	Infiltration (AN); Caudal Black; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intracardiac; Intraperitoneal; Intravenous; Nerve Block; Topical
Sodium Lactate, L-	Epidural; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intracardiac; Nerve Block
Sodium Laureth-2 Sulfate	Topical
Sodium Laureth-3 Sulfate	Topical
Sodium Laureth-5 Sulfate	Topical
Sodium Lauroyl Sarcosinate	Topical
Sodium Lauryl Sulfate	Dental; Respiratory (Inhalation); Topical; Vaginal
Sodium Lauryl Sulfoacetate	Topical
Sodium Metabisulfite	Intrathecal (AN, CNBLK); Infiltration (AN); Cardal Block; Dental; Epidural; Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Inflitration; Inhalation; Intra-articular; Initrabursal; Intracardiac; Intramuscular; Intraperitoneal; Intravenous; Iontophoresis; Nerve Block; Ophthalmic; Peridural; Rectal; Submucosal; Topical; Vaginal
Sodium Nitrate	Ophthalmic
Sodium Phosphate	Intramuscular (IM); Infusion (IV); Intra-articular; Intrabursal; Intradermal; Intralesional; Nasal; Nerve Block; Ophthalmic; Soft tissue; Subcutanesou; Topical

Sodium Phosphate Dihydrate	Intramuscular (IM); Subcutaneous (SC); Ophthalmic
Sodium Phosphate, Dibasic	Intramuscular (IM); Infusion (IV); Intradermal; Intralesional; Intrasynovial; Intravenous; Nasal; Ophthalmic; Soft tissue; Topical; Subcutaneous (SC)
Sodium Phosphate, Dibasic, Anhydrous	Auricular (Otic); Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intra-articular; Intralesional; Intramuscular; Intravenous; Intravesical; Nasal; Ophthalmic; Topical; Vaginal
Sodium Phosphate, Dibasic, Dihydrate	Intramuscular (IM); Infusion (IV); Intravenous; Nasal; Ophthalmic; Subcutaneous; Topical
Sodium Phosphate, Dibasic, Dodecahydrate	Nasal
Sodium Phosphate, Dibasic, Heptahydrate	Infiltration (AN); Auricular (Otic); Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Iintra-articular; Intrabursal; Intradermal; Intralesional; Intramuscular; Intrasynovial; Intravenous; Intravitreal; Nasal; Nerve Block; Ophthalmic; Soft tissue; Topical; Urethral
Sodium Phosphate, Monobasic	Intramuscular (IM); Infusion (IV); Intralesional; Intrasynovial; Iontophoresis; Ophthalmic; Soft tissue; Subcutaneous; Topical
Sodium Phosphate, Monobasic, Anhydrous	Auricular (Otic); Intramuscular (IM); Infusion (IV); Intrabursal; Intradermal; Intralesional; Intrasynovial; Intravascular; Intravenous; Intravesical; Nasal; Ophthalmic; Soft tissue; Subcutaneous; Topical; Vaginal
Sodium Phosphate, Monobasic, Dihydrate	Intravenous; Infusion (IV); Nasal; Ophthalmic; Subcutaneous; Topical
Sodium Phosphate, Monobasic, Monohydrate	Intramuscular (IM); Infusion (IV); Intra-articular; Intralesional; Intravascular; Intravenous; Intravitreal; Ophthalmic; Subcutaneous; Topical
Sodium Polyacrylate (2500000 Mw)	Topical
Sodium Pyrophosphate	Intravenous
Sodium Pyrrolidone Carboxylate	Topical
Sodium Starch Glycolate	Transmucosal
Sodium Succinate Hexahydrate	Intravenous
Sodium Sulfate	Intramuscular (IM); Infusion (IV); Ophthalmic
Sodium Sulfate Anhydrous	Inhalation; Iintramuscular; Ophthalmic
Sodium Sulfate Decahydrate	Ophthalmic
Sodium Sulfite	Auricular (Otic); Epidural; Intramuscular (IM); Infusion (IV); Inhalation; Intra-articular; Intralesional; Intravenous; Ophthalmic; Soft tissue; Subcutaneous; Topical
Sodium Sulfosuccinated Undecyclenic Monoalkylolamide	Topical
Sodium Tartrate	Intramuscual (IM); Infusion (IV); Intravenous
Sodium Thioglycolate	Subcutaneous
Sodium Thiomalate	Intramuscular (IM); Infusion (IV)
Sodium Thiosulfate	Intravenous; Ophthalmic; Topical
Sodium Thiosulfate Anhydrous	Intravenous
Sodium Trimetaphosphate	Intravenous
Sodium Xylenesulfonate	Topical
Somay 44	Topical
Sorbic Acid	Ophthalmic; Topical; Vaginal
Sorbitan	Topical
Sorbitan Isostearate	Topical
Sorbitan Monolaurate	Ophthalmic; Topical
Sorbitan Monooleate	Rectal; Topical; Transdermal

Sorbitan Monopalmitate	Intramuscular; Topical
Sorbitan Monostearate	Topical; Vaginal
Sorbitan Sesquioleate	Rectal; Topical
Sorbitan Trioleate	Inhalation; Nasal
Sorbitan Tristearate	Topical
Sorbitol	Dental; Intra-articular; Intralesional; Intramuscular; Intrasynovial; Intravenous; Infusion (IV); Nasal; Ophthalmic; Rectal; Topical; Vaginal
Sorbitol Solution	Intra-articular; Intralesional; Intramuscular; Intravenous; Infusion (IV); Nasal; Ophthalmic; Rectal; Topical; Vaginal
Soybean Flour	Topical
Soybean Oil	Intraveous; Infusion (IV); Topical
Spearmint Oil	Topical
Spermaceti	Topical; Vaginal
Squalane	Topical
Stabilized Oxychloro Complex	Ophthalmic
Stannous 2-Ethylhexanoate	Vaginl
Stannous Chloride	Intravenous; Infusion (IV)
Stannous Chloride Anhydrous	Intravenous; Infusion (IV)
Stannous Fluoride	Intravenous
Stannous Tartrate	Intravenous
Starch	Intramuscular; Rectal; Topical; Vaginal
Starch 1500, Pregelatinized	Vaginl
Starch, Corn	Vaginl
Stearalkonium Chloride	Topical
Stearalkonium Hectorite/Propylene Carbonate	Transdermal
Stearamidoethyl Diethylamine	Topical; Vaginal
Steareth-10	Rectal; Topical
Steareth-100	Topical
Steareth-2	Topical
Steareth-20	Topical
Steareth-21	Topical
Steareth-40	Topical; Rectal
Stearic Acid	Implantation; Subcutaneous; Topical; Vaginal
Stearic Diethanolamide	Topical
Stearoxytrimethylsilane	Topical
Steartrimonium Hydrolyzed Animal Collagen	Topical
Stearyl Alcohol	Topical; Vaginal
Sterile Water For Inhalation	Infusion (IV)
Styrene/Isoprene/Styrene Block Copolymer	Topical
Succimer	Intravenous
Succinic Acid	Intramuscular (IM); Infusion (IV); Intravenous
Sucralose	Nasa
Sucrose	Intramuscular; Intravenous; Infusion (IV); Rectal; Subcutaneous; Topical
Sucrose Distearate	Topical
Sucrose Polyesters	Topical
Sulfacetamide Sodium	Topical
Sulfobutylether .Beta.-Cyclodextrin	Intramuscular; Intravenous; Infusion (IV)
Sulfur Dioxide	Infusion (IV)
Sulfuric Acid	Auricular (Otic); Epidural; Intramuscular (IM); Infusion (IV); Inhalation; Intraperitoneal; Intravenous; Irrigation; Nasal; Ophthalmic; Respiratory (Inhalation); Topical

Sulfurous Acid	Intramuscular
Surfactol Qs	Topical
Tagatose, D-	Rectal
Talc	Topical
Tall Oil	Topical
Tallow Glycerides	Topical
Tartaric Acid	Intramuscular; Intravenous; Infusion (IV); Topical
Tartaric Acid, Dl-	Intramuscular (IM); Infusion (IV); Intravenous; Rectal; Vaginal
Tenox	Topical
Tenox-2	Topical
Tert-Butyl Alcohol	Intravenous; Infusion (IV); Topical
Tert-Butyl Hydroperoxide	Topical
Tert-Butylhydroquinone	Vaginl
Tetrakis(2-Methoxyisobutylisocyanide)Copper(I) Tetrafluoroborate	Intravenous
Tetrapropyl Orthosilicate	Vaginl
Tetrofosmin	Infusion (IV)
Theophylline	Intravenous; Infusion (IV)
Thimerosal	Auricular (Otic); Intramuscular (IM); Infusion (IV); Subcutaneous (SC); Intravenous; Ophthalmic; Topical
Threonine	Intravenous; Infusion (IV)
Thymol	Inhalation
Tin	Intravenous
Titanium Dioxide	Dental; Intrauterine; Ophthalmic; Respiratory (Inhalation); Topical; Transdermal
Tocopherol	Topical
Tocophersolan	Ophthalmic; Topical
Triacetin	Endocervical; Transdermal
Tricaprylin	Epidural; Infiltration
Trichloromonofluoromethane	Inhalation; Nasal; Topical
Trideceth-10	Topical
Triethanolamine Lauryl Sulfate	Topical
Trifluoroacetic Acid	Infusion (IV)
Triglycerides, Medium Chain	Topical
Trihydroxystearin	Topical
Trilaneth-4 Phosphate	Topical
Trilaureth-4 Phosphate	Topical
Trisodium Citrate Dihydrate	Intramuscular (IM); Infusion (IV); Intravenous; Intravitreal; Nasal; Ophthalmic; Topical
Trisodium Hedta	Topical
Triton 720	Ophthalmic
Triton X-200	Topical
Trolamine	Rectal; Topical; Transdermal; Vaginal
Tromantadine	Intramuscular; Intravenous
Tromethamine	Intramuscular (IM); Infusion (IV); Intra-arterial; Intrathecal; Intratracheal; Intravasular; Intravenous; Ophthalmic; Rectal; Respiratory (Inhalation); Subcutaneous; Topical; Transdermal; Urethral
Tryptophan	Infusion (IV)
Tyloxapol	Ophthalmic; Topical
Tyrosine	Infusion (IV)
Undecylenic Acid	Topical
Union 76 Amsco-Res 6038	Transdermal

Urea	Intramuscular; Vaginal
Valine	Infusion (IV)
Vegetable Oil	Topical
Vegetable Oil Glyceride, Hydrogenated	Rectal
Vegetable Oil, Hydrogenated	Rectal; Topical; Vaginal
Versetamide	Intravenous
Viscarin	Topical
Viscose/Cotton	Transdermal
Vitamin E	Topical
Wax, Emulsifying	Rectal; Topical
Wecobee Fs	Topical; Vaginal
White Ceresin Wax	Vagianl
White Wax	Rectal; Topical; Vaginal
Xanthan Gum	Rectal; Topical
Zinc	Subcutaneous
Zinc Acetate	Subcutaneous, Topical
Zinc Carbonate	Subcutaneous
Zinc Chloride	Intramuscular (IM); Subcutaneous (SC); Intradermal; Ophthalmic
Zinc Oxide	Intramuscular (IM); Subcutaneous (SC); Rectal; Respiratory (Inhalation)

Delivery

[0627] The present disclosure encompasses the delivery of NAVs for any of therapeutic, pharmaceutical, diagnostic or imaging by any appropriate route taking into consideration likely advances in the sciences of drug delivery. Delivery may be naked or formulated.

Formulated Delivery

[0628] The NAVs of the present invention may be formulated, using the methods described herein. The formulations contain polynucleotides which are modified. The formulations may further include, but are not limited to, cell penetration agents, a pharmaceutically acceptable carrier, a delivery agent, a bioerodible or biocompatible polymer, a solvent, and a sustained-release delivery depot. The formulated NAVs may be delivered to the cell using routes of administration known in the art and described herein.

[0629] The compositions may also be formulated for direct delivery to an organ or tissue in any of several ways in the art including, but not limited to, direct soaking or bathing, via a catheter, by gels, powder, ointments, creams, gels, lotions, and/or drops, by using substrates such as fabric or biodegradable materials coated or impregnated with the compositions, and the like.

Administration

[0630] The NAVs of the present invention may be administered by any route which results in a therapeutically effective outcome. These include, but are not limited to enteral (into the intestine), gastroenteral, epidural (into the dura matter), oral (by way of the mouth), transdermal, peridural, intracerebral (into the cerebrum), intracerebroventricular (into the cerebral ventricles), epicutaneous (application onto the skin), intradermal, (into the skin itself), subcutaneous (under the skin), nasal administration (through the nose), intravenous (into a vein), intravenous bolus, intravenous drip, intraarterial (into an artery), intramuscular (into a muscle), intracardiac (into the heart), intraosseous infusion (into the bone marrow), intrathecal (into the spinal canal), intraperitoneal, (infusion or injection into the peritoneum), intravesical infusion, intravitreal, (through the eye), intracavernous injection (into a pathologic cavity) intracavitary (into the base of the penis), intravaginal administration, intrauterine, extra-amniotic administration, transdermal (diffusion through the intact skin for systemic distribution), transmucosal (diffusion through a mucous membrane), transvaginal, insufflation (snorting), sublingual, sublabial, enema, eye drops (onto the conjunctiva), in ear drops, auricular (in or by way of the ear), buccal (directed toward the cheek), conjunctival, cutaneous, dental (to a tooth or teeth), electro-osmosis, endocervical, endosi-

nusial, endotracheal, extracorporeal, hemodialysis, infiltration, interstitial, intra-abdominal, intra-amniotic, intra-articular, intrabiliary, intrabronchial, intrabursal, intracartilaginous (within a cartilage), intracaudal (within the cauda equine), intracisternal (within the cisterna magna cerebellomedularis), intracorneal (within the cornea), dental intracornal, intracoronary (within the coronary arteries), intracorporus cavernosum (within the dilatable spaces of the corporus cavernosa of the penis), intradiscal (within a disc), intraductal (within a duct of a gland), intraduodenal (within the duodenum), intradural (within or beneath the dura), intraepidermal (to the epidermis), intraesophageal (to the esophagus), intragastric (within the stomach), intragingival (within the gingivae), intraileal (within the distal portion of the small intestine), intralesional (within or introduced directly to a localized lesion), intraluminal (within a lumen of a tube), intralymphatic (within the lymph), intramedullary (within the marrow cavity of a bone), intrameningeal (within the meninges), intraocular (within the eye), intraovarian (within the ovary), intrapericardial (within the pericardium), intrapleural (within the pleura), intraprostatic (within the prostate gland), intrapulmonary (within the lungs or its bronchi), intrasinal (within the nasal or periorbital sinuses), intraspinal (within the vertebral column), intrasynovial (within the synovial cavity of a joint), intratendinous (within a tendon), intratesticular (within the testicle), intrathecal (within the cerebrospinal fluid at any level of the cerebrospinal axis), intrathoracic (within the thorax), intratubular (within the tubules of an organ), intratumor (within a tumor), intratympanic (within the aurus media), intravascular (within a vessel or vessels), intraventricular (within a ventricle), iontophoresis (by means of electric current where ions of soluble salts migrate into the tissues of the body), irrigation (to bathe or flush open wounds or body cavities), laryngeal (directly upon the larynx), nasogastric (through the nose and into the stomach), occlusive dressing technique, ophthalmic (to the external eye), oropharyngeal (directly to the mouth and pharynx), parenteral, percutaneous, periarticular, peridural, perineural, periodontal, rectal, respiratory (within the respiratory tract by inhaling orally or nasally for local or systemic effect), retrobulbar (behind the pons or behind the eyeball), intramyocardial (entering the myocardium), soft tissue, subarachnoid, subconjunctival, submucosal, transplacental (through or across the placenta), transtracheal (through the wall of the trachea), transtympanic (across or through the tympanic cavity), ureteral (to the ureter), urethral (to the urethra), vaginal, caudal block, diagnostic, nerve block, biliary perfusion, cardiac perfusion, photopheresis or spinal. In specific instances, compositions may be administered in a way which allows them cross the blood-brain barrier, vascular barrier, or other epithelial barrier. In one instance, a formulation for a route of administration may include at least one inactive ingredient. Non-limiting examples of routes of administration and inactive ingredients which may be included in formulations for the specific route of administration is shown in Table 20. In Table 27, "AN" means anesthetic, "CNBLK" means cervical nerve block, "NBLK" means nerve block, "IV" means intravenous, "IM" means intramuscular and "SC" means subcutaneous.

Table 27. Routes of Adminsitration and Inactive Ingredients

Route of Administration	Inactive Ingredient
Intrathecal (AN, CNBLK)	Acetone Sodium Bisulfite; Citric Acid; Hydrochloric Acid; Sodium Chloride; Sodium Hydroxide; Sodium Metabisulfite
Infiltration (AN)	Acetic Acid; Acetone Sodium Bisulfite; Ascorbic Acid; Benzyl Alcohol; Calcium Chloride;

	Carbon Dioxide; Chlorobutanol; Citric Acid; Citric Acid Monohydrate; Edetate Calcium Disodium; Edetate Disodium; Hydrochloric Acid; Hydrochloric Acid, Diluted; Lactic Acid; Methylparaben; Monothioglycerol; Nitrogen; Potassium Chloride; Potassium Metabisulfite; Potassium Phosphate, Monobasic; Propylparaben; Sodium Bisulfite; Sodium Carbonate; Sodium Chlorate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Lactate; Sodium Metabisulfite; Sodium Phosphate, Dibasic, Heptahydrate
Sympathetic NBLK (AN)	Hydrochloric Acid; Sodium Chloride; Sodium Hydroxide
Auricular (Otic)	Acetic Acid; Aluminum Acetate; Aluminum Sulfate Anhydrous; Benzalkonium Chloride; Benzethonium Chloride; Benzyl Alcohol; Boric Acid; Calcium Carbonate; Cetyl Alcohol; Chlorobutanol; Chloroxylenol; Citric Acid; Creatinine; Cupric Sulfate; Cupric Sulfate Anhydrous; Edetate Disodium; Edetic Acid; Glycerin; Glyceryl Stearate; Hydrochloric Acid; Hydrocortisone; Hydroxyethyl Cellulose; Isopropyl Myristate; Lactic Acid; Lecithin, Hydrogenated; Methylparaben; Mineral Oil; Petrolatum; Petrolatum, White; Phenylethyl Alcohol; Polyoxyl 40 Stearate; Polyoxyl Stearate; Polysorbate 20; Polysorbate 80; Polyvinyl Alcohol; Potassium Metabisulfite; Potassium Phosphate, Monobasic; Povidone K90f; Povidones; Propylene Glycol; Propylene Glycol Diacetate; Propylparaben; Sodium Acetate; Sodium Bisulfite; Sodium Borate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic, Anhydrous; Sodium Sulfite; Sulfuric Acid; Thimerosal
Caudal Block	Ascorbic Acid; Calcium Chloride; Citric Acid; Edetate Calcium Disodium; Edetate Disodium; Hydrochloric Acid; Methylparaben; Monothioglycerol; Nitrogen; Potassium Chloride; Sodium Chloride; Sodium Hydroxide; Sodium Lactate; Sodium Metabisulfite
Dental	Acetone Sodium Bisulfite; Alcohol; Alcohol, Dehydrated; Alcohol, Denatured; Anethole; Benzyl Alcohol; Carboxymethylcellulose Sodium; Carrageenan; D&C Yellow No. 10; Dimethicone Medical Fluid 360; Eucalyptol; Fd&C Blue No. 1; Fd&C Green No. 3; Flavor 89-186; Flavor 89-259; Flavor Df-119; Flavor Df-1530; Flavor Enhancer; Gelatin; Gelatin, Crosslinked; Glycerin; Glyceryl Stearate; High Density Polyethylene; Hydrocarbon Gel, Plasticized; Hydrochloric Acid; Menthol; Mineral Oil; Nitrogen; Pectin; Peg-40 Sorbitan Diisostearate; Peppermint Oil; Petrolatum, White; Plastibase-50w; Polyethylene Glycol 1540; Polyglactin; Polyols; Polyoxyl 40 Hydrogenated Castor Oil; Polyoxyl 40 Stearate; Propylene Glycol; Pvm/Ma Copolymer; Saccharin Sodium; Silica, Dental; Silicon Dioxide; Sodium Benzoate; Sodium Chloride; Sodium Hydroxide; Sodium Lauryl Sulfate; Sodium Metabisulfite; Sorbitol; Titanium Dioxide
Diagnostic	Hydrochloric Acid
Endocervical	Colloidal Silicon Dioxide; Triacetin
Epidural	1,2-Dioleoyl-Sn-Glycero-3-Phosphocholine; 1,2-Dipalmitoyl-Sn-Glycero-3-(Phospho-Rac-(1-Glycerol)); Ascorbic Acid; Benzyl Alcohol; Calcium Chloride; Cholesterol; Citric Acid; Edetate Calcium Disodium; Edetate Disodium; Glyceryl Trioleate; Hydrochloric Acid; Isotonic Sodium Chloride Solution; Methylparaben; Monothioglycerol; Nitrogen; Potassium Chloride; Sodium Bisulfite; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Lactate, L-; Sodium Metabisulfite; Sodium Sulfite; Sulfuric Acid; Tricaprylin
Extracorporeal	Acetic Acid; Alcohol, Dehydrated; Benzyl Alcohol; Hydrochloric Acid; Propylene Glycol; Sodium Acetate; Sodium Chloride; Sodium Hydroxide
Intramuscular-Intravenous	Acetic Acid; Alcohol; Alcohol, Dehydrated; Alcohol, Diluted; Anhydrous Dextrose; Anhydrous Lactose; Anhydrous Trisodium Citrate; Arginine; Ascorbic Acid; Benzethonium Chloride; Benzoic Acid; Benzyl Alcohol; Calcium Chloride; Carbon Dioxide; Chlorobutanol; Citric Acid; Citric Acid Monohydrate; Creatinine; Dextrose; Edetate Calcium Disodium; Edetate Disodium; Edetate Sodium; Gluconolactone; Glycerin; Hydrochloric Acid; Hydrochloric Acid, Diluted; Lactic Acid; Lactic Acid, Dl-; Lactose; Lactose Monchydrate; Lactose, Hydrous; Lysine; Mannitol; Methylparaben; Monothioglycerol; Niacinamide; Nitrogen; Phenol; Phenol, Liquefied; Phosphoric Acid; Polyethylene Glycol 300; Polyethylene Glycol 400; Polypropylene Glycol; Polysorbate 40; Potassium Metabisulfite; Potassium Phosphate, Monobasic; Propylene Glycol; Propylparaben; Saccharin Sodium; Saccharin Sodium Anhydrous; Silicone; Simethicone; Sodium Acetate; Sodium Acetate Anhydrous; Sodium Benzoate; Sodium Bicarbonate; Sodium Bisulfate; Sodium Bisulfite; Sodium Carbonate; Sodium Chloride; Sodium Citrate; Sodium Formaldehyde Sulfoxylate; Sodium Hydroxide; Sodium Lactate, L-; Sodium Metabisulfite; Sodium Phosphate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Dihydrate; Sodium Phosphate, Dibasic, Heptahydrate; Sodium

	Phosphate, Monobasic; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Monohydrate; Sodium Sulfate; Sodium Sulfite; Sodium Tartrate; Sodium Thiomalate; Succinic Acid; Sulfuric Acid; Tartaric Acid, Dl-; Thimerosal; Trisodium Citrate Dihydrate; Tromethamine
Intramuscular-Intravenous-Subcutaneous	Acetic Acid; Alcohol; Alcohol, Dehydrated; Benzyl Alcohol; Chlorobutanol; Citric Acid; Citric Acid Monohydrate; Citric Acid, Hydrous; Creatinine; Dextrose; Edetate Disodium; Edetate Sodium; Gelatin; Glycerin; Glycine; Hydrochloric Acid; Hydrochloric Acid, Diluted; Lactic Acid; Lactose; Lactose Monohydrate; Metacresol; Methanesulfonic Acid; Methylparaben; Monothioglycerol; Nitrogen; Phenol; Phosphoric Acid; Polyoxyethylene Fatty Acid Esters; Propylparaben; Sodium Acetate; Sodium Bisulfate; Sodium Bisulfite; Sodium Chloride; Sodium Citrate; Sodium Dithionite; Sodium Hydroxide; Sodium Lactate; Sodium Lactate, L-; Sodium Metabisulfite; Sodium Phosphate, Dibasic, Heptahydrate; Thimerosal
Intramuscular –Subcutaneous	Acetic Acid; Anhydrous Dextrose; Benzyl Alcohol; Chlorobutanol; Citric Acid; Cysteine; Edetate Disodium; Gelatin; Glycerin; Glycine; Hydrochloric Acid; Lactose Monohydrate; Mannitol; Metacresol; Methylparaben; Nitrogen; Peg Vegetable Oil; Peg-40 Castor Oil; Phenol; Phenol, Liquefied; Phosphoric Acid; Polyoxyethylene Fatty Acid Esters; Polysorbate 20; Propylparaben; Protamine Sulfate; Sesame Oil; Sodium Acetate; Sodium Acetate Anhydrous; Sodium Chloride; Sodium Citrate; Sodium Formaldehyde Sulfoxylate; Sodium Hydroxide; Sodium Phosphate Dihydrate; Sodium Phosphate, Dibasic, Heptahydrate; Sulfuric Acid; Thimerosal; Zinc Chloride; Zinc Oxide
Implantation	Acetone; Crospovidone; Dimethylsiloxane/Methylvinylsiloxane Copolymer; Ethylene Vinyl Acetate Copolymer; Magnesium Stearate; Poly(Bis(P-Carboxyphenoxy)Propane Anhydride):Sebacic Acid; Polyglactin; Silastic Brand Medical Grade Tubing; Silastic Medical Adhesive,Silicone Type A; Stearic Acid
Infiltration	Cholesterol; Citric Acid; Diethyl Pyrocarbonate; Dipalmitoylphosphatidylglycerol, Dl-; Hydrochloric Acid; Nitrogen; Phosphoric Acid; Sodium Chloride; Sodium Hydroxide; Sodium Metabisulfite; Tricaprylin
Inhalation	Acetone Sodium Bisulfite; Acetylcysteine; Alcohol; Alcohol, Dehydrated; Ammonia; Ascorbic Acid; Benzalkonium Chloride; Carbon Dioxide; Cetylpyridinium Chloride; Chlorobutanol; Citric Acid; D&C Yellow No. 10; Dichlorodifluoromethane; Dichlorotetrafluoroethane; Edetate Disodium; Edetate Sodium; Fd&C Yellow No. 6; Fluorochlorohydrocarbons; Glycerin; Hydrochloric Acid; Hydrochloric Acid, Diluted; Lactose; Lecithin; Lecithin, Hydrogenated Soy; Lecithin, Soybean; Menthol; Methylparaben; Nitric Acid; Nitrogen; Norflurane; Oleic Acid; Propylene Glycol; Propylparaben; Saccharin; Saccharin Sodium; Sodium Bisulfate; Sodium Bisulfite; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Metabisulfite; Sodium Sulfate Anhydrous; Sodium Sulfite; Sorbitan Tricleate; Sulfuric Acid; Thymol; Trichloromonofluoromethane
Interstitial	Benzyl Alcohol; Dextrose; Hydrochloric Acid; Sodium Acetate; Sodium Hydroxide
Intra-amniotic	Citric Acid; Edetate Disodium Anhydrous; Hydrochloric Acid; Sodium Hydroxide
Intra-arterial	Anhydrous Trisodium Citrate; Benzyl Alcohol; Carbon Dioxide; Citric Acid; Diatrizoic Acid; Edetate Calcium Disodium; Edetate Disodium; Hydrochloric Acid; Hydrochloric Acid, Diluted; Iodine; Meglumine; Methylparaben; Nitrogen; Propylparaben; Sodium Bisulfite; Sodium Carbonate; Sodium Carbonate Monohydrate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Tromethamine
Intra-articular	Acetic Acid; Anhydrous Trisodium Citrate; Benzalkonium Chloride; Benzyl Alcohol; Carboxymethylcellulose; Carboxymethylcellulose Sodium; Cellulose, Microcrystalline; Citric Acid; Creatine; Creatinine; Crospovidone; Diatrizoic Acid; Edetate Calcium Disodium; Edetate Disodium; Hyaluronate Sodium; Hydrochloric Acid; Iodine; Meglumine; Methylcelluloses; Methylparaben; Myristyl-.Gamma.-Picolinium Chloride; Niacinamide; Phenol; Phosphoric Acid; Polyethylene Glycol 3350; Polyethylene Glycol 4000; Polysorbate 80; Potassium Phosphate, Dibasic; Potassium Phosphate, Monobasic; Propylparaben; Sodium Acetate; Sodium Bisulfite; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Metabisulfite; Sodium Phosphate; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Monohydrate; Sodium Sulfite; Sorbitol; Sorbitol Solution
Intrabursal	Anhydrous Trisodium Citrate; Benzalkonium Chloride; Benzyl Alcohol; Carboxymethylcellulose; Carboxymethylcellulose Sodium; Citric Acid; Creatinine; Edetate Disodium; Hydrochloric Acid;

	Methylparaben; Polysorbate 80; Propylparaben; Sodium Bisulfite; Sodium Chloride; Sodium Hydroxide; Sodium Metabisulfite; Sodium Phosphate; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic, Anhydrous
Intracardiac	Carbon Dioxide; Citric Acid; Citric Acid Monohydrate; Diatrizoic Acid; Edetate Calcium Disodium; Edetate Disodium; Hydrochloric Acid; Iodine; Lactic Acid; Meglumine; Sodium Bisulfite; Sodium Carbonate Monohydrate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Lactate; Sodium Lactate, L-; Sodium Metabisulfite
Intracaudal	Hydrochloric Acid; Sodium Chloride; Sodium Hydroxide
Intracavitary	Alcohol, Dehydrated; Alfadex; Anhydrous Lactose; Benzyl Alcohol; Dextrose; Hydrochloric Acid; Lactose; Lactose Monohydrate; Nitrogen; Sodium Acetate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide
Intradermal	Benzalkonium Chloride; Benzyl Alcohol; Carboxymethylcellulose Sodium; Creatinine; Edetate Disodium; Glycerin; Hydrochloric Acid; Metacresol; Methylparaben; Phenol; Polysorbate 80; Protamine Sulfate; Sodium Acetate; Sodium Bisulfite; Sodium Chloride; Sodium Hydroxide; Sodium Phosphate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic, Anhydrous; Zinc Chloride
Intradiscal	Cysteine Hydrochloride Anhydrous; Cysteine, Dl-; Diatrizoic Acid; Edetate Calcium Disodium; Edetate Disodium; Iodine; Meglumine; Sodium Bisulfite; Sodium Hydroxide
Intralesional	Acetic Acid; Benzalkonium Chloride; Benzyl Alcohol; Carboxymethylcellulose; Carboxymethylcellulose Sodium; Citric Acid; Creatine; Creatinine; Edetate Disodium; Hydrochloric Acid; Methylcelluloses; Methylparaben; Myristyl-.Gamma.-Picolinium Chloride; Niacinamide; Phenol; Phosphoric Acid; Polyethylene Glycol 3350; Polyethylene Glycol 4000; Polysorbate 80; Propylparaben; Sodium Acetate; Sodium Bisulfite; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Phosphate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Monohydrate; Sodium Sulfite; Sorbitol; Sorbitol Solution
Intralymphatic	Poppy Seed Oil
Intramuscular	Acetic Acid; Activated Charcoal; Adipic Acid; Alcohol; Alcohol, Dehydrated; Ammonium Acetate; Anhydrous Dextrose; Ascorbic Acid; Benzalkonium Chloride; Benzethonium Chloride; Benzoic Acid; Benzyl Alcohol; Benzyl Benzoate; Butylated Hydroxyanisole; Butylated Hydroxytoluene; Butylparaben; Calcium; Calcium Chloride; Carbon Dioxide; Carboxymethylcellulose; Carboxymethylcellulose Sodium; Castor Oil; Cellulose, Microcrystalline; Chlorobutanol; Chlorobutanol Hemihydrate; Chlorobutanol, Anhydrous; Citric Acid; Citric Acid Monohydrate; Corn Oil; Cottonseed Oil; Creatine; Creatinine; Croscarmellose Sodium; Crospovidone; Dextrose; Diatrizoic Acid; Docusate Sodium; Edetate Calcium Disodium; Edetate Disodium; Edetate Disodium Anhydrous; Edetate Sodium; Ethyl Acetate; Gelatin; Glutathione; Glycerin; Glycine; Hyaluronate Sodium; Hydrochloric Acid; Hydroxide Ion; Lactic Acid; Lactic Acid, Dl-; Lactose; Lactose Monohydrate; Lactose, Hydrous; Lecithin; Magnesium Chloride; Maleic Acid; Mannitol; Meglumine; Metacresol; Methionine; Methylcelluloses; Methylparaben; Monothioglycerol; Myristyl-.Gamma.-Picolinium Chloride; N,N-Dimethylacetamide; Niacinamide; Nitrogen; Peanut Oil; Peg-20 Sorbitan Isostearate; Phenol; Phenylmercuric Nitrate; Phosphoric Acid; Polyethylene Glycol 200; Polyethylene Glycol 300; Polyethylene Glycol 3350; Polyethylene Glycol 4000; Polyglactin; Polylactide; Polysorbate 20; Polysorbate 40; Polysorbate 80; Polyvinyl Alcohol; Potassium Phosphate, Dibasic; Potassium Phosphate, Monobasic; Povidones; Propyl Gallate; Propylene Glycol; Propylparaben; Saccharin Sodium; Saccharin Sodium Anhydrous; Sesame Oil; Sodium Acetate; Sodium Acetate Anhydrous; Sodium Benzoate; Sodium Bicarbonate; Sodium Bisulfite; Sodium Carbonate; Sodium Chlorate; Sodium Chloride; Sodium Chloride Injection; Sodium Citrate; Sodium Formaldehyde Sulfoxylate; Sodium Hydroxide; Sodium Metabisulfite; Sodium Phosphate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Monohydrate; Sodium Sulfate Anhydrous; Sodium Sulfite; Sodium Tartrate; Sorbitan Monopalmitate; Sorbitol; Sorbitol Solution; Starch; Sucrose; Sulfobutylether .Beta.-Cyclodextrin; Sulfuric Acid; Sulfurous Acid; Tartaric Acid; Thimerosal; Tromantadine; Tromethamine; Urea
Intraocular	Benzalkonium Chloride; Calcium Chloride; Citric Acid Monohydrate; Hydrochloric Acid;

	Magnesium Chloride; Polyvinyl Alcohol; Potassium Chloride; Sodium Acetate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide
Intraperitoneal	Benzyl Alcohol; Calcium Chloride; Dextrose; Edetate Calcium Disodium; Hydrochloric Acid; Magnesium Chloride; Sodium Acetate; Sodium Bicarbonate; Sodium Bisulfite; Sodium Carbonate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Lactate; Sodium Metabisulfite; Sulfuric Acid
Intrapleural	Benzyl Alcohol; Citric Acid; Dextrose; Dichlorodifluoromethane; Hydrochloric Acid; Sodium Acetate; Sodium Carbonate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide
Intraspinal	Dextrose; Hydrochloric Acid; Sodium Hydroxide
Intrasynovial	Acetic Acid; Benzyl Alcohol; Carboxymethylcellulose Sodium; Citric Acid; Creatinine; Edetate Disodium; Hydrochloric Acid; Methylcelluloses; Methylparaben; Myristyl-.Gamma.-Picolinium Chloride; Niacinamide; Phenol; Polyethylene Glycol 3350; Polyethylene Glycol 4000; Polysorbate 80; Propylparaben; Sodium Acetate; Sodium Bisulfite; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic; Sodium Phosphate, Monobasic, Anhydrous; Sorbitol
Intrathecal	Benzyl Alcohol; Carbon Dioxide; Citric Acid; Edetate Calcium Disodium; Hydrochloric Acid; Methionine; Nitrogen; Pentetate Calcium Trisodium; Pentetic Acid; Sodium Bicarbonate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sulfuric Acid; Tromethamine
Intratracheal	Acetic Acid; Benzyl Alcohol; Carboxymethylcellulose Sodium; Hydrochloric Acid; Isotonic Sodium Chloride Solution; Peanut Oil; Sodium Bicarbonate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Tromethamine
Intratumor	Benzyl Alcohol; Hydrochloric Acid; Nitrogen; Sodium Carbonate; Sodium Chloride; Sodium Hydroxide
Intrauterine	Barium Sulfate; Crospovidone; Diatrizoic Acid; Dimethylsiloxane/Methylvinylsiloxane Copolymer; Edetate Calcium Disodium; Edetate Disodium; Ethylene Vinyl Acetate Copolymer; High Density Polyethylene; Meglumine; Polyethylene High Density Containing Ferric Oxide Black (<1%); Polyethylene Low Density Containing Barium Sulfate (20-24%); Polyethylene T; Polypropylene; Poppy Seed Oil; Potassium Phosphate, Monobasic; Silicone; Sodium Citrate; Sodium Hydroxide; Titanium Dioxide
Intravascular	Alcohol; Alcohol, Dehydrated; Calcium Chloride; Carbon Dioxide; Citric Acid; Diatrizoic Acid; Edetate Calcium Disodium; Edetate Disodium; Hydrochloric Acid; Hydrochloric Acid, Diluted; Iodine; Meglumine; Nitrogen; Potassium Hydroxide; Sodium Carbonate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Monohydrate; Tromethamine
Intravenous	Alpha-Tocopherol; Alpha-Tocopherol, Dl-; 1,2-Dimyristoyl-Sn-Glycero-3-Phosphocholine; 1,2-Distearoyl-Sn-Glycero-3-(Phospho-Rac-(1-Glycerol)); 1,2-Distearoyl-Sn-Glycero-3-Phosphocholine; Acetic Acid; Acetic Acid, Glacial; Acetic Anhydride; Acetylated Monoglycerides; Acetyltryptophan, Dl-; Activated Charcoal; Albumin Aggregated; Albumin Colloidal; Albumin Human; Alcohol; Alcohol, Dehydrated; Alcohol, Denatured; Ammonium Acetate; Ammonium Hydroxide; Ammonium Sulfate; Anhydrous Citric Acid; Anhydrous Dextrose; Anhydrous Lactose; Anhydrous Trisodium Citrate; Arginine; Ascorbic Acid; Benzenesulfonic Acid; Benzethonium Chloride; Benzoic Acid; Benzyl Alcohol; Benzyl Chloride; Bibapcitide; Boric Acid; Butylated Hydroxytoluene; Calcium Chloride; Calcium Gluceptate; Calcium Hydroxide; Calcobutrol; Caldiamide Sodium; Caloxetate Trisodium; Calteridol Calcium; Captisol; Carbon Dioxide; Cellulose, Microcrystalline; Chlorobutanol; Chlorobutanol Hemihydrate; Chlorobutanol, Anhydrous; Cholesterol; Citrate; Citric Acid; Citric Acid Monohydrate; Citric Acid, Hydrous; Cysteine; Cysteine Hydrochloride; Dalfampridine; Dextran; Dextran 40; Dextrose; Dextrose Monohydrate; Dextrose Solution; Diatrizoic Acid; Dimethicone Medical Fluid 360; Edetate Calcium Disodium; Edetate Disodium; Edetate Disodium Anhydrous; Egg Phospholipids; Ethanolamine Hydrochloride; Ethylenediamine; Exametazime; Ferric Chloride; Gadolinium Oxide; Gamma Cyclodextrin; Gelatin; Gentisic Acid; Gluceptate Sodium; Gluceptate Sodium Dihydrate; Gluconolactone; Glucuronic Acid; Glycerin; Glycine; Guanidine Hydrochloride; Hetastarch; Histidine; Human Albumin Microspheres; Hydrochloric Acid; Hydrochloric Acid, Diluted; Hydroxyethylpiperazine Ethane Sulfonic Acid; Hydroxypropyl-Bcyclodextrin; Iodine; Iodoxamic Acid; Iofetamine Hydrochloride; Isopropyl Alcohol; Isotonic Sodium Chloride Solution; Lactic Acid; Lactic Acid, Dl-; Lactic Acid, L-; Lactobionic Acid;

	Lactose; Lactose Monohydrate; Lactose, Hydrous; Lecithin, Egg; Lecithin, Hydrogenated Soy; Lidofenin; Mannitol; Mebrofenin; Medronate Disodium; Medronic Acid; Meglumine; Methionine; Methylboronic Acid; Methylene Blue; Methylparaben; Monothioglycerol; N-(Carbamoyl-Methoxy Peg-40)-1,2-Distearoyl-Cephalin Sodium; N,N-Dimethylacetamide; Nioxime; Nitrogen; Octanoic Acid; Oxidronate Disodium; Oxyquinoline; Pentasodium Pentetate; Pentetate Calcium Trisodium; Pentetic Acid; Perflutren; Phenol; Phenol, Liquefied; Phosphatidyl Glycerol, Egg; Phospholipid, Egg; Phosphoric Acid; Poloxamer 188; Polyethylene Glycol 300; Polyethylene Glycol 400; Polyethylene Glycol 600; Polysiloxane; Polysorbate 20; Polysorbate 80; Potassium Bisulfite; Potassium Chloride; Potassium Hydroxide; Potassium Metabisulfite; Potassium Phosphate, Dibasic; Potassium Phosphate, Monobasic; Povidones; Propylene Glycol; Propylparaben; Saccharin Sodium; Sodium Acetate; Sodium Acetate Anhydrous; Sodium Ascorbate; Sodium Benzoate; Sodium Bicarbonate; Sodium Bisulfite; Sodium Carbonate; Sodium Carbonate Decahydrate; Sodium Carbonate Monohydrate; Sodium Chloride; Sodium Chloride Injection, Bacteriostatic; Sodium Citrate; Sodium Dithionite; Sodium Gluconate; Sodium Hydroxide; Sodium Iodide; Sodium Lactate; Sodium Metabisulfite; Sodium Phosphate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Dihydrate; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Dihydrate; Sodium Phosphate, Monobasic, Monohydrate; Sodium Pyrophosphate; Sodium Succinate Hexahydrate; Sodium Sulfite; Sodium Tartrate; Sodium Thiosulfate; Sodium Thiosulfate Anhydrous; Sodium Trimetaphosphate; Sorbitol; Sorbitol Solution; Soybean Oil; Stannous Chloride; Stannous Chloride Anhydrous; Stannous Fluoride; Stannous Tartrate; Succimer; Succinic Acid; Sucrose; Sulfobutylether .Beta.-Cyclodextrin; Sulfuric Acid; Tartaric Acid; Tartaric Acid, Dl-; Tert-Butyl Alcohol; Tetrakis(2-Methoxyisobutylisocyanide)Copper(I) Tetrafluoroborate; Theophylline; Thimerosal; Threonine; Tin; Trisodium Citrate Dihydrate; Tromantadine; Tromethamine; Versetamide
Intravenous Bolus	Sodium Chloride
Intravesical	Alcohol, Dehydrated; Edetate Calcium Disodium; Hydrochloric Acid; Nitrogen; Polyoxyl 35 Castor Oil; Potassium Phosphate, Monobasic; Sodium Chloride; Sodium Hydroxide; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Monobasic, Anhydrous
Intravitreal	Calcium Chloride; Carboxymethylcellulose Sodium; Cellulose, Microcrystalline; Hyaluronate Sodium; Hydrochloric Acid; Magnesium Chloride; Magnesium Stearate; Polysorbate 80; Polyvinyl Alcohol; Potassium Chloride; Sodium Acetate; Sodium Bicarbonate; Sodium Carbonate; Sodium Chloride; Sodium Hydroxide; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic, Monohydrate; Trisodium Citrate Dihydrate
Iontophoresis	Cetylpyridinium Chloride; Citric Acid; Edetate Disodium; Glycerin; Hydrochloric Acid; Methylparaben; Phenonip; Polacrilin; Polyvinyl Alcohol; Povidone Hydrogel; Sodium Bisulfite; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Metabisulfite; Sodium Phosphate, Monobasic
Irrigation	Acetic Acid; Activated Charcoal; Benzoic Acid; Hydrochloric Acid; Hypromelloses; Methylparaben; Nitrogen; Sodium Bisulfite; Sodium Citrate; Sodium Hydroxide; Sulfuric Acid
Intravenous – Subcutaneous	Acetic Acid; Alcohol; Benzyl Alcohol; Calcium Hydroxide; Chlorobutanol; Glycerin; Hydrochloric Acid; Lactose Monohydrate; Methylparaben; Nitrogen; Phenol; Phenol, Liquefied; Phosphoric Acid; Propylparaben; Sodium Acetate; Sodium Carbonate; Sodium Chloride; Sodium Hydroxide
Intravenous (Infusion)	1,2-Dimyristoyl-Sn-Glycero-3-(Phospho-S-(1-Glycerol)); 1,2-Dimyristoyl-Sn-Glycero-3-Phosphocholine; Acetic Acid; Acetic Acid, Glacial; Activated Charcoal; Alanine; Albumin Human; Alcohol; Alcohol, Dehydrated; Ammonium Acetate; Anhydrous Citric Acid; Anhydrous Dextrose; Anhydrous Lactose; Anhydrous Trisodium Citrate; Arginine; Ascorbic Acid; Aspartic Acid; Benzenesulfonic Acid; Benzethonium Chloride; Benzoic Acid; Benzyl Alcohol; Brocrinat; Butylated Hydroxyanisole; Butylated Hydroxytoluene; Carbon Dioxide; Chlorobutanol; Citric Acid; Citric Acid Monohydrate; Citric Acid, Hydrous; Cysteine; Cysteine Hydrochloride; Deoxycholic Acid; Dextrose; Dextrose Solution; Diatrizoic Acid; Diethanolamine; Dimethyl Sulfoxide; Disodium Sulfosalicylate; Disofenin; Edetate Calcium Disodium; Edetate Disodium; Edetate Disodium Anhydrous; Edetate Sodium; Egg Phospholipids; Ethylenediamine; Fructose; Gelatin; Gentisic Acid Ethanolamide; Glycerin; Glycine; Histidine; Hydrochloric Acid; Hydrochloric Acid, Diluted; Hydroxide Ion; Hydroxypropyl-Bcyclodextrin; Isoleucine; Isotonic Sodium Chloride Solution; Lactic Acid; Lactic Acid, Dl-; Lactobionic Acid; Lactose; Lactose Monohydrate; Lactose, Hydrous; Leucine; Lysine; Lysine Acetate; Magnesium Chloride; Maleic

	Acid; Mannitol; Meglumine; Metacresol; Metaphosphoric Acid; Methanesulfonic Acid; Methionine; Methylparaben; Monothioglycerol; N,N-Dimethylacetamide; Nitric Acid; Nitrogen; Peg Vegetable Oil; Peg-40 Castor Oil; Peg-60 Castor Oil; Pentetate Calcium Trisodium; Phenol; Phenylalanine; Phospholipid; Phospholipid, Egg; Phosphoric Acid; Polyethylene Glycol 300; Polyethylene Glycol 400; Polyoxyl 35 Castor Oil; Polysorbate 20; Polysorbate 80; Potassium Chloride; Potassium Hydroxide; Potassium Metabisulfite; Potassium Phosphate, Dibasic; Potassium Phosphate, Monobasic; Povidones; Proline; Propylene Glycol; Propylparaben; Saccharin Sodium; Saccharin Sodium Anhydrous; Serine; Sodium Acetate; Sodium Acetate Anhydrous; Sodium Benzoate; Sodium Bicarbonate; Sodium Bisulfite; Sodium Carbonate; Sodium Chlorate; Sodium Chloride; Sodium Cholesteryl Sulfate; Sodium Citrate; Sodium Desoxycholate; Sodium Dithionite; Sodium Formaldehyde Sulfoxylate; Sodium Gluconate; Sodium Hydroxide; Sodium Hypochlorite; Sodium Lactate; Sodium Lactate, L-; Sodium Metabisulfite; Sodium Phosphate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Dihydrate; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Dihydrate; Sodium Phosphate, Monobasic, Monohydrate; Sodium Sulfite; Sodium Tartrate; Sorbitol; Sorbitol Solution; Soybean Oil; Stannous Chloride; Stannous Chloride Anhydrous; Sterile Water For Inhalation; Sucrose; Sulfobutylether .Beta.-Cyclodextrin; Sulfur Dioxide; Sulfuric Acid; Tartaric Acid; Tartaric Acid, Dl-; Tert-Butyl Alcohol; Tetrofosmin; Theophylline; Threonine; Trifluoroacetic Acid; Trisodium Citrate Dihydrate; Tromethamine; Tryptophan; Tyrosine; Valine
Any Delivery Route	Alcohol; Benzyl Alcohol; Citric Acid Monohydrate; Gelfoam Sponge; Hydrochloric Acid; Methylparaben; Poly(Dl-Lactic-Co-Glycolic Acid), (50:50; Poly(Dl-Lactic-Co-Glycolic Acid), Ethyl Ester Terminated, (50:50; Polyquaternium-7 (70/30 Acrylamide/Dadmac ; Propylene Glycol; Propylparaben; Sodium Chloride; Sodium Citrate ; Sodium Hydroxide; Sodium Lactate ; Sodium Phosphate, Monobasic, Monohydrate
Nasal	Acetic Acid; Alcohol, Dehydrated; Allyl .Alpha.-Ionone; Anhydrous Dextrose; Anhydrous Trisodium Citrate; Benzalkonium Chloride; Benzethonium Chloride; Benzyl Alcohol; Butylated Hydroxyanisole; Butylated Hydroxytoluene; Caffeine; Carbon Dioxide; Carboxymethylcellulose Sodium; Cellulose, Microcrystalline; Chlorobutanol; Citric Acid; Citric Acid Monohydrate; Dextrose; Dichlorodifluoromethane; Dichlorotetrafluoroethane; Edetate Disodium; Glycerin; Glycerol Ester Of Hydrogenated Rosin; Hydrochloric Acid; Hypromellose 2910 (15000 Mpa.S); Methylcelluloses; Methylparaben; Nitrogen; Norflurane; Oleic Acid; Petrolatum, White; Phenylethyl Alcohol; Polyethylene Glycol 3350; Polyethylene Glycol 400; Polyoxyl 400 Stearate; Polysorbate 20; Polysorbate 80; Potassium Phosphate, Monobasic; Potassium Sorbate; Propylene Glycol; Propylparaben; Sodium Acetate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Phosphate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Dihydrate; Sodium Phosphate, Dibasic, Dodecahydrate; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Dihydrate; Sorbitan Trioleate; Sorbitol; Sorbitol Solution; Sucralose; Sulfuric Acid; Trichloromonofluoromethane; Trisodium Citrate Dihydrate
Nerve Block	Acetic Acid; Acetone Sodium Bisulfite; Ascorbic Acid; Benzyl Alcohol; Calcium Chloride; Carbon Dioxide; Chlorobutanol; Citric Acid; Citric Acid Monohydrate; Edetate Calcium Disodium; Edetate Disodium; Hydrochloric Acid; Hydrochloric Acid, Diluted; Lactic Acid; Methylparaben; Monothioglycerol; Nitrogen; Potassium Chloride; Potassium Metabisulfite; Potassium Phosphate, Monobasic; Propylparaben; Sodium Bisulfite; Sodium Carbonate; Sodium Chlorate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Lactate; Sodium Lactate, L-; Sodium Metabisulfite; Sodium Phosphate; Sodium Phosphate, Dibasic, Heptahydrate
Ophthalmic	Acetic Acid; Alcohol; Alcohol, Dehydrated; Alginic Acid; Amerchol-Cab; Ammonium Hydroxide; Anhydrous Trisodium Citrate; Antipyrine; Benzalkonium Chloride; Benzethonium Chloride; Benzododecinium Bromide; Boric Acid; Caffeine; Calcium Chloride; Carbomer 1342; Carbomer 934p; Carbomer 940; Carbomer Homopolymer Type B (Allyl Pentaerythritol Crosslinked); Carboxymethylcellulose Sodium; Castor Oil; Cetyl Alcohol; Chlorobutanol; Chlorobutanol, Anhydrous; Cholesterol; Citric Acid; Citric Acid Monohydrate; Creatinine; Diethanolamine; Diethylhexyl Phthalate **See Cder Guidance: Limiting The Use Of Certain Phthalates As Excipients In Cder-Regulated Products; Divinylbenzene Styrene Copolymer; Edetate Disodium; Edetate Disodium Anhydrous; Edetate Sodium; Ethylene Vinyl Acetate Copolymer; Gellan Gum

	(Low Acyl); Glycerin; Glyceryl Stearate; High Density Polyethylene; Hydrocarbon Gel, Plasticized; Hydrochloric Acid; Hydrochloric Acid, Diluted; Hydroxyethyl Cellulose; Hydroxypropyl Methylcellulose 2906; Hypromellose 2910 (15000 Mpa.S); Hypromelloses; Jelene; Lanolin; Lanolin Alcohols; Lanolin Anhydrous; Lanolin Nonionic Derivatives; Lauralkonium Chloride; Lauroyl Sarcosine; Light Mineral Oil; Magnesium Chloride; Mannitol; Methylcellulose (4000 Mpa.S); Methylcelluloses; Methylparaben; Mineral Oil; Nitric Acid; Nitrogen; Nonoxynol-9; Octoxynol-40; Octylphenol Polymethylene; Petrolatum; Petrolatum, White; Phenylethyl Alcohol; Phenylmercuric Acetate; Phenylmercuric Nitrate; Phosphoric Acid; Polidronium Chloride; Poloxamer 188; Poloxamer 407; Polycarbophil; Polyethylene Glycol 300; Polyethylene Glycol 400; Polyethylene Glycol 8000; Polyoxyethylene - Polyoxypropylene 1800; Polyoxyl 35 Castor Oil; Polyoxyl 40 Hydrogenated Castor Oil; Polyoxyl 40 Stearate; Polypropylene Glycol; Polysorbate 20; Polysorbate 60; Polysorbate 80; Polyvinyl Alcohol; Potassium Acetate; Potassium Chloride; Potassium Phosphate, Monobasic; Potassium Sorbate; Povidone K29/32; Povidone K30; Povidone K90; Povidones; Propylene Glycol; Propylparaben; Soda Ash; Sodium Acetate; Sodium Bisulfate; Sodium Bisulfite; Sodium Borate; Sodium Borate Decahydrate; Sodium Carbonate; Sodium Carbonate Monohydrate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Metabisulfite; Sodium Nitrate; Sodium Phosphate; Sodium Phosphate Dihydrate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Dihydrate; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Dihydrate; Sodium Phosphate, Monobasic, Monohydrate; Sodium Sulfate; Sodium Sulfate Anhydrous; Sodium Sulfate Decahydrate; Sodium Sulfite; Sodium Thiosulfate; Sorbic Acid; Sorbitan Monolaurate; Sorbitol; Sorbitol Solution; Stabilized Oxychloro Complex; Sulfuric Acid; Thimerosal; Titanium Dioxide; Tocophersolan; Trisodium Citrate Dihydrate; Triton 720; Tromethamine; Tyloxapol; Zinc Chloride
Parenteral	Hydrochloric Acid; Mannitol; Nitrogen; Sodium Acetate; Sodium Chloride; Sodium Hydroxide
Percutaneous	Duro-Tak 87-2287; Silicone Adhesive 4102
Perfusion, Biliary	Glycerin
Perfusion, Cardiac	Hydrochloric Acid; Sodium Hydroxide
Periarticular	Diatrizoic Acid; Edetate Calcium Disodium; Iodine; Meglumine
Peridural	Citric Acid; Hydrochloric Acid; Methylparaben; Sodium Chloride; Sodium Hydroxide; Sodium Metabisulfite
Perineural	Hydrochloric Acid; Sodium Chloride; Sodium Hydroxide
Periodontal	Ethylene Vinyl Acetate Copolymer; Hydrochloric Acid; Methyl Pyrrolidone; Poloxamer 188; Poloxamer 407; Polylactide
Photopheresis	Acetic Acid; Alcohol, Dehydrated; Propylene Glycol; Sodium Acetate; Sodium Chloride; Sodium Hydroxide
Rectal	Alcohol; Alcohol, Dehydrated; Aluminum Subacetate; Anhydrous Citric Acid; Aniseed Oil; Ascorbic Acid; Ascorbyl Palmitate; Balsam Peru; Benzoic Acid; Benzyl Alcohol; Bismuth Subgallate; Butylated Hydroxyanisole; Butylated Hydroxytoluene; Butylparaben; Caramel; Carbomer 934; Carbomer 934p; Carboxypolymethylene; Cerasynt-Se; Cetyl Alcohol; Cocoa Butter; Coconut Oil, Hydrogenated; Coconut Oil/Palm Kernel Oil Glycerides, Hydrogenated; Cola Nitida Seed Extract; D&C Yellow No. 10; Dichlorodifluoromethane; Dichlorotetrafluoroethane; Dimethyldioctadecylammonium Bentonite; Edetate Calcium Disodium; Edetate Disodium; Edetic Acid; Epilactose; Ethylenediamine; Fat, Edible; Fat, Hard; Fd&C Blue No. 1; Fd&C Green No. 3; Fd&C Yellow No. 6; Flavor Fig 827118; Flavor Raspberry Pfc-8407; Fructose; Galactose; Glycerin; Glyceryl Palmitate; Glyceryl Stearate; Glyceryl Stearate/Peg Stearate; Glyceryl Stearate/Peg-40 Stearate; Glycine; Hydrocarbon; Hydrochloric Acid; Hydrogenated Palm Oil; Hypromelloses; Lactose; Lanolin; Lecithin; Light Mineral Oil; Magnesium Aluminum Silicate; Magnesium Aluminum Silicate Hydrate; Methylparaben; Nitrogen; Palm Kernel Oil; Paraffin; Petrolatum, White; Polyethylene Glycol 1000; Polyethylene Glycol 1540; Polyethylene Glycol 3350; Polyethylene Glycol 400; Polyethylene Glycol 4000; Polyethylene Glycol 6000; Polyethylene Glycol 8000; Polysorbate 60; Polysorbate 80; Potassium Acetate; Potassium Metabisulfite; Propylene Glycol; Propylparaben; Saccharin Sodium; Saccharin Sodium Anhydrous; Silicon Dioxide, Colloidal; Simethicone; Sodium Benzoate; Sodium Carbonate; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Metabisulfite; Sorbitan Monooleate; Sorbitan Sesquioleate; Sorbitol; Sorbitol Solution; Starch;

	Steareth-10; Steareth-40; Sucrose; Tagatose, D-; Tartaric Acid, Dl-; Trolamine; Tromethamine; Vegetable Oil Glyceride, Hydrogenated; Vegetable Oil, Hydrogenated; Wax, Emulsifying; White Wax; Xanthan Gum; Zinc Oxide
Respiratory (Inhalation)	Alcohol; Alcohol, Dehydrated; Apaflurane; Benzalkonium Chloride; Calcium Carbonate; Edetate Disodium; Gelatin; Glycine; Hydrochloric Acid; Lactose Monohydrate; Lysine Monohydrate; Mannitol; Norflurane; Oleic Acid; Polyethylene Glycol 1000; Povidone K25; Silicon Dioxide, Colloidal; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Lauryl Sulfate; Sulfuric Acid; Titanium Dioxide; Tromethamine; Zinc Oxide
Retrobulbar	Hydrochloric Acid; Sodium Hydroxide
Soft Tissue	Acetic Acid; Anhydrous Trisodium Citrate; Benzyl Alcohol; Carboxymethylcellulose; Carboxymethylcellulose Sodium; Citric Acid; Creatinine; Edetate Disodium; Hydrochloric Acid; Methylcelluloses; Methylparaben; Myristyl-.Gamma.-Picolinium Chloride; Phenol; Phosphoric Acid; Polyethylene Glycol 3350; Polyethylene Glycol 4000; Polysorbate 80; Propylparaben; Sodium Acetate; Sodium Bisulfite; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Phosphate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic; Sodium Phosphate, Monobasic, Anhydrous; Sodium Sulfite
Spinal	Anhydrous Dextrose; Dextrose; Hydrochloric Acid; Sodium Hydroxide
Subarachnoid	Hydrochloric Acid; Sodium Chloride; Sodium Hydroxide
Subconjunctival	Benzyl Alcohol; Hydrochloric Acid; Sodium Hydroxide
Subcutaneous	Acetic Acid; Acetic Acid, Glacial; Albumin Human; Ammonium Hydroxide; Ascorbic Acid; Benzyl Alcohol; Calcium Chloride; Carboxymethylcellulose Sodium; Chlorobutanol; Cresol; Diatrizoic Acid; Dimethyl Sulfoxide; Edetate Calcium Disodium; Edetate Disodium; Ethylene Vinyl Acetate Copolymer; Glycerin; Glycine; Glycine Hydrochloride; Histidine; Hydrochloric Acid; Lactic Acid; Lactic Acid, L-; Lactose; Magnesium Chloride; Magnesium Stearate; Mannitol; Metacresol; Methanesulfonic Acid; Methionine; Methyl Pyrrolidone; Methylparaben; Nitrogen; Phenol; Phenol, Liquefied; Phosphoric Acid; Poloxamer 188; Polyethylene Glycol 3350; Polyglactin; Polysorbate 20; Polysorbate 80; Potassium Phosphate, Dibasic; Potassium Phosphate, Monobasic; Povidone K17; Povidones; Propylene Glycol; Propylparaben; Protamine Sulfate; Sodium Acetate; Sodium Acetate Anhydrous; Sodium Bicarbonate; Sodium Bisulfite; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sodium Metabisulfite; Sodium Phosphate; Sodium Phosphate Dihydrate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Dihydrate; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Dihydrate; Sodium Phosphate, Monobasic, Monohydrate; Sodium Sulfite; Sodium Thioglycolate; Stearic Acid; Sucrose; Thimerosal; Tromethamine; Zinc; Zinc Acetate; Zinc Carbonate; Zinc Chloride; Zinc Oxide
Sublingual	Alcohol, Dehydrated
Submucosal	Acetic Acid; Edetic Acid; Mannitol; Nitrogen; Sodium Acetate; Sodium Chloride; Sodium Hydroxide; Sodium Metabisulfite
Topical	.Alpha.-Terpineol; .Alpha.-Tocopherol; .Alpha.-Tocopherol Acetate, Dl-; .Alpha.-Tocopherol, Dl-; 1,2,6-Hexanetriol; 1-O-Tolylbiguanide; 2-Ethyl-1,6-Hexanediol; Acetic Acid; Acetone; Acetylated Lanolin Alcohols; Acrylates Copolymer; Adhesive Tape; Alcohol; Alcohol, Dehydrated; Alcohol, Denatured; Alcohol, Diluted; Alkyl Ammonium Sulfonic Acid Betaine; Alkyl Aryl Sodium Sulfonate; Allantoin; Almond Oil; Aluminum Acetate; Aluminum Chlorhydroxy Allantoinate; Aluminum Hydroxide; Aluminum Hydroxide - Sucrose, Hydrated; Aluminum Hydroxide Gel; Aluminum Hydroxide Gel F 500; Aluminum Hydroxide Gel F 5000; Aluminum Monostearate; Aluminum Oxide; Aluminum Silicate; Aluminum Starch Octenylsuccinate; Aluminum Stearate; Aluminum Sulfate Anhydrous; Amerchol C; Amerchol-Cab; Aminomethylpropanol; Ammonia Solution; Ammonia Solution, Strong; Ammonium Hydroxide; Ammonium Lauryl Sulfate; Ammonium Nonoxynol-4 Sulfate; Ammonium Salt Of C-12-C-15 Linear Primary Alcohol Ethoxylate; Ammonyx; Amphoteric-2; Amphoteric-9; Anhydrous Citric Acid; Anhydrous Trisodium Citrate; Anoxid Sbn; Antifoam; Apricot Kernel Oil Peg-6 Esters; Aquaphor; Arlacel; Ascorbic Acid; Ascorbyl Palmitate; Beeswax; Beeswax, Synthetic; Beheneth-10; Bentonite; Benzalkonium Chloride; Benzoic Acid; Benzyl Alcohol; Betadex; Boric Acid; Butane; Butyl Alcohol; Butyl Ester Of Vinyl Methyl Ether/Maleic Anhydride Copolymer (125000 Mw); Butyl Stearate; Butylated Hydroxyanisole; Butylated Hydroxytoluene; Butylene Glycol; Butylparaben; C20-40 Pareth-24; Calcium Chloride; Calcium Hydroxide; Canada

	Balsam; Caprylic/Capric Triglyceride; Caprylic/Capric/Stearic Triglyceride; Captan; Caramel; Carbomer 1342; Carbomer 1382; Carbomer 934; Carbomer 934p; Carbomer 940; Carbomer 941; Carbomer 980; Carbomer 981; Carbomer Homopolymer Type B (Allyl Pentaerythritol Crosslinked); Carbomer Homopolymer Type C (Allyl Pentaerythritol Crosslinked); Carboxy Vinyl Copolymer; Carboxymethylcellulose; Carboxymethylcellulose Sodium; Carboxypolymethylene; Carrageenan; Carrageenan Salt; Castor Oil; Cedar Leaf Oil; Cellulose; Cerasynt-Se; Ceresin; Ceteareth-12; Ceteareth-15; Ceteareth-30; Cetearyl Alcohol/Ceteareth-20; Cetearyl Ethylhexanoate; Ceteth-10; Ceteth-2; Ceteth-20; Ceteth-23; Cetostearyl Alcohol; Cetrimonium Chloride; Cetyl Alcohol; Cetyl Esters Wax; Cetyl Palmitate; Chlorobutanol; Chlorocresol; Chloroxylenol; Cholesterol; Choleth-24; Citric Acid; Citric Acid Monohydrate; Cocamide Ether Sulfate; Cocamine Oxide; Coco Betaine; Coco Diethanolamide; Coco Monoethanolamide; Cocoa Butter; Coco-Glycerides; Coconut Oil; Cocoyl Caprylocaprate; Collagen; Coloring Suspension; Cream Base; Creatinine; Crospovidone; Cyclomethicone; Cyclomethicone/Dimethicone Copolyol; D&C Red No. 28; D&C Red No. 33; D&C Red No. 36; D&C Red No. 39; D&C Yellow No. 10; Decyl Methyl Sulfoxide; Dehydag Wax Sx; Dehydroacetic Acid; Dehymuls E; Denatonium Benzoate; Dextrin; Diazolidinyl Urea; Dichlorobenzyl Alcohol; Dichlorodifluoromethane; Dichlorotetrafluoroethane; Diethanolamine; Diethyl Sebacate; Diethylene Glycol Monoethyl Ether; Dihydroxyaluminum Aminoacetate; Diisopropanolamine; Diisopropyl Adipate; Diisopropyl Dilinoleate; Dimethicone 350; Dimethicone Copolyol; Dimethicone Medical Fluid 360; Dimethyl Isosorbide; Dimethyl Sulfoxide; Dinoseb Ammonium Salt; Disodium Cocoamphodiacetate; Disodium Laureth Sulfosuccinate; Disodium Lauryl Sulfosuccinate; Dmdm Hydantoin; Docosanol; Docusate Sodium; Edetate Disodium; Edetate Sodium; Edetic Acid; Entsufon; Entsufon Sodium; Epitetracycline Hydrochloride; Essence Bouquet 9200; Ethyl Acetate; Ethylcelluloses; Ethylene Glycol; Ethylenediamine; Ethylenediamine Dihydrochloride; Ethylhexyl Hydroxystearate; Ethylparaben; Fatty Acid Pentaerythriol Ester; Fatty Acids; Fatty Alcohol Citrate; Fd&C Blue No. 1; Fd&C Red No. 4; Fd&C Red No. 40; Fd&C Yellow No. 10 (Delisted); Fd&C Yellow No. 5; Fd&C Yellow No. 6; Ferric Oxide; Flavor Rhodia Pharmaceutical No. Rf 451; Formaldehyde; Formaldehyde Solution; Fractionated Coconut Oil; Fragrance 3949-5; Fragrance 520a; Fragrance 6.007; Fragrance 91-122; Fragrance 9128-Y; Fragrance 93498g; Fragrance Balsam Pine No. 5124; Fragrance Bouquet 10328; Fragrance Chemoderm 6401-B; Fragrance Chemoderm 6411; Fragrance Cream No. 73457; Fragrance Cs-28197; Fragrance Felton 066m; Fragrance Firmenich 47373; Fragrance Givaudan Ess 9090/1c; Fragrance H-6540; Fragrance Herbal 10396; Fragrance Nj-1085; Fragrance P O Fl-147; Fragrance Pa 52805; Fragrance Pera Derm D; Fragrance Rbd-9819; Fragrance Shaw Mudge U-7776; Fragrance Tf 044078; Fragrance Ungerer Honeysuckle K 2771; Fragrance Ungerer N5195; Gelatin; Gluconolactone; Glycerin; Glyceryl Citrate; Glyceryl Isostearate; Glyceryl Monostearate; Glyceryl Oleate; Glyceryl Oleate/Propylene Glycol; Glyceryl Palmitate; Glyceryl Ricinoleate; Glyceryl Stearate; Glyceryl Stearate - Laureth-23; Glyceryl Stearate/Peg-100 Stearate; Glyceryl Stearate-Stearamidoethyl Diethylamine; Glycol Distearate; Glycol Stearate; Guar Gum; Hair Conditioner (18n195-1m); Hexylene Glycol; High Density Polyethylene; Hyaluronate Sodium; Hydrocarbon Gel, Plasticized; Hydrochloric Acid; Hydrochloric Acid, Diluted; Hydrogen Peroxide; Hydrogenated Castor Oil; Hydrogenated Palm/Palm Kernel Oil Peg-6 Esters; Hydroxyethyl Cellulose; Hydroxymethyl Cellulose; Hydroxyoctacosanyl Hydroxystearate; Hydroxypropyl Cellulose; Hypromelloses; Imidurea; Irish Moss Extract; Isobutane; Isoceteth-20; Isooctyl Acrylate; Isopropyl Alcohol; Isopropyl Isostearate; Isopropyl Myristate; Isopropyl Myristate - Myristyl Alcohol; Isopropyl Palmitate; Isopropyl Stearate; Isostearic Acid; Isostearyl Alcohol; Jelene; Kaolin; Kathon Cg; Kathon Cg Ii; Lactate; Lactic Acid; Lactic Acid, Dl-; Laneth; Lanolin; Lanolin Alcohol - Mineral Oil; Lanolin Alcohols; Lanolin Anhydrous; Lanolin Cholesterols; Lanolin, Ethoxylated; Lanolin, Hydrogenated; Lauramine Oxide; Laurdimonium Hydrolyzed Animal Collagen; Laureth Sulfate; Laureth-2; Laureth-23; Laureth-4; Lauric Diethanolamide; Lauric Myristic Diethanolamide; Lauryl Sulfate; Lavandula Angustifolia Flowering Top; Lecithin; Lecithin Unbleached; Lemon Oil; Light Mineral Oil; Light Mineral Oil (85 Ssu); Limonene, (+/-)-; Lipocol Sc-15; Magnesium Aluminum Silicate; Magnesium Aluminum Silicate Hydrate; Magnesium Nitrate; Magnesium Stearate; Mannitol; Maprofix; Medical Antiform A-F Emulsion; Menthol; Methyl Gluceth-10; Methyl Gluceth-20; Methyl Gluceth-20 Sesquistearate; Methyl Glucose Sesquistearate; Methyl Salicylate; Methyl Stearate; Methylcelluloses; Methylchloroisothiazolinone; Methylisothiazolinone; Methylparaben; Microcrystalline Wax;

	Mineral Oil; Mono And Diglyceride; Monostearyl Citrate; Multisterol Extract; Myristyl Alcohol; Myristyl Lactate; Niacinamide; Nitric Acid; Nitrogen; Nonoxynol Iodine; Nonoxynol-15; Nonoxynol-9; Oatmeal; Octadecene-1/Maleic Acid Copolymer; Octoxynol-1; Octoxynol-9; Octyldodecanol; Oleic Acid; Oleth-10/Oleth-5; Oleth-2; Oleth-20; Oleyl Alcohol; Oleyl Oleate; Olive Oil; Palmitamine Oxide; Parabens; Paraffin; Paraffin, White Soft; Parfum Creme 45/3; Peanut Oil; Peanut Oil, Refined; Pectin; Peg 6-32 Stearate/Glycol Stearate; Peg-100 Stearate; Peg-12 Glyceryl Laurate; Peg-120 Glyceryl Stearate; Peg-120 Methyl Glucose Dioleate; Peg-15 Cocamine; Peg-150 Distearate; Peg-2 Stearate; Peg-22 Methyl Ether/Dodecyl Glycol Copolymer; Peg-25 Propylene Glycol Stearate; Peg-4 Dilaurate; Peg-4 Laurate; Peg-45/Dodecyl Glycol Copolymer; Peg-5 Oleate; Peg-50 Stearate; Peg-54 Hydrogenated Castor Oil; Peg-6 Isostearate; Peg-60 Hydrogenated Castor Oil; Peg-7 Methyl Ether; Peg-75 Lanolin; Peg-8 Laurate; Peg-8 Stearate; Pegoxol 7 Stearate; Pentaerythritol Cocoate; Peppermint Oil; Perfume 25677; Perfume Bouquet; Perfume E-1991; Perfume Gd 5604; Perfume Tana 90/42 Scba; Perfume W-1952-1; Petrolatum; Petrolatum, White; Petroleum Distillates; Phenonip; Phenoxyethanol; Phenylmercuric Acetate; Phosphoric Acid; Pine Needle Oil (Pinus Sylvestris); Plastibase-50w; Polidronium Chloride; Poloxamer 124; Poloxamer 181; Poloxamer 182; Poloxamer 188; Poloxamer 237; Poloxamer 407; Polycarbophil; Polyethylene Glycol 1000; Polyethylene Glycol 1450; Polyethylene Glycol 1500; Polyethylene Glycol 1540; Polyethylene Glycol 200; Polyethylene Glycol 300; Polyethylene Glycol 300-1600; Polyethylene Glycol 3350; Polyethylene Glycol 400; Polyethylene Glycol 4000; Polyethylene Glycol 540; Polyethylene Glycol 600; Polyethylene Glycol 6000; Polyethylene Glycol 8000; Polyethylene Glycol 900; Polyhydroxyethyl Methacrylate; Polyisobutylene; Polyisobutylene (1100000 Mw); Polyoxyethylene - Polyoxypropylene 1800; Polyoxyethylene Alcohols; Polyoxyethylene Fatty Acid Esters; Polyoxyethylene Propylene; Polyoxyl 20 Cetostearyl Ether; Polyoxyl 40 Hydrogenated Castor Oil; Polyoxyl 40 Stearate; Polyoxyl 400 Stearate; Polyoxyl 6 And Polyoxyl 32 Palmitostearate; Polyoxyl Distearate; Polyoxyl Glyceryl Stearate; Polyoxyl Lanolin; Polyoxyl Stearate; Polypropylene; Polyquaternium-10; Polysorbate 20; Polysorbate 40; Polysorbate 60; Polysorbate 65; Polysorbate 80; Polyvinyl Alcohol; Potash; Potassium Citrate; Potassium Hydroxide; Potassium Soap; Potassium Sorbate; Povidone Acrylate Copolymer; Povidone Hydrogel; Povidone K90; Povidone/Eicosene Copolymer; Povidones; Ppg-12/Smdi Copolymer; Ppg-15 Stearyl Ether; Ppg-20 Methyl Glucose Ether Distearate; Ppg-26 Oleate; Product Wat; Promulgen D; Promulgen G; Propane; Propellant A-46; Propyl Gallate; Propylene Carbonate; Propylene Glycol; Propylene Glycol Diacetate; Propylene Glycol Dicaprylate; Propylene Glycol Monopalmitostearate; Propylene Glycol Palmitostearate; Propylene Glycol Ricinoleate; Propylene Glycol/Diazolidinyl Urea/Methylparaben/Propylparben; Propylparaben; Protein Hydrolysate; Quaternium-15; Quaternium-15 Cis-Form; Quaternium-52; Saccharin; Saccharin Sodium; Safflower Oil; Sd Alcohol 3a; Sd Alcohol 40; Sd Alcohol 40-2; Sd Alcohol 40b; Sepineo P 600; Shea Butter; Silicon; Silicon Dioxide; Silicone; Silicone Adhesive Bio-Psa Q7-4201; Silicone Adhesive Bio-Psa Q7-4301; Silicone Emulsion; Simethicone; Simethicone Emulsion; Sipon Ls 20np; Sodium Acetate; Sodium Acetate Anhydrous; Sodium Alkyl Sulfate; Sodium Benzoate; Sodium Bisulfite; Sodium Borate; Sodium Cetostearyl Sulfate; Sodium Chloride; Sodium Citrate; Sodium Cocoyl Sarcosinate; Sodium Dodecylbenzenesulfonate; Sodium Formaldehyde Sulfoxylate; Sodium Hydroxide; Sodium Iodide; Sodium Lactate; Sodium Laureth-2 Sulfate; Sodium Laureth-3 Sulfate; Sodium Laureth-5 Sulfate; Sodium Lauroyl Sarcosinate; Sodium Lauryl Sulfate; Sodium Lauryl Sulfoacetate; Sodium Metabisulfite; Sodium Phosphate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Dihydrate; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate, Monobasic, Dihydrate; Sodium Phosphate, Monobasic, Monohydrate; Sodium Polyacrylate (2500000 Mw); Sodium Pyrrolidone Carboxylate; Sodium Sulfite; Sodium Sulfosuccinated Undecyclenic Monoalkylolamide; Sodium Thiosulfate; Sodium Xylenesulfonate; Somay 44; Sorbic Acid; Sorbitan; Sorbitan Isostearate; Sorbitan Monolaurate; Sorbitan Monooleate; Sorbitan Monopalmitate; Sorbitan Monostearate; Sorbitan Sesquioleate; Sorbitan Tristearate; Sorbitol; Sorbitol Solution; Soybean Flour; Soybean Oil; Spearmint Oil; Spermaceti; Squalane; Starch; Stearalkonium Chloride; Stearamidoethyl Diethylamine; Steareth-10; Steareth-100; Steareth-2; Steareth-20; Steareth-21; Steareth-40; Stearic Acid; Stearic Diethanolamide; Stearoxytrimethylsilane; Steartrimonium Hydrolyzed Animal Collagen; Stearyl Alcohol; Styrene/Isoprene/Styrene Block Copolymer; Sucrose; Sucrose Distearate; Sucrose Polyesters;

	Sulfacetamide Sodium; Sulfuric Acid; Surfactol Qs; Talc; Tall Oil; Tallow Glycerides; Tartaric Acid; Tenox; Tenox-2; Tert-Butyl Alcohol; Tert-Butyl Hydroperoxide; Thimerosal; Titanium Dioxide; Tocopherol; Tocophersolan; Trichloromonofluoromethane; Trideceth-10; Triethanolamine Lauryl Sulfate; Triglycerides, Medium Chain; Trihydroxystearin; Trilaneth-4 Phosphate; Trilaureth-4 Phosphate; Trisodium Citrate Dihydrate; Trisodium Hedta; Triton X-200; Trolamine; Tromethamine; Tyloxapol; Undecylenic Acid; Vegetable Oil; Vegetable Oil, Hydrogenated; Viscarin; Vitamin E; Wax, Emulsifying; Wecobee Fs; White Wax; Xanthan Gum; Zinc Acetate
Transdermal	Acrylates Copolymer; Acrylic Acid-Isooctyl Acrylate Copolymer; Acrylic Adhesive 788; Adcote 72a103; Aerotex Resin 3730; Alcohol; Alcohol, Dehydrated; Aluminum Polyester; Bentonite; Butylated Hydroxytoluene; Butylene Glycol; Butyric Acid; Caprylic/Capric Triglyceride; Carbomer 1342; Carbomer 940; Carbomer 980; Carrageenan; Cetylpyridinium Chloride; Citric Acid; Crospovidone; Daubert 1-5 Pestr (Matte) 164z; Diethylene Glycol Monoethyl Ether; Diethylhexyl Phthalate **See Cder Guidance: Limiting The Use Of Certain Phthalates As Excipients In Cder-Regulated Products; Dimethicone Copolyol; Dimethicone Mdx4-4210; Dimethicone Medical Fluid 360; Dimethylaminoethyl Methacrylate - Butyl Methacrylate - Methyl Methacrylate Copolymer; Dipropylene Glycol; Duro-Tak 280-2516; Duro-Tak 387-2516; Duro-Tak 80-1196; Duro-Tak 87-2070; Duro-Tak 87-2194; Duro-Tak 87-2287; Duro-Tak 87-2296; Duro-Tak 87-2888; Duro-Tak 87-2979; Edetate Disodium; Ethyl Acetate; Ethyl Oleate; Ethylcelluloses; Ethylene Vinyl Acetate Copolymer; Ethylene-Propylene Copolymer; Fatty Acid Esters; Gelva 737; Glycerin; Glyceryl Laurate; Glyceryl Oleate; Heptane; High Density Polyethylene; Hydrochloric Acid; Hydrogenated Polybutene 635-690; Hydroxyethyl Cellulose; Hydroxypropyl Cellulose; Isopropyl Myristate; Isopropyl Palmitate; Lactose; Lanolin Anhydrous; Lauryl Lactate; Lecithin; Levulinic Acid; Light Mineral Oil; Medical Adhesive Modified S-15; Methyl Alcohol; Methyl Laurate; Mineral Oil; Nitrogen; Octisalate; Octyldodecanol; Oleic Acid; Oleyl Alcohol; Oleyl Oleate; Pentadecalactone; Petrolatum, White; Polacrilin; Polyacrylic Acid (250000 Mw); Polybutene (1400 Mw); Polyester; Polyester Polyamine Copolymer; Polyester Rayon; Polyethylene Terephthalates; Polyisobutylene; Polyisobutylene (1100000 Mw); Polyisobutylene (35000 Mw); Polyisobutylene 178-236; Polyisobutylene 241-294; Polyisobutylene 35-39; Polyisobutylene Low Molecular Weight; Polyisobutylene Medium Molecular Weight; Polyisobutylene/Polybutene Adhesive; Polypropylene; Polyvinyl Acetate; Polyvinyl Alcohol; Polyvinyl Chloride; Polyvinyl Chloride-Polyvinyl Acetate Copolymer; Polyvinylpyridine; Povidone K29/32; Povidones; Propylene Glycol; Propylene Glycol Monolaurate; Ra-2397; Ra-3011; Silicon; Silicon Dioxide, Colloidal; Silicone; Silicone Adhesive 4102; Silicone Adhesive 4502; Silicone Adhesive Bio-Psa Q7-4201; Silicone Adhesive Bio-Psa Q7-4301; Silicone/Polyester Film Strip; Sodium Chloride; Sodium Citrate; Sodium Hydroxide; Sorbitan Monooleate; Stearalkonium Hectorite/Propylene Carbonate; Titanium Dioxide; Triacetin; Trolamine; Tromethamine; Union 76 Amsco-Res 6038; Viscose/Cotton
Transmucosal	Magnesium Stearate; Mannitol; Potassium Bicarbonate; Sodium Starch Glycolate
Ureteral	Benzyl Alcohol; Diatrizoic Acid; Edetate Calcium Disodium; Edetate Disodium; Hydrochloric Acid; Meglumine; Methylparaben; Propylparaben; Sodium Citrate; Sodium Hydroxide
Urethral	Diatrizoic Acid; Edetate Calcium Disodium; Edetate Disodium; Hydrochloric Acid; Meglumine; Methylparaben; Polyethylene Glycol 1450; Propylparaben; Sodium Hydroxide; Sodium Phosphate, Dibasic, Heptahydrate; Tromethamine
Vaginal	Adipic Acid; Alcohol, Denatured; Allantoin; Anhydrous Lactose; Apricot Kernel Oil Peg-6 Esters; Barium Sulfate; Beeswax; Bentonite; Benzoic Acid; Benzyl Alcohol; Butylated Hydroxyanisole; Butylated Hydroxytoluene; Calcium Lactate; Carbomer 934; Carbomer 934p; Cellulose, Microcrystalline; Ceteth-20; Cetostearyl Alcohol; Cetyl Alcohol; Cetyl Esters Wax; Cetyl Palmitate; Cholesterol; Choleth; Citric Acid; Citric Acid Monohydrate; Coconut Oil/Palm Kernel Oil Glycerides, Hydrogenated; Crospovidone; Edetate Disodium; Ethylcelluloses; Ethylene-Vinyl Acetate Copolymer (28% Vinyl Acetate); Ethylene-Vinyl Acetate Copolymer (9% Vinylacetate); Fatty Alcohols; Fd&C Yellow No. 5; Gelatin; Glutamic Acid, Dl-; Glycerin; Glyceryl Isostearate; Glyceryl Monostearate; Glyceryl Stearate; Guar Gum; High Density Polyethylene; Hydrogel Polymer; Hydrogenated Palm Oil; Hypromellose 2208 (15000 Mpa.S); Hypromelloses; Isopropyl Myristate; Lactic Acid; Lactic Acid, Dl-; Lactose; Lactose Monohydrate; Lactose, Hydrous; Lanolin; Lanolin Anhydrous; Lecithin; Lecithin, Soybean; Light Mineral Oil; Magnesium Aluminum Silicate; Magnesium Aluminum Silicate Hydrate; Magnesium Stearate; Methyl Stearate; Methylparaben; Microcrystalline Wax; Mineral Oil; Nitric Acid;

	Octyldodecanol; Peanut Oil; Peg 6-32 Stearate/Glycol Stearate; Peg-100 Stearate; Peg-120 Glyceryl Stearate; Peg-2 Stearate; Peg-5 Oleate; Pegoxol 7 Stearate; Petrolatum, White; Phenylmercuric Acetate; Phospholipon 90g; Phosphoric Acid; Piperazine Hexahydrate; Poly(Dimethylsiloxane/Methylvinylsiloxane/Methylhydrogensiloxane) Dimethylvinyl Or Dimethylhydroxy Or Trimethyl Endblocked; Polycarbophil; Polyester; Polyethylene Glycol 1000; Polyethylene Glycol 3350; Polyethylene Glycol 400; Polyethylene Glycol 4000; Polyethylene Glycol 6000; Polyethylene Glycol 8000; Polyglyceryl-3 Oleate; Polyglyceryl-4 Oleate; Polyoxyl Palmitate; Polysorbate 20; Polysorbate 60; Polysorbate 80; Polyurethane; Potassium Alum; Potassium Hydroxide; Povidone K29/32; Povidones; Promulgen D; Propylene Glycol; Propylene Glycol Monopalmitostearate; Propylparaben; Quaternium-15 Cis-Form; Silicon Dioxide; Silicon Dioxide, Colloidal; Silicone; Sodium Bicarbonate; Sodium Citrate; Sodium Hydroxide; Sodium Lauryl Sulfate; Sodium Metabisulfite; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Monobasic, Anhydrous; Sorbic Acid; Sorbitan Monostearate; Sorbitol; Sorbitol Solution; Spermaceti; Stannous 2-Ethylhexanoate; Starch; Starch 1500, Pregelatinized; Starch, Corn; Stearamidoethyl Diethylamine; Stearic Acid; Stearyl Alcohol; Tartaric Acid, Dl-; Tert-Butylhydroquinone; Tetrapropyl Orthosilicate; Trolamine; Urea; Vegetable Oil, Hydrogenated; Wecobee Fs; White Ceresin Wax; White Wax

[0631] Non-limiting routes of administration for the NAVs of the present invention are described below.

Parenteral and Injectable Administration

[0632] Liquid dosage forms for parenteral administration include, but are not limited to, pharmaceutically acceptable emulsions, microemulsions, solutions, suspensions, syrups, and/or elixirs. In addition to active ingredients, liquid dosage forms may comprise inert diluents commonly used in the art such as, for example, wafer or other solvents, solubilizing agents and emulsifiers such as ethyl alcohol, isopropyl alcohol, ethyl carbonate, ethyl acetate, benzyl alcohol, benzyl benzoate, propylene glycol, 1 ,3-butylene glycol, dimethylformamide, oils (in particular, cottonseed, groundnut, corn, germ, olive, castor, and sesame oils), glycerol, tetrahydrofurfuryl alcohol, polyethylene glycols and fatty acid esters of sorbitan, and mixtures thereof. Besides inert diluents, oral compositions can include adjuvants such as wetting agents, emulsifying and suspending agents, sweetening, flavoring, and/or perfuming agents. In certain embodiments for parenteral administration, compositions are mixed with solubilizing agents such as CREMOPHOR®, alcohols, oils, modified oils, glycols, polysorbates, cyclodextrins, polymers, and/or combinations thereof.

[0633] A pharmaceutical composition for parenteral administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for parenteral administration includes hydrochloric acid, mannitol, nitrogen, sodium acetate, sodium chloride and sodium hydroxide.

[0634] Injectable preparations, for example, sterile injectable aqueous or oleaginous suspensions may be formulated according to the known art using suitable dispersing agents, wetting agents, and/or suspending agents. Sterile injectable preparations may be sterile injectable solutions, suspensions, and/or emulsions in nontoxic parenterally acceptable diluents and/or solvents, for example, as a solution in 1,3-butanediol. Among the acceptable vehicles and solvents that may be employed are water, Ringer's solution, U.S.P., and isotonic sodium chloride solution. Sterile, fixed oils are conventionally employed as a solvent or suspending medium. For this purpose any bland fixed oil can be employed including synthetic mono- or diglycerides. Fatty acids such as oleic acid can be used in the preparation of injectables. The sterile formulation may also comprise adjuvants such as local anesthetics, preservatives and buffering agents.

[0635] Injectable formulations can be sterilized, for example, by filtration through a bacterial-retaining filter, and/or by incorporating sterilizing agents in the form of sterile solid compositions which can be dissolved or dispersed in sterile wafer or other sterile injectable medium prior to use.

[0636] Injectable formulations may be for direct injection into a region of a tissue, organ and/or subject. As a non-limiting example, a tissue, organ and/or subject may be directly injected a formulation by intramyocardial injection into the ischemic region. (See e.g., Zangi et al. Nature Biotechnology 2013).

[0637] In order to prolong the effect of an active ingredient, it is often desirable to slow the absorption of the active ingredient from subcutaneous or intramuscular injection. This may be accomplished by the use of a liquid suspension of crystalline or amorphous material with poor wafer solubility. The rate of absorption of the drug then depends upon its rate of dissolution which, in turn, may depend upon crystal size and crystalline form. Alternatively, delayed absorption of a parenterally administered drug form is accomplished by dissolving or suspending the drug in an oil vehicle. Injectable depot forms are made by forming microencapsule matrices of the drug in biodegradable polymers such as polylactide-polyglycolide. Depending upon the ratio of drug to polymer and the nature of the particular polymer employed, the rate

of drug release can be controlled. Examples of other biodegradable polymers include poly(orthoesters) and poly(anhydrides). Depot injectable formulations are prepared by entrapping the drug in liposomes or microemulsions which are compatible with body tissues.

Rectal and Vaginal Administration

[0638] Compositions for rectal or vaginal (e.g., transvaginal) administration are typically suppositories which can be prepared by mixing compositions with suitable non-irritating excipients such as cocoa buffer, polyethylene glycol or a suppository wax which are solid at ambient temperature but liquid at body temperature and therefore melt in the rectum or vaginal cavity and release the active ingredient.

[0639] As a non-limiting example, the formulations for rectal and/or vaginal administration may be prepared by mixing the drug with a suitable non-irritating excipient that is solid at ordinary temperatures but liquid at the rectal temperature and will therefore melt in the rectum and/or vagina to release the drug. Such materials include cocoa buffer and polyethylene glycols.

[0640] A pharmaceutical composition for rectal administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for rectal administration includes alcohol, alcohol, dehydrated, aluminum subacetate, anhydrous citric acid, aniseed oil, ascorbic acid, ascorbyl palmitate, balsam peru, benzoic acid, benzyl alcohol, bismuth subgallate, butylated hydroxyanisole, butylated hydroxytoluene, butylparaben, caramel, carbomer 934, carbomer 934p, carboxypolymethylene, cerasynt-se, cetyl alcohol, cocoa buffer, coconut oil, hydrogenated, coconut oil/palm kernel oil glycerides, hydrogenated, cola nitida seed extract, d&c yellow no. 10, dichlorodifluoromethane, dichlorotetrafluoroethane, dimethyldioctadecylammonium bentonite, edetate calcium disodium, edetate disodium, edetic acid, epilactose, ethylenediamine, fat, edible, fat, hard, fd&c blue no. 1, fd&c green no. 3, fd&c yellow no. 6, flavor fig 827118, flavor raspberry pfc-8407, fructose, galactose, glycerin, glyceryl palmitate, glyceryl stearate, glyceryl stearate/peg stearate, glyceryl stearate/peg-40 stearate, glycine, hydrocarbon, hydrochloric acid, hydrogenated palm oil, hypromelloses, lactose, lanolin, lecithin, light mineral oil, magnesium aluminum silicate, magnesium aluminum silicate hydrate, methylparaben, nitrogen, palm kernel oil, paraffin, petrolatum, white, polyethylene glycol 1000, polyethylene glycol 1540, polyethylene glycol 3350, polyethylene glycol 400, polyethylene glycol 4000, polyethylene glycol 6000, polyethylene glycol 8000, polysorbate 60, polysorbate 80, potassium acetate, potassium metabisulfite, propylene glycol, propylparaben, saccharin sodium, saccharin sodium anhydrous, silicon dioxide, colloidal, simethicone, sodium benzoate, sodium carbonate, sodium chloride, sodium citrate, sodium hydroxide, sodium metabisulfite, sorbitan monooleate, sorbitan sesquioleate, sorbitol, sorbitol solution, starch, steareth-10, steareth-40, sucrose, tagatose, d-, tartaric acid, dl-, trolamine, tromethamine, vegetable oil glyceride, hydrogenated, vegetable oil, hydrogenated, wax, emulsifying, white wax, xanthan gum and zinc oxide.

[0641] A pharmaceutical composition for vaginal administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for vaginal administration includes adipic acid, alcohol, denatured, allantoin, anhydrous lactose, apricot kernel oil peg-6 esters, barium sulfate, beeswax, bentonite, benzoic acid, benzyl alcohol, butylated hydroxyanisole, butylated hydroxytoluene, calcium lactate, carbomer 934, carbomer 934p, cellulose, microcrystalline, ceteth-20, cetostearyl alcohol, cetyl alcohol, cetyl esters wax, cetyl palmitate, cholesterol, choleth, citric acid, citric acid monohydrate, coconut oil/palm kernel oil glycerides, hydrogenated, crospovidone, edetate disodium, ethylcelluloses, ethylene-vinyl acetate copolymer (28% vinyl acetate), ethylene-vinyl acetate copolymer (9% vinyl acetate), fatty alcohols, fd&c yellow no. 5, gelatin, glutamic acid, dl-, glycerin, glyceryl isostearate, glyceryl monostearate, glyceryl stearate, guar gum, high density polyethylene, hydrogel polymer, hydrogenated palm oil, hypromellose 2208 (15000 mpa.s), hypromelloses, isopropyl myristate, lactic acid, lactic acid, dl-, lactose, lactose monohydrate, lactose, hydrous, lanolin, lanolin anhydrous, lecithin, lecithin, soybean, light mineral oil, magnesium aluminum silicate, magnesium aluminum silicate hydrate, magnesium stearate, methyl stearate, methylparaben, microcrystalline wax, mineral oil, nitric acid, octyldodecanol, peanut oil, peg 6-32 stearate/glycol stearate, peg-100 stearate, peg-120 glyceryl stearate, peg-2 stearate, peg-5 oleate, pegoxol 7 stearate, petrolatum, white, phenylmercuric acetate, phospholipon 90g, phosphoric acid, piperazine hexahydrate, poly(dimethylsiloxane/methylvinylsiloxane/methylhydrogensiloxane) dimethylvinyl or dimethylhydroxy or trimethyl endblocked, polycarbophil, polyester, polyethylene glycol 1000, polyethylene glycol 3350, polyethylene glycol 400, polyethylene glycol 4000, polyethylene glycol 6000, polyethylene glycol 8000, polyglyceryl-3 oleate, polyglyceryl-4 oleate, polyoxyl palmitate, polysorbate 20, polysorbate 60, polysorbate 80, polyurethane, potassium alum, potassium hydroxide, povidone k29/32, povidones, promulgen d, propylene glycol, propylene glycol monopalmitostearate, propylparaben, quaternium-15 cis-form, silicon dioxide, silicon dioxide, colloidal, silicone, sodium bicarbonate, sodium citrate, sodium hydroxide, sodium lauryl sulfate, sodium metabisulfite, sodium phosphate, dibasic, anhydrous, sodium phosphate, monobasic, anhydrous, sorbic acid, sorbitan monostearate, sorbitol, sorbitol solution, spermaceti, stannous 2-ethylhexanoate, starch, starch 1500, pregelatinized, starch, corn,

stearamidoethyl diethylamine, stearic acid, stearyl alcohol, tartaric acid, dl-, tert-butylhydroquinone, tetrapropyl orthosilicate, trolamine, urea, vegetable oil, hydrogenated, wecobee fs, white ceresin wax and white wax

Oral Administration

[0642] Liquid dosage forms for oral administration include, but are not limited to, pharmaceutically acceptable emulsions, microemulsions, solutions, suspensions, syrups, and/or elixirs. In addition to active ingredients, liquid dosage forms may comprise inert diluents and/or excipients commonly used in the art such as, for example, wafer or other solvents, solubilizing agents and emulsifiers such as ethyl alcohol, isopropyl alcohol, ethyl carbonate, ethyl acetate, benzyl alcohol, benzyl benzoate, propylene glycol, 1 ,3-butylene glycol, dimethylformamide, oils (in particular, cottonseed, groundnut, corn, germ, olive, castor, and sesame oils), glycerol, tetrahydrofurfuryl alcohol, polyethylene glycols and fatty acid esters of sorbitan, and mixtures thereof. Besides inert diluents, oral compositions can include adjuvants such as wetting agents, emulsifying and suspending agents, sweetening, flavoring, and/or perfuming agents. In certain embodiments for parenteral administration, compositions are mixed with solubilizing agents such as CREMOPHOR®, alcohols, oils, modified oils, glycols, polysorbates, cyclodextrins, polymers, and/or combinations thereof.

[0643] Syrups and elixirs can be formulated with sweetening agents, for example glycerol, propylene glycol, sorbitol, glucose or sucrose. Such formulations can also contain a demulcent, a preservative and flavoring and coloring agents. The pharmaceutical compositions can be in the form of a sterile injectable aqueous or oleaginous suspension. This suspension can be formulated according to the known art using those suitable dispersing or wetting agents and suspending agents that have been mentioned above. The sterile injectable preparation can also be a sterile injectable solution or suspension in a non-toxic parentally acceptable diluent or solvent, for example as a solution in 1 ,3-butanediol. Among the acceptable vehicles and solvents that can be employed are water, Ringer's solution and isotonic sodium chloride solution. In addition, sterile, fixed oils are conventionally employed as a solvent or suspending medium. For this purpose, any bland fixed oil can be employed including synthetic mono- or diglycerides. In addition, fatty acids such as oleic acid find use in the preparation of injectables.

[0644] Suspensions for oral dosage may contain the active materials in a mixture with excipients suitable for the manufacture of aqueous suspensions. Such excipients may be suspending agents, as a non-limiting example the suspending agents may be sodium carboxymethylcellulose, methylcellulose, hydropropyl-methylcellulose, sodium alginate, polyvinylpyrrolidone, gum tragacanth and gum acacia; dispersing or wetting agents can be a naturally-occurring phosphatide, for example, lecithin, or condensation products of an alkylene oxide with fatty acids, for example polyoxyethylene stearate; or condensation products of ethylene oxide with long chain aliphatic alcohols, for example heptadecaethyleneoxycetanol, or condensation products of ethylene oxide with partial esters derived from fatty acids and a hexitol such as polyoxyethylene sorbitol monooleate, or condensation products of ethylene oxide with partial esters derived from fatty acids and hexitol anhydrides, for example polyethylene sorbitan monooleate. The aqueous suspensions may also contain one or more preservatives, for example ethyl, or n-propyl p-hydroxybenzoate, one or more coloring agents, one or more flavoring agents, and one or more sweetening agents, such as sucrose or saccharin.

[0645] Oily suspensions for oral dosage can be formulated by suspending the active ingredients in a vegetable oil, for example arachis oil, olive oil, sesame oil or coconut oil, or in a mineral oil such as liquid paraffin. The oily suspensions can contain a thickening agent, for example beeswax, hard paraffin or cetyl alcohol. Sweetening agents and flavoring agents can be added to provide palatable oral preparations. These compositions can be preserved by the addition of an anti-oxidant such as ascorbic acid

[0646] The oral dosage may also be in the form of oil-in-water emulsions. The oily phase can be a vegetable oil or a mineral oil or mixtures of these. Suitable emulsifying agents can be naturally-occurring gums, for example gum acacia or gum tragacanth, naturally-occurring phosphatides, for example soy bean, lecithin, and esters or partial esters derived from fatty acids and hexitol, anhydrides, for example sorbitan monooleate, and condensation products of the said partial esters with ethylene oxide, for example polyoxyethylene sorbitan monooleate. The emulsions may also contain sweetening and flavoring agents.

[0647] Solid dosage forms for oral administration include capsules, tablets, pills, powders, and granules. In such solid dosage forms, an active ingredient is mixed with at least one inert, pharmaceutically acceptable excipient such as sodium citrate or dicalcium phosphate and/or fillers or extenders (e.g. starches, lactose, sucrose, glucose, mannitol, and silicic acid), binders (e.g. carboxymethylcellulose, alginates, gelatin, polyvinylpyrrolidinone, sucrose, and acacia), humectants (e.g. glycerol), disintegrating agents (e.g. agar, calcium carbonate, potato or tapioca starch, alginic acid, certain silicates, and sodium carbonate), solution retarding agents (e.g. paraffin), absorption accelerators (e.g. quaternary ammonium compounds), wetting agents (e.g. cetyl alcohol and glycerol monostearate), absorbents (e.g. kaolin and bentonite clay), and lubricants (e.g. talc, calcium stearate, magnesium stearate, solid polyethylene glycols, sodium lauryl sulfate), and mixtures thereof. In the case of capsules, tablets and pills, the dosage form may comprise buffering agents. The solid dosage forms may also dissolve once they come in contact with liquid such as, but not limited to, salvia and bile.

[0648] Compositions intended for oral use can be prepared according to any method known to the art for the manu-

facture of pharmaceutical compositions and such compositions can contain one or more such sweetening agents, flavoring agents, coloring agents or preservative agents in order to provide pharmaceutically elegant and palatable preparations.

[0649] Solid dosage forms may be uncoated or they can be coated by known techniques. In some cases such coatings can be prepared by known techniques to delay disintegration and absorption in the gastrointestinal tract and thereby provide a sustained action over a longer period. For example, a time delay material such as glyceryl monosterate or glyceryl distearate can be employed.

[0650] Formulations for oral use can also be presented as hard gelatin capsules wherein the active ingredient is mixed with an inert solid diluent, for example, calcium carbonate, calcium phosphate or kaolin, or as soft gelatin capsules wherein the active ingredient is mixed with water or an oil medium, for example peanut oil, liquid paraffin or olive oil.

[0651] Dosage forms for oral delivery may also be chewable or may be suckable (e.g., lozenge form). The chewable dosages forms may be sustained release formulations such as, but not limited to, the sustained release compositions described in International Publication No WO2013082470 and US Publication No US20130142876. The chewable dosage forms may comprise amphipathic lipids such as, but not limited to, those described in International Publication No WO2013082470 and US Publication No US20130142876.

Topical or Transdermal Administration

[0652] As described herein, compositions containing the NAVs of the invention may be formulated for administration transdermally. The skin may be an ideal target site for delivery as it is readily accessible. Gene expression may be restricted not only to the skin, potentially avoiding nonspecific toxicity, but also to specific layers and cell types within the skin.

[0653] The site of cutaneous expression of the delivered compositions will depend on the route of nucleic acid delivery. Two routes are commonly considered to deliver NAVs to the skin: (ii) intradermal injection; and (iii) systemic delivery (e.g. for treatment of dermatologic diseases that affect both cutaneous and extracutaneous regions). NAVs can be delivered to the skin by several different approaches known in the art. After delivery of the nucleic acid, gene products have been detected in a number of different skin cell types, including, but not limited to, basal keratinocytes, sebaceous gland cells, dermal fibroblasts and dermal macrophages.

[0654] In one instance, the disclosure provides for the NAV compositions to be delivered in more than one injection.

[0655] In one instance, before transdermal administration at least one area of tissue, such as skin, may be subjected to a device and/or solution which may increase permeability In one instance, the tissue may be subjected to an abrasion device to increase the permeability of the skin (see U.S. Patent Publication No. 20080275468). In another instance, the tissue may be subjected to an ultrasound enhancement device. An ultrasound enhancement device may include, but is not limited to, the devices described in U.S. Publication No. 20040236268 and U.S. Patent Nos. 6,491,657 and 6,234,990. Methods of enhancing the permeability of tissue are described in U.S. Publication Nos. 20040171980 and 20040236268 and U.S. Pat. No. 6,190,315.

[0656] In one instance, a device may be used to increase permeability of tissue before delivering formulations of modified mRNA described herein. The permeability of skin may be measured by methods known in the art and/or described in U.S. Patent No. 6,190,315. As a non-limiting example, a modified mRNA formulation may be delivered by the drug delivery methods described in U.S. Patent No. 6,190,315.

[0657] In another non-limiting example tissue may be treated with a eutectic mixture of local anesthetics (EMLA) cream before, during and/or after the tissue may be subjected to a device which may increase permeability. Katz et al. (Anesth Analg (2004); 98:371-76) showed that using the EMLA cream in combination with a low energy, an onset of superficial cutaneous analgesia was seen as fast as 5 minutes after a pretreatment with a low energy ultrasound.

[0658] In one instance, enhancers may be applied to the tissue before, during, and/or after the tissue has been treated to increase permeability. Enhancers include, but are not limited to, transport enhancers, physical enhancers, and cavitation enhancers. Non-limiting examples of enhancers are described in U.S. Patent No. 6,190,315.

[0659] In one instance, a device may be used to increase permeability of tissue before delivering formulations of NAVs described herein, which may further contain a substance that invokes an immune response. In another non-limiting example, a formulation containing a substance to invoke an immune response may be delivered by the methods described in U.S. Publication Nos. 20040171980 and 20040236268.

[0660] Dosage forms for transdermal administration of a composition may include ointments, pastes, creams, lotions, gels, powders, solutions, sprays, inhalants and/or patches. Generally, an active ingredient is admixed under sterile conditions with a pharmaceutically acceptable excipient and/or any needed preservatives and/or buffers as may be required.

[0661] Additionally, the present disclosure contemplates the use of transdermal patches, which often have the added advantage of providing controlled delivery of a compound to the body. Such dosage forms may be prepared, for example, by dissolving and/or dispensing the compound in the proper medium. Alternatively or additionally, rate may be controlled

by either providing a rate controlling membrane and/or by dispersing the compound in a polymer matrix and/or gel.

[0662] A pharmaceutical NAV composition for transdermal administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for transdermal administration includes acrylates copolymer, acrylic acid-isooctyl acrylate copolymer, acrylic adhesive 788, adcote 72a103, aerotex resin 3730, alcohol, alcohol, dehydrated, aluminum polyester, bentonite, butylated hydroxytoluene, butylene glycol, butyric acid, caprylic/capric triglyceride, carbomer 1342, carbomer 940, carbomer 980, carrageenan, cetylpyridinium chloride, citric acid, crospovidone, daubert 1-5 pestr (matte) 164z, diethylene glycol monoethyl ether, diethylhexyl phthalate, dimethicone copolyol, dimethicone mdx4-4210, dimethicone medical fluid 360, dimethylaminoethyl methacrylate - butyl methacrylate - methyl methacrylate copolymer, dipropylene glycol, duro-tak 280-2516, duro-tak 387-2516, duro-tak 80-1196, duro-tak 87-2070, duro-tak 87-2194, duro-tak 87-2287, duro-tak 87-2296, duro-tak 87-2888, duro-tak 87-2979, edetate disodium, ethyl acetate, ethyl oleate, ethylcelluloses, ethylene vinyl acetate copolymer, ethylene-propylene copolymer, fatty acid esters, gelva 737, glycerin, glyceryl laurate, glyceryl oleate, heptane, high density polyethylene, hydrochloric acid, hydrogenated polybutene 635-690, hydroxyethyl cellulose, hydroxypropyl cellulose, isopropyl myristate, isopropyl palmitate, lactose, lanolin anhydrous, lauryl lactate, lecithin, levulinic acid, light mineral oil, medical adhesive modified s-15, methyl alcohol, methyl laurate, mineral oil, nitrogen, octisalate, octyldodecanol, oleic acid, oleyl alcohol, oleyl oleate, pentadecalactone, petrolatum, white, polacrilin, polyacrylic acid (250000 mw), polybutene (1400 mw), polyester, polyester polyamine copolymer, polyester rayon, polyethylene terephthalates, polyisobutylene, polyisobutylene (1100000 mw), polyisobutylene (35000 mw), polyisobutylene 178-236, polyisobutylene 241-294, polyisobutylene 35-39, polyisobutylene low molecular weight, polyisobutylene medium molecular weight, polyisobutylene/polybutene adhesive, polypropylene, polyvinyl acetate, polyvinyl alcohol, polyvinyl chloride, polyvinyl chloride-polyvinyl acetate copolymer, polyvinylpyridine, povidone k29/32, povidones, propylene glycol, propylene glycol monolaurate, ra-2397, ra-3011, silicon, silicon dioxide, colloidal, silicone, silicone adhesive 4102, silicone adhesive 4502, silicone adhesive bio-psa q7-4201, silicone adhesive bio-psa q7-4301, silicone/polyester film strip, sodium chloride, sodium citrate, sodium hydroxide, sorbitan monooleate, stearalkonium hectorite/propylene carbonate, titanium dioxide, triacetin, trolamine, tromethamine, union 76 amsco-res 6038 and viscose/cotton.

[0663] A pharmaceutical NAV composition for intradermal administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for intradermal administration includes benzalkonium chloride, benzyl alcohol, carboxymethylcellulose sodium, creatinine, edetate disodium, glycerin, hydrochloric acid, metacresol, methylparaben, phenol, polysorbate 80, protamine sulfate, sodium acetate, sodium bisulfite, sodium chloride, sodium hydroxide, sodium phosphate, sodium phosphate, dibasic, sodium phosphate, dibasic, heptahydrate, sodium phosphate, monobasic, anhydrous and zinc chloride.

Depot Administration

[0664] As described herein, in some instances, the composition is formulated in depots for extended release. Generally, a specific organ or tissue (a "target tissue") is targeted for administration.

[0665] In some aspects of the disclosure, the NAVs are spatially retained within or proximal to a target tissue. Provided are method of providing a composition to a target tissue of a mammalian subject by contacting the target tissue (which contains one or more target cells) with the composition under conditions such that the composition, in particular the nucleic acid component(s) of the composition, is substantially retained in the target tissue, meaning that at least 10, 20, 30, 40, 50, 60, 70, 80, 85, 90, 95, 96, 97, 98, 99, 99.9, 99.99 or greater than 99.99% of the composition is retained in the target tissue. Advantageously, retention is determined by measuring the amount of the nucleic acid present in the composition that enters one or more target cells. For example, at least 1, 5, 10, 20, 30, 40, 50, 60, 70, 80, 85, 90, 95, 96, 97, 98, 99, 99.9, 99.99 or greater than 99.99% of the nucleic acids administered to the subject are present intracellularly at a period of time following administration. For example, intramuscular injection to a mammalian subject is performed using an aqueous composition containing a ribonucleic acid and a transfection reagent, and retention of the composition is determined by measuring the amount of the ribonucleic acid present in the muscle cells.

[0666] Aspects of the disclosure are directed to methods of providing a composition to a target tissue of a mammalian subject, by contacting the target tissue (containing one or more target cells) with the composition under conditions such that the composition is substantially retained in the target tissue. The composition contains an effective amount of a polynucleotides such that the polypeptide of interest is produced in at least one target cell. The compositions generally contain a cell penetration agent, although "naked" NAV (such as nucleic acids without a cell penetration agent or other agent) is also contemplated, and a pharmaceutically acceptable carrier.

[0667] In some circumstances, the amount of a protein produced by cells in a tissue is desirably increased. Preferably, this increase in protein production is spatially restricted to cells within the target tissue. Thus, provided are methods of increasing production of a protein of interest in a tissue of a mammalian subject. A composition is provided that contains

polynucleotides characterized in that a unit quantity of composition has been determined to produce the polypeptide of interest in a substantial percentage of cells contained within a predetermined volume of the target tissue.

[0668] In some embodiments, the NAV composition includes a plurality of different polynucleotides, where one or more than one of the polynucleotides encodes a polypeptide of interest. Optionally, the composition also contains a cell penetration agent to assist in the intracellular delivery of the composition. A determination is made of the dose of the composition required to produce the polypeptide of interest in a substantial percentage of cells contained within the predetermined volume of the target tissue (generally, without inducing significant production of the polypeptide of interest in tissue adjacent to the predetermined volume, or distally to the target tissue). Subsequent to this determination, the determined dose is introduced directly into the tissue of the mammalian subject.

[0669] In one instance, the disclosure provides for the NAVs to be delivered in more than one injection or by split dose injections.

[0670] In one instance, the disclosure may be retained near target tissue using a small disposable drug reservoir, patch pump or osmotic pump. Non-limiting examples of patch pumps include those manufactured and/or sold by BDO (Franklin Lakes, NJ), Insulet Corporation (Bedford, MA), SteadyMed Therapeutics (San Francisco, CA), Medtronic (Minneapolis, MN) (e.g., MiniMed), UniLife (York, PA), Valeritas (Bridgewater, NJ), and SpringLeaf Therapeutics (Boston, MA). A non-limiting example of an osmotic pump include those manufactured by DURECT® (Cupertino, CA) (e.g., DUROS® and ALZET ®).

Pulmonary Administration

[0671] A pharmaceutical composition may be prepared, packaged, and/or sold in a formulation suitable for pulmonary administration via the buccal cavity. Such a formulation may comprise dry particles which comprise the active ingredient and which have a diameter in the range from about 0.5 nm to about 7 nm or from about 1 nm to about 6 nm. Such compositions are suitably in the form of dry powders for administration using a device comprising a dry powder reservoir to which a stream of propellant may be directed to disperse the powder and/or using a self propelling solvent/powder dispensing container such as a device comprising the active ingredient dissolved and/or suspended in a low-boiling propellant in a sealed container. Such powders comprise particles wherein at least 98% of the particles by weight have a diameter greater than 0.5 nm and at least 95% of the particles by number have a diameter less than 7 nm. Alternatively, at least 95% of the particles by weight have a diameter greater than 1 nm and at least 90% of the particles by number have a diameter less than 6 nm. Dry powder compositions may include a solid fine powder diluent such as sugar and are conveniently provided in a unit dose form.

[0672] Low boiling propellants generally include liquid propellants having a boiling point of below 65 °F at atmospheric pressure. Generally the propellant may constitute 50% to 99.9% (w/w) of the composition, and active ingredient may constitute 0.1% to 20% (w/w) of the composition. A propellant may further comprise additional ingredients such as a liquid non-ionic and/or solid anionic surfactant and/or a solid diluent (which may have a particle size of the same order as particles comprising the active ingredient).

[0673] As a non-limiting example, the NAVs described herein may be formulated for pulmonary delivery by the methods described in U.S. Pat. No. 8,257,685.

[0674] Pharmaceutical NAV compositions formulated for pulmonary delivery may provide an active ingredient in the form of droplets of a solution and/or suspension. Such formulations may be prepared, packaged, and/or sold as aqueous and/or dilute alcoholic solutions and/or suspensions, optionally sterile, comprising active ingredient, and may conveniently be administered using any nebulization and/or atomization device. Such formulations may further comprise one or more additional ingredients including, but not limited to, a flavoring agent such as saccharin sodium, a volatile oil, a buffering agent, a surface active agent, and/or a preservative such as methyl hydroxybenzoate. Droplets provided by this route of administration may have an average diameter in the range from about 0.1 nm to about 200 nm.

[0675] The compositions and formulations provided herein which may be used for pulmonary delivery may further comprise one or more surfactants. Suitable surfactants or surfactant components for enhancing the uptake of the compositions of the invention include synthetic and natural as well as full and truncated forms of surfactant protein A, surfactant protein B, surfactant protein C, surfactant protein D and surfactant Protein E, di-saturated phosphatidylcholine (other than dipalmitoyl), dipalmitoylphosphatidylcholine, phosphatidylcholine, phosphatidylglycerol, phosphatidylinositol, phosphatidylethanolamine, phosphatidylserine; phosphatidic acid, ubiquinones, lysophosphatidylethanolamine, lysophosphatidylcholine, palmitoyl-lysophosphatidylcholine, dehydroepiandrosterone, dolichols, sulfatidic acid, glycerol-3-phosphate, dihydroxyacetone phosphate, glycerol, glycero-3-phosphocholine, dihydroxyacetone, palmitate, cytidine diphosphate (CDP) diacylglycerol, CDP choline, choline, choline phosphate, as well as natural and artificial lamellar bodies which are the natural carrier vehicles for the components of surfactant, omega-3 fatty acids, polyenic acid, polyenoic acid, lecithin, palmitinic acid, non-ionic block copolymers of ethylene or propylene oxides, polyoxypropylene, monomeric and polymeric, polyoxyethylene, monomeric and polymeric, poly(vinyl amine) with dextran and/or alkanoyl side chains, Brij 35, Triton X-100 and synthetic surfactants ALEC, Exosurf, Survan and Atovaquone, among others. These surfactants

can be used either as single or part of a multiple component surfactant in a formulation, or as covalently bound additions to the 5' and/or 3' ends of the nucleic acid component of a pharmaceutical composition herein.

Intranasal, nasal and buccal Administration

[0676] Formulations described herein as being useful for pulmonary delivery are useful for intranasal delivery of a NAV pharmaceutical composition. Another formulation suitable for intranasal administration is a coarse powder comprising the active ingredient and having an average particle from about 0.2 μm to 500 μm. Such a formulation is administered in the manner in which snuff is taken, *i.e.* by rapid inhalation through the nasal passage from a container of the powder held close to the nose.

[0677] Formulations suitable for nasal administration may, for example, comprise from about as little as 0.1% (w/w) and as much as 100% (w/w) of active ingredient, and may comprise one or more of the additional ingredients described herein. A pharmaceutical composition may be prepared, packaged, and/or sold in a formulation suitable for buccal administration. Such formulations may, for example, be in the form of tablets and/or lozenges made using conventional methods and may, for example, 0.1% to 20% (w/w) active ingredient, the balance comprising an orally dissolvable and/or degradable composition and, optionally, one or more of the additional ingredients described herein. Alternately, formulations suitable for buccal administration may comprise a powder and/or an aerosolized and/or atomized solution and/or suspension comprising active ingredient. Such powdered, aerosolized, and/or aerosolized formulations, when dispersed, may have an average particle and/or droplet size in the range from about 0.1 nm to about 200 nm, and may further comprise one or more of any additional ingredients described herein.

[0678] A pharmaceutical NAV composition for inhalation (respiratory) administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for inhalation (respiratory) administration includes acetone sodium bisulfite, acetylcysteine, alcohol, alcohol, dehydrated, ammonia, apaflurane, ascorbic acid, benzalkonium chloride, calcium carbonate, carbon dioxide, cetylpyridinium chloride, chlorobutanol, citric acid, d&c yellow no. 10, dichlorodifluoromethane, dichlorotetrafluoroethane, edetate disodium, edetate sodium, fd&c yellow no. 6, fluorochlorohydrocarbons, gelatin, glycerin, glycine, hydrochloric acid, hydrochloric acid, diluted, lactose, lactose monohydrate, lecithin, lecithin, hydrogenated soy, lecithin, soybean, lysine monohydrate, mannitol, menthol, methylparaben, nitric acid, nitrogen, norflurane, oleic acid, polyethylene glycol 1000, povidone k25, propylene glycol, propylparaben, saccharin, saccharin sodium, silicon dioxide, colloidal, sodium bisulfate, sodium bisulfite, sodium chloride, sodium citrate, sodium hydroxide, sodium lauryl sulfate, sodium metabisulfite, sodium sulfate anhydrous, sodium sulfite, sorbitan trioleate, sulfuric acid, thymol, titanium dioxide, trichloromonofluoromethane, tromethamine and zinc oxide.

[0679] A pharmaceutical NAV composition for nasal administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for nasal administration includes acetic acid, alcohol, dehydrated, allyl alpha-ionone, anhydrous dextrose, anhydrous trisodium citrate, benzalkonium chloride, benzethonium chloride, benzyl alcohol, butylated hydroxyanisole, butylated hydroxytoluene, caffeine, carbon dioxide, carboxymethylcellulose sodium, cellulose, microcrystalline, chlorobutanol, citric acid, citric acid monohydrate, dextrose, dichlorodifluoromethane, dichlorotetrafluoroethane, edetate disodium, glycerin, glycerol ester of hydrogenated rosin, hydrochloric acid, hypromellose 2910 (15000 mpa.s), methylcelluloses, methylparaben, nitrogen, norflurane, oleic acid, petrolatum, white, phenylethyl alcohol, polyethylene glycol 3350, polyethylene glycol 400, polyoxyl 400 stearate, polysorbate 20, polysorbate 80, potassium phosphate, monobasic, potassium sorbate, propylene glycol, propylparaben, sodium acetate, sodium chloride, sodium citrate, sodium hydroxide, sodium phosphate, sodium phosphate, dibasic, sodium phosphate, dibasic, anhydrous, sodium phosphate, dibasic, dihydrate, sodium phosphate, dibasic, dodecahydrate, sodium phosphate, dibasic, heptahydrate, sodium phosphate, monobasic, anhydrous, sodium phosphate, monobasic, dihydrate, sorbitan trioleate, sorbitol, sorbitol solution, sucralose, sulfuric acid, trichloromonofluoromethane and trisodium citrate dihydrate.

Ophthalmic and Auricular (Otic) Administration

[0680] A pharmaceutical NAV composition may be prepared, packaged, and/or sold in a form ulation suitable for delivery to and/or around the eye and/or delivery to the ear (e.g., auricular (otic) administration). Non-limiting examples of route of administration for delivery to and/or around the eye include retrobulbar, conjuctival, intracorneal, intraocular, intravitreal, ophthlamic and subconjuctiva. Such formulations may, for example, be in the form of eye drops or ear drops including, for example, a 0.1/1.0% (wlw) solution and/or suspension of the active ingredient in an aqueous or oily liquid excipient. Such drops may further comprise buffering agents, salts, and/or one or more other of any additional ingredients described herein. Other ophthalmically-administrable formulations which are useful include those which comprise the active ingredient in microcrystalline form and/or in a liposomal preparation. Ear drops and/or eye drops are contemplated

as being within the scope of this invention. A multilayer thin film device may be prepared to contain a pharmaceutical composition for delivery to the eye and/or surrounding tissue.

[0681] A pharmaceutical NAV composition for ophthalmic administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for ophthalmic administration includes acetic acid, alcohol, alcohol, dehydrated, alginic acid, amerchol-cab, ammonium hydroxide, anhydrous trisodium citrate, antipyrine, benzalkonium chloride, benzethonium chloride, benzododecinium bromide, boric acid, caffeine, calcium chloride, carbomer 1342, carbomer 934p, carbomer 940, carbomer homopolymer type b (allyl pentaerythritol crosslinked), carboxymethylcellulose sodium, castor oil, cetyl alcohol, chlorobutanol, chlorobutanol, anhydrous, cholesterol, citric acid, citric acid monohydrate, creatinine, diethanolamine, diethylhexyl phthalate, divinylbenzene styrene copolymer, edetate disodium, edetate disodium anhydrous, edetate sodium, ethylene vinyl acetate copolymer, gellan gum (low acyl), glycerin, glyceryl stearate, high density polyethylene, hydrocarbon gel, plasticized, hydrochloric acid, hydrochloric acid, diluted, hydroxyethyl cellulose, hydroxypropyl methylcellulose 2906, hypromellose 2910 (15000 mpa.s), hypromelloses, jelene, lanolin, lanolin alcohols, lanolin anhydrous, lanolin nonionic derivatives, lauralkonium chloride, lauroyl sarcosine, light mineral oil, magnesium chloride, mannitol, methylcellulose (4000 mpa.s), methylcelluloses, methylparaben, mineral oil, nitric acid, nitrogen, nonoxynol-9, octoxynol-40, octylphenol polymethylene, petrolatum, petrolatum, white, phenylethyl alcohol, phenylmercuric acetate, phenylmercuric nitrate, phosphoric acid, polidronium chloride, poloxamer 188, poloxamer 407, polycarbophil, polyethylene glycol 300, polyethylene glycol 400, polyethylene glycol 8000, polyoxyethylene - polyoxypropylene 1800, polyoxyl 35 castor oil, polyoxyl 40 hydrogenated castor oil, polyoxyl 40 stearate, polypropylene glycol, polysorbate 20, polysorbate 60, polysorbate 80, polyvinyl alcohol, potassium acetate, potassium chloride, potassium phosphate, monobasic, potassium sorbate, povidone k29/32, povidone k30, povidone k90, povidones, propylene glycol, propylparaben, soda ash, sodium acetate, sodium bisulfate, sodium bisulfite, sodium borate, sodium borate decahydrate, sodium carbonate, sodium carbonate monohydrate, sodium chloride, sodium citrate, sodium hydroxide, sodium metabisulfite, sodium nitrate, sodium phosphate, sodium phosphate dihydrate, sodium phosphate, dibasic, sodium phosphate, dibasic, anhydrous, sodium phosphate, dibasic, dihydrate, sodium phosphate, dibasic, heptahydrate, sodium phosphate, monobasic, sodium phosphate, monobasic, anhydrous, sodium phosphate, monobasic, dihydrate, sodium phosphate, monobasic, monohydrate, sodium sulfate, sodium sulfate anhydrous, sodium sulfate decahydrate, sodium sulfite, sodium thiosulfate, sorbic acid, sorbitan monolaurate, sorbitol, sorbitol solution, stabilized oxychloro complex, sulfuric acid, thimerosal, titanium dioxide, tocophersolan, trisodium citrate dihydrate, triton 720, tromethamine, tyloxapol and zinc chloride.

[0682] A pharmaceutical NAV composition for retrobulbar administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for retrobulbar administration includes hydrochloric acid and sodium hydroxide.

[0683] A pharmaceutical NAV composition for intraocular administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for intraocular administration includes benzalkonium chloride, calcium chloride, citric acid monohydrate, hydrochloric acid, magnesium chloride, polyvinyl alcohol, potassium chloride, sodium acetate, sodium chloride, sodium citrate and sodium hydroxide.

[0684] A pharmaceutical NAV composition for intravitreal administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for intravitreal administration includes calcium chloride, carboxymethylcellulose sodium, cellulose, microcrystalline, hyaluronate sodium, hydrochloric acid, magnesium chloride, magnesium stearate, polysorbate 80, polyvinyl alcohol, potassium chloride, sodium acetate, sodium bicarbonate, sodium carbonate, sodium chloride, sodium hydroxide, sodium phosphate dibasic heptahydrate, sodium phosphate monobasic monohydrate and trisodium citrate dehydrate.

[0685] A pharmaceutical NAV composition for subconjunctival administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for subconjunctival administration includes benzyl alcohol, hydrochloric acid and sodium hydroxide.

[0686] A pharmaceutical NAV composition for auricular administration may comprise at least one inactive ingredient. Any or none of the inactive ingredients used may have been approved by the US Food and Drug Administration (FDA). A non-exhaustive list of inactive ingredients for use in pharmaceutical compositions for auricular administration includes acetic acid, aluminum acetate, aluminum sulfate anhydrous, benzalkonium chloride, benzethonium chloride, benzyl alcohol, boric acid, calcium carbonate, cetyl alcohol, chlorobutanol, chloroxylenol, citric acid, creatinine, cupric sulfate, cupric sulfate anhydrous, edetate disodium, edetic acid, glycerin, glyceryl stearate, hydrochloric acid, hydrocortisone, hydroxyethyl cellulose, isopropyl myristate, lactic acid, lecithin, hydrogenated, methylparaben, mineral oil, petrolatum, petrolatum, white, phenylethyl alcohol, polyoxyl 40 stearate, polyoxyl stearate, polysorbate 20, polysorbate 80, polyvinyl

alcohol, potassium metabisulfite, potassium phosphate, monobasic, povidone k90f, povidones, propylene glycol, propylene glycol diacetate, propylparaben, sodium acetate, sodium bisulfite, sodium borate, sodium chloride, sodium citrate, sodium hydroxide, sodium phosphate, dibasic, anhydrous, sodium phosphate, dibasic, heptahydrate, sodium phosphate, monobasic, anhydrous, sodium sulfite, sulfuric acid and thimerosal.

Combinations

[0687] The NAVs may be used in combination with one or more other therapeutic, prophylactic, diagnostic, or imaging agents. By "in combination with," it is not intended to imply that the agents must be administered at the same time and/or formulated for delivery together, although these methods of delivery are within the scope of the present disclosure. Compositions can be administered concurrently with, prior to, or subsequent to, one or more other desired therapeutics or medical procedures. In general, each agent will be administered at a dose and/or on a time schedule determined for that agent. In some instances, the present disclosure encompasses the delivery of pharmaceutical, prophylactic, diagnostic, or imaging compositions in combination with agents that may improve their bioavailability, reduce and/or modify their metabolism, inhibit their excretion, and/or modify their distribution within the body

[0688] The combinations can conveniently be presented for use in the form of a pharmaceutical formulation and thus pharmaceutical compositions comprising a combination as defined above together with a pharmaceutically acceptable diluent or carrier represent a further aspect of the disclosure.

[0689] The individual compounds of such combinations can be administered either sequentially or simultaneously in separate or combined pharmaceutical formulations. In one instance, the individual compounds will be administered simultaneously in a combined pharmaceutical formulation.

[0690] It will further be appreciated that therapeutically, prophylactically, diagnostically, or imaging active agents utilized in combination may be administered together in a single composition or administered separately in different compositions In general, it is expected that agents utilized in combination with be utilized at levels that do not exceed the levels at which they are utilized individually. In some instances, the levels utilized in combination will be lower than those utilized individually. In one instance, the combinations, each or together may be administered according to the split dosing regimens described herein.

Dosing

[0691] The present disclosure provides methods comprising administering NAVs and in accordance with the disclosure to a subject in need thereof. The exact amount required will vary from subject to subject, depending on the species, age, and general condition of the subject, the severity of the disease, the particular composition, its mode of administration, its mode of activity, and the like. Compositions in accordance with the disclosure are typically formulated in dosage unit form for ease of administration and uniformity of dosage. It will be understood, however, that the total daily usage of the compositions of the present disclosure may be decided by the attending physician within the scope of sound medical judgment. The specific therapeutically effective, prophylactically effective, or appropriate imaging dose level for any particular patient will depend upon a variety of factors including the disorder being treated and the severity of the disorder; the activity of the specific compound employed; the specific composition employed; the age, body weight, general health, sex and diet of the patient; the time of administration, route of administration, and rate of excretion of the specific compound employed; the duration of the treatment, drugs used in combination or coincidental with the specific compound employed; and like factors well known in the medical arts.

[0692] In certain instances, compositions in accordance with the present disclosure may be administered at dosage levels sufficient to deliver from about 0.0001 mg/kg to about 100 mg/kg, from about 0.001 mg/kg to about 0.05 mg/kg, from about 0.005 mg/kg to about 0.05 mg/kg, from about 0.001 mg/kg to about 0.005 mg/kg, from about 0.05 mg/kg to about 0.5 mg/kg, from about 0.01 mg/kg to about 50 mg/kg, from about 0.1 mg/kg to about 40 mg/kg, from about 0.5 mg/kg to about 30 mg/kg, from about 0.01 mg/kg to about 10 mg/kg, from about 0.1 mg/kg to about 10 mg/kg, or from about 1 mg/kg to about 25 mg/kg, of subject body weight per day, one or more times a day, to obtain the desired therapeutic, diagnostic, prophylactic, or imaging effect (see e.g., the range of unit doses described in International Publication No WO2013078199). The desired dosage may be delivered three times a day, two times a day, once a day, every other day, every third day, every week, every two weeks, every three weeks, or every four weeks. In certain instances, the desired dosage may be delivered using multiple administrations (e.g., two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen, fourteen, or more administrations). When multiple administrations are employed, split dosing regimens such as those described herein may be used.

[0693] According to the present disclosure, NAVs may be administered in split-dose regimens. As used herein, a "split dose" is the division of single unit dose or total daily dose into two or more doses, e.g, two or more administrations of the single unit dose. As used herein, a "single unit dose" is a dose of any therapeutic administed in one dose/at one time/single route/single point of contact, i.e., single administration event. As used herein, a "total daily dose" is an amount

given or prescribed in 24 hr period. It may be administered as a single unit dose. In one instance, the NAVs of the present disclosure are administed to a subject in split doses. The NAVs may be formulated in buffer only or in a formulation described herein.

Dosage Forms

[0694] A NAV pharmaceutical composition described herein can be formulated into a dosage form described herein, such as an intranasal, intratracheal, or injectable (e.g., intravenous, intraocular, intravitreal, intramuscular, intracardiac, intraperitoneal, subcutaneous).

Liquid dosage forms

[0695] Liquid dosage forms for parenteral administration include, but are not limited to, pharmaceutically acceptable emulsions, microemulsions, solutions, suspensions, syrups, and/or elixirs. In addition to active ingredients, liquid dosage forms may comprise inert diluents commonly used in the art including, but not limited to, wafer or other solvents, solubilizing agents and emulsifiers such as ethyl alcohol, isopropyl alcohol, ethyl carbonate, ethyl acetate, benzyl alcohol, benzyl benzoate, propylene glycol, 1,3-butylene glycol, dimethylformamide, oils (in particular, cottonseed, groundnut, corn, germ, olive, castor, and sesame oils), glycerol, tetrahydrofurfuryl alcohol, polyethylene glycols and fatty acid esters of sorbitan, and mixtures thereof. In certain instances for parenteral administration, compositions may be mixed with solubilizing agents such as CREMOPHOR®, alcohols, oils, modified oils, glycols, polysorbates, cyclodextrins, polymers, and/or combinations thereof.

Injectable

[0696] Injectable preparations, for example, sterile injectable aqueous or oleaginous suspensions may be formulated according to the known art and may include suitable dispersing agents, wetting agents, and/or suspending agents. Sterile injectable preparations may be sterile injectable solutions, suspensions, and/or emulsions in nontoxic parenterally acceptable diluents and/or solvents, for example, a solution in 1,3-butanediol. Among the acceptable vehicles and solvents that may be employed include, but are not limited to, wafer, Ringer's solution, U.S.P., and isotonic sodium chloride solution. Sterile, fixed oils are conventionally employed as a solvent or suspending medium. For this purpose any bland fixed oil can be employed including synthetic mono- or diglycerides. Fatty acids such as oleic acid can be used in the preparation of injectables.

[0697] Injectable formulations can be sterilized, for example, by filtration through a bacterial-retaining filter, and/or by incorporating sterilizing agents in the form of sterile solid compositions which can be dissolved or dispersed in sterile wafer or other sterile injectable medium prior to use.

[0698] In order to prolong the effect of an active ingredient, it may be desirable to slow the absorption of the active ingredient from subcutaneous or intramuscular injection. This may be accomplished by the use of a liquid suspension of crystalline or amorphous material with poor wafer solubility. The rate of absorption of the NAVs then depends upon its rate of dissolution which, in turn, may depend upon crystal size and crystalline form. Alternatively, delayed absorption of a parenterally administered NAV may be accomplished by dissolving or suspending the NAVs in an oil vehicle. Injectable depot forms are made by forming microencapsule matrices of the NAVs in biodegradable polymers such as polylactide-polyglycolide. Depending upon the ratio of NAVs to polymer and the nature of the particular polymer employed, the rate of polynucleotides release can be controlled. Examples of other biodegradable polymers include, but are not limited to, poly(orthoesters) and poly(anhydrides). Depot injectable formulations may be prepared by entrapping the NAVs in liposomes or microemulsions which are compatible with body tissues.

Pulmonary

[0699] Formulations described herein as being useful for pulmonary delivery may also be used for intranasal delivery of a pharmaceutical composition. Another formulation suitable for intranasal administration may be a coarse powder comprising the active ingredient and having an average particle from about 0.2 μm to 500 μm. Such a formulation may be administered in the manner in which snuff is taken, i.e. by rapid inhalation through the nasal passage from a container of the powder held close to the nose.

[0700] Formulations suitable for nasal administration may, for example, comprise from about as little as 0.1% (w/w) and as much as 100% (w/w) of active ingredient, and may comprise one or more of the additional ingredients described herein. A pharmaceutical composition may be prepared, packaged, and/or sold in a formulation suitable for buccal administration. Such formulations may, for example, be in the form of tablets and/or lozenges made using conventional methods, and may, for example, contain about 0.1% to 20% (w/w) active ingredient, where the balance may comprise

an orally dissolvable and/or degradable composition and, optionally, one or more of the additional ingredients described herein. Alternately, formulations suitable for buccal administration may comprise a powder and/or an aerosolized and/or atomized solution and/or suspension comprising active ingredient. Such powdered, aerosolized, and/or aerosolized formulations, when dispersed, may have an average particle and/or droplet size in the range from about 0.1 nm to about 200 nm, and may further comprise one or more of any additional ingredients described herein.

[0701] General considerations in the formulation and/or manufacture of pharmaceutical agents may be found, for example, in Remington: The Science and Practice of Pharmacy 21st ed., Lippincott Williams & Wilkins, 2005.

Coatings or Shells

[0702] Solid dosage forms of tablets, dragees, capsules, pills, and granules can be prepared with coatings and shells such as enteric coatings and other coatings well known in the pharmaceutical formulating art. They may optionally comprise opacifying agents and can be of a composition that they release the active ingredient(s) only, or preferentially, in a certain part of the intestinal tract, optionally, in a delayed manner. Examples of embedding compositions which can be used include polymeric substances and waxes. Solid compositions of a similar type may be employed as fillers in soft and hard-filled gelatin capsules using such excipients as lactose or milk sugar as well as high molecular weight polyethylene glycols and the like.

Multi-dose and repeat-dose administration

[0703] In some instances, NAV compounds and/or compositions of the present disclosure may be administered in two or more doses (referred to herein as "multi-dose administration"). Such doses may comprise the same components or may comprise components not included in a previous dose. Such doses may comprise the same mass and/or volume of components or an altered mass and/or volume of components in comparison to a previous dose. In some instances, multi-dose administration may comprise repeat-dose administration. As used herein, the term "repeat-dose administration" refers to two or more doses administered consecutively or within a regimen of repeat doses comprising substantially the same components provided at substantially the same mass and/or volume. In some instances, subjects may display a repeat-dose response. As used herein, the term "repeat-dose response" refers to a response in a subject to a repeat-dose that differs from that of another dose administered within a repeat-dose administration regimen. In some instances, such a response may be the expression of a protein in response to a repeat-dose comprising NAV. In such instances, protein expression may be elevated in comparison to another dose administered within a repeat-dose administration regimen or protein expression may be reduced in comparison to another dose administered within a repeat-dose administration regimen. Alteration of protein expression may be from about 1% to about 20%, from about 5% to about 50% from about 10% to about 60%, from about 25% to about 75%, from about 40% to about 100% and/or at least 100%. A reduction in expression of mRNA administered as part of a repeat-dose regimen, wherein the level of protein translated from the administered RNA is reduced by more than 40% in comparison to another dose within the repeat-dose regimen is referred to herein as "repeat-dose resistance."

Properties of the Pharmaceutical Compositions

[0704] The NAV pharmaceutical compositions described herein can be characterized by one or more of the following properties:

Bioavailability

[0705] The NAVs, when formulated into a composition with a delivery agent as described herein, can exhibit an increase in bioavailability as compared to a composition lacking a delivery agent as described herein. As used herein, the term "bioavailability" refers to the systemic availability of a given amount of NAVs administered to a mammal. Bioavailability can be assessed by measuring the area under the curve (AUC) or the maximum serum or plasma concentration (C_{max}) of the unchanged form of a compound following administration of the compound to a mammal. AUC is a determination of the area under the curve plotting the serum or plasma concentration of a compound along the ordinate (Y-axis) against time along the abscissa (X-axis). Generally, the AUC for a particular compound can be calculated using methods known to those of ordinary skill in the art and as described in G. S. Banker, Modern Pharmaceutics, Drugs and the Pharmaceutical Sciences, v. 72, Marcel Dekker, New York, Inc., 1996.

[0706] The C_{max} value is the maximum concentration of the compound achieved in the serum or plasma of a mammal following administration of the compound to the mammal. The C_{max} value of a particular compound can be measured using methods known to those of ordinary skill in the art. The phrases "increasing bioavailability" or "improving the pharmacokinetics," as used herein mean that the systemic availability of a first NAV, measured as AUC, C_{max}, or C_{min}

in a mammal is greater, when co-administered with a delivery agent as described herein, than when such co-administration does not take place. In some embodiments, the bioavailability of the NAVs can increase by at least about 2%, at least about 5%, at least about 10%, at least about 15%, at least about 20%, at least about 25%, at least about 30%, at least about 35%, at least about 40%, at least about 45%, at least about 50%, at least about 55%, at least about 60%, at least about 65%, at least about 70%, at least about 75%, at least about 80%, at least about 85%, at least about 90%, at least about 95%, or about 100%.

[0707] In some embodiments, liquid formulations of NAVs may have varying in vivo half-life, requiring modulation of doses to yield a therapeutic effect. To address this, in some embodiments of the present invention, NAV formulations may be designed to improve bioavailability and/or therapeutic effect during repeat administrations. Such formulations may enable sustained release of NAVs and/or reduce NAV degradation rates by nucleases. In some embodiments, suspension formulations are provided comprising NAVs, water immiscible oil depots, surfactants and/or co-surfactants and/or co-solvents. Combinations of oils and surfactants may enable suspension formulation with NAVs. Delivery of NAVs in a wafer immiscible depot may be used to improve bioavailability through sustained release of polynucleotides from the depot to the surrounding physiologic environment and/or prevent polynucleotide degradation by nucleases.

[0708] In some embodiments, cationic nanoparticles comprising combinations of divalent and monovalent cations may be formulated with NAVs. Such nanoparticles may form spontaneously in solution over a given period (e.g. hours, days, etc). Such nanoparticles do not form in the presence of divalent cations alone or in the presence of monovalent cations alone. The delivery of NAVs in cationic nanoparticles or in one or more depot comprising cationic nanoparticles may improve NAV bioavailability by acting as a long-acting depot and/or reducing the rate of degradation by nucleases.

Therapeutic Window

[0709] The NAVs, when formulated into a composition with a delivery agent as described herein, can exhibit an increase in the therapeutic window of the administered NAV composition as compared to the therapeutic window of the administered NAV composition lacking a delivery agent as described herein. As used herein "therapeutic window" refers to the range of plasma concentrations, or the range of levels of therapeutically active substance at the site of action, with a high probability of eliciting a therapeutic effect. In some instances, the therapeutic window of the NAVs when co-administered with a delivery agent as described herein can increase by at least about 2%, at least about 5%, at least about 10%, at least about 15%, at least about 20%, at least about 25%, at least about 30%, at least about 35%, at least about 40%, at least about 45%, at least about 50%, at least about 55%, at least about 60%, at least about 65%, at least about 70%, at least about 75%, at least about 80%, at least about 85%, at least about 90%, at least about 95%, or about 100%.

Volume of Distribution

[0710] The NAVs, when formulated into a composition with a delivery agent as described herein, can exhibit an improved volume of distribution (V_{dist}), e.g., reduced or targeted, relative to a composition lacking a delivery agent as described herein. The volume of distribution (Vdist) relates the amount of the drug in the body to the concentration of the drug in the blood or plasma. As used herein, the term "volume of distribution" refers to the fluid volume that would be required to contain the total amount of the drug in the body at the same concentration as in the blood or plasma: Vdist equals the amount of drug in the body/concentration of drug in blood or plasma. For example, for a 10 mg dose and a plasma concentration of 10 mg/L, the volume of distribution would be 1 liter. The volume of distribution reflects the extent to which the drug is present in the extravascular tissue. A large volume of distribution reflects the tendency of a compound to bind to the tissue components compared with plasma protein binding. In a clinical setting, Vdist can be used to determine a loading dose to achieve a steady state concentration. In some instances, the volume of distribution of the NAVs when co-administered with a delivery agent as described herein can decrease at least about 2%, at least about 5%, at least about 10%, at least about 15%, at least about 20%, at least about 25%, at least about 30%, at least about 35%, at least about 40%, at least about 45%, at least about 50%, at least about 55%, at least about 60%, at least about 65%, at least about 70%.

Biological Effect

[0711] In one instance, the biological effect of the NAV delivered to the animals may be categorized by analyzing the protein expression in the animals. The protein expression may be determined from analyzing a biological sample collected from a mammal administered the NAV of the present invention

Detection of Polynucleotides by Mass Spectrometry

[0712] Mass spectrometry (MS) is an analytical technique that can provide structural and molecular mass/concentration information on molecules after their conversion to ions. The molecules are first ionized to acquire positive or negative charges and then they travel through the mass analyzer to arrive at different areas of the detector according to their mass/charge (m/z) ratio.

[0713] Mass spectrometry is performed using a mass spectrometer which includes an ion source for ionizing the fractionated sample and creating charged molecules for further analysis. For example ionization of the sample may be performed by electrospray ionization (ESI), atmospheric pressure chemical ionization (APCI), photoionization, electron ionization, fast atom bombardment (FAB)lliquid secondary ionization (LSIMS), matrix assisted laser desorption/ionization (MALDI), field ionization, field desorption, thermospray/plasmaspray ionization, and particle beam ionization. The skilled artisan will understand that the choice of ionization method can be determined based on the analyte to be measured, type of sample, the type of detector, the choice of positive versus negative mode, etc.

[0714] After the sample has been ionized, the positively charged or negatively charged ions thereby created may be analyzed to determine a mass-to-charge ratio (i.e., m/z). Suitable analyzers for determining mass-to-charge ratios include quadropole analyzers, ion traps analyzers, and time-of-flight analyzers. The ions may be detected using several detection modes. For example, selected ions may be detected (i.e., using a selective ion monitoring mode (SIM)), or alternatively, ions may be detected using a scanning mode, e.g., multiple reaction monitoring (MRM) or selected reaction monitoring (SRM).

[0715] Liquid chromatography-multiple reaction monitoring (LC-MS/MRM) coupled with stable isotope labeled dilution of peptide standards has been shown to be an effective method for protein verification (e.g., Keshishian et al., Mol Cell Proteomics 2009 8: 2339-2349; Kuhn et al., Clin Chem 2009 55:1108-1117; Lopez et al., Clin Chem 2010 56:281-290). Unlike untargeted mass spectrometry frequently used in biomarker discovery studies, targeted MS methods are peptide sequence-based modes of MS that focus the full analytical capacity of the instrument on tens to hundreds of selected peptides in a complex mixture. By restricting detection and fragmentation to only those peptides derived from proteins of interest, sensitivity and reproducibility are improved dramatically compared to discovery-mode MS methods. This method of mass spectrometry-based multiple reaction monitoring (MRM) quantitation of proteins can dramatically impact the discovery and quantitation of biomarkers via rapid, targeted, multiplexed protein expression profiling of clinical samples.

[0716] In one instance, a biological sample which may contain at least one protein encoded by at least one modified mRNA of the present disclosure may be analyzed by the method of MRM-MS. The quantification of the biological sample may further include, but is not limited to, isotopically labeled peptides or proteins as internal standards.

[0717] According to the present disclosure, the biological sample, once obtained from the subject, may be subjected to enzyme digestion. As used herein, the term "digest" means to break apart into shorter peptides. As used herein, the phrase "treating a sample to digest proteins" means manipulating a sample in such a way as to break down proteins in a sample. These enzymes include, but are not limited to, trypsin, endoproteinase Glu-C and chymotrypsin. In one instance, a biological sample which may contain at least one protein encoded by at least one modified mRNA of the present disclosure may be digested using enzymes.

[0718] In one instance, a biological sample which may contain protein encoded by modified mRNA of the present disclosure may be analyzed for protein using electrospray ionization. Electrospray ionization (ESI) mass spectrometry (ESIMS) uses electrical energy to aid in the transfer of ions from the solution to the gaseous phase before they are analyzed by mass spectrometry. Samples may be analyzed using methods known in the art (e.g., Ho et al., Clin Biochem Rev. 2003 24(1):3-12). The ionic species contained in solution may be transferred into the gas phase by dispersing a fine spray of charge droplets, evaporating the solvent and ejecting the ions from the charged droplets to generate a mist of highly charged droplets. The mist of highly charged droplets may be analyzed using at least 1, at least 2, at least 3 or at least 4 mass analyzers such as, but not limited to, a quadropole mass analyzer. Further, the mass spectrometry method may include a purification step. As a non-limiting example, the first quadrapole may be set to select a single m/z ratio so it may filter out other molecular ions having a different m/z ratio which may eliminate complicated and time-consuming sample purification procedures prior to MS analysis.

[0719] In one instance, a biological sample which may contain protein encoded by modified mRNA of the present disclosure may be analyzed for protein in a tandem ESIMS system (e.g., MS/MS). As non-limiting examples, the droplets may be analyzed using a product scan (or daughter scan) a precursor scan (parent scan) a neutral loss or a multiple reaction monitoring.

[0720] In one instance, a biological sample which may contain protein encoded by modified mRNA of the present disclosure may be analyzed using matrix-assisted laser desorption/ionization (MALDI) mass spectrometry (MALDIMS). MALDI provides for the nondestructive vaporization and ionization of both large and small molecules, such as proteins. In MALDI analysis, the analyte is first co-crystallized with a large molar excess of a matrix compound, which may also include, but is not limited to, an ultraviolet absorbing weak organic acid. Non-limiting examples of matrices used in MALDI

are α-cyano-4-hydroxycinnamic acid, 3,5-dimethoxy-4-hydroxycinnamic acid and 2,5-dihydroxybenzoic acid. Laser radiation of the analyte-matrix mixture may result in the vaporization of the matrix and the analyte. The laser induced desorption provides high ion yields of the intact analyte and allows for measurement of compounds with high accuracy. Samples may be analyzed using methods known in the art (e.g., Lewis, Wei and Siuzdak, Encyclopedia of Analytical Chemistry 2000:5880-5894). As non-limiting examples, mass analyzers used in the MALDI analysis may include a linear time-of-flight (TOF), a TOF reflectron or a Fourier transform mass analyzer.

[0721] In one instance, the analyte-matrix mixture may be formed using the dried-droplet method. A biologic sample is mixed with a matrix to create a saturated matrix solution where the matrix-to-sample ratio is approximately 5000:1. An aliquot (approximately 0.5-2.0 uL) of the saturated matrix solution is then allowed to dry to form the analyte-matrix mixture.

[0722] In one instance, the analyte-matrix mixture may be formed using the thin-layer method. A matrix homogeneous film is first formed and then the sample is then applied and may be absorbed by the matrix to form the analyte-matrix mixture.

[0723] In one instance, the analyte-matrix mixture may be formed using the thick-layer method. A matrix homogeneous film is formed with a nitro-cellulose matrix additive. Once the uniform nitro-cellulose matrix layer is obtained the sample is applied and absorbed into the matrix to form the analyte-matrix mixture.

[0724] In one instance, the analyte-matrix mixture may be formed using the sandwich method. A thin layer of matrix crystals is prepared as in the thin-layer method followed by the addition of droplets of aqueous trifluoroacetic acid, the sample and matrix. The sample is then absorbed into the matrix to form the analyte-matrix mixture.

VI. Kits and Devices

Kits

[0725] The disclosure provides a variety of kits for conveniently and/or effectively carrying out methods of the present disclosure. Typically kits will comprise sufficient amounts and/or numbers of components to allow a user to perform multiple treatments of a subject(s) and/or to perform multiple experiments.

[0726] In one aspect, the present disclosure provides kits comprising the NAV molecules (including any proteins or polynucleotides) of the disclosure. In one instance, the kit comprises one or more functional antigens or function fragments thereof.

[0727] The kits can be for protein production, comprising a first polynucleotides comprising a translatable region of an antigen. The kit may further comprise packaging and instructions and/or a delivery agent to form a formulation composition. The delivery agent may comprise a saline, a buffered solution, a lipidoid or any delivery agent disclosed herein.

[0728] In one instance, the buffer solution may include sodium chloride, calcium chloride, phosphate and/or EDTA. In another instance, the buffer solution may include, but is not limited to, saline, saline with 2mM calcium, 5% sucrose, 5% sucrose with 2mM calcium, 5% Mannitol, 5% Mannitol with 2mM calcium, Ringer's lactate, sodium chloride, sodium chloride with 2mM calcium and mannose (See e.g., U.S. Pub. No. 20120258046). In a futher instance, the buffer solutions may be precipitated or it may be lyophilized. The amount of each component may be varied to enable consistent, reproducible higher concentration saline or simple buffer formulations.

[0729] The components may also be varied in order to increase the stability of polynucleotides in the buffer solution over a period of time and/or under a variety of conditions.

[0730] In one aspect, the present disclosure provides kits for protein production, comprising: a polynucleotide comprising a translatable region, provided in an amount effective to produce a desired amount of a protein encoded by the translatable region when introduced into a target cell; a second polynucleotide comprising an inhibitory nucleic acid, provided in an amount effective to substantially inhibit the innate immune response of the cell; and packaging and instructions.

[0731] In one aspect, the present disclosure provides kits for protein production, comprising a polynucleotide comprising a translatable region, wherein the polynucleotide exhibits reduced degradation by a cellular nuclease, and packaging and instructions.

[0732] In one aspect, the present disclosure provides kits for protein production, comprising a polynucleotide comprising a translatable region, wherein the polynucleotide exhibits reduced degradation by a cellular nuclease, and a mammalian cell suitable for translation of the translatable region of the first nucleic acid.

Devices

[0733] The present disclosure provides for devices which may incorporate RNAVs comprising polynucleotides that encode polypeptides of interest, e.g., encode antigenic polypeptides. These devices contain in a stable formulation the

reagents to synthesize a polynucleotide in a formulation available to be immediately delivered to a subject in need thereof, such as a human patient.

[0734] Devices for administration may be employed to deliver the NAVs of the present disclosure according to single, multi- or split-dosing regimens taught herein. Such devices are taught in, for example, International Application PCT/US2013/30062 filed March 9, 2013.

[0735] Method and devices known in the art for multi-administration to cells, organs and tissues are contemplated for use in conjunction with the methods and compositions disclosed herein as instances of the present disclosure. These include, for example, those methods and devices having multiple needles, hybrid devices employing for example lumens or catheters as well as devices utilizing heat, electric current or radiation driven mechanisms.

[0736] According to the present disclosure, these multi-administration devices may be utilized to deliver the single, multi- or split doses contemplated herein. Such devices are taught for example in, International Application PCT/US2013/30062 filed March 9, 2013.

[0737] In one instance, the NAV is administered subcutaneously or intramuscularly via at least 3 needles to three different, optionally adjacent, sites simultaneously, or within a 60 minutes period (e.g., administration to 4 5, 6, 7, 8, 9, or 10 sites simultaneously or within a 60 minute period).

Methods and Devices utilizing catheters and/or lumens

[0738] Methods and devices using catheters and lumens may be employed to administer the NAVs of the present disclosure on a single, multi- or split dosing schedule. Such methods and devices are described in International Application PCT/US2013/30062 filed March 9, 2013.

Methods and Devices utilizing electrical current

[0739] Methods and devices utilizing electric current may be employed to deliver the NAVs of the present disclosure according to the single, multi- or split dosing regimens taught herein. Such methods and devices are described in International Application PCT/US2013/30062 filed March 9, 2013.

VII. DEFINITIONS

[0740] At various places in the present specification, substituents of compounds of the present disclosure are disclosed in groups or in ranges. It is specifically intended that the present disclosure include each and every individual subcombination of the members of such groups and ranges.. For example, the term "C_{1-6} alkyl" is specifically intended to individually disclose methyl, ethyl, C_3 alkyl, C_4 alkyl, C_5 alkyl, and C_6 alkyl. Herein a phrase of the form "optionally substituted X" (*e.g.,* optionally substituted alkyl) is intended to be equivalent to "X, wherein X is optionally substituted" (*e.g.,* "alkyl, wherein said alkyl is optionally substituted"). It is not intended to mean that the feature "X" *(e.g.* alkyl) *per se* is optional.

[0741] *About* As used herein, the term "about" means +/- 10% of the recited value.

[0742] *Administered in combination:* As used herein, the term "administered in combination" or "combined administration" means that two or more agents are administered to a subject at the same time or within an interval such that there may be an overlap of an effect of each agent on the patient. In some instances, they are administered within about 60, 30, 15, 10, 5, or 1 minute of one another. In some instances, the administrations of the agents are spaced sufficiently closely together such that a combinatorial *(e.g.,* a synergistic) effect is achieved.

[0743] *Adjuvant*: As used herein, the term "adjuvant" means a substance that enhances a subject's immune response to an antigen. The NAVs of the present invention may optionally comprise one or more adjuvants.

[0744] *Animal:* As used herein, the term "animal" refers to any member of the animal kingdom. In some embodiments, "animal" refers to humans at any stage of development. In some embodiments, "animal" refers to non-human animals at any stage of development. In certain embodiments, the non-human animal is a mammal *(e.g.,* a rodent, a mouse, a rat, a rabbit, a monkey, a dog, a cat, a sheep, cattle, a primate, or a pig). In some embodiments, animals include, but are not limited to, mammals, birds, reptiles, amphibians, fish, and worms. In some embodiments, the animal is a transgenic animal, genetically-engineered animal, or a clone.

[0745] *Antigen:* As defined herein, the term "antigen" or "antibody generator" ("Ag") refers to a composition, for example, a substance or agent which causes an immune response in an organism, *e.g.,* causes the immune response of the organism to produce antibodies against the substance or agent in particular, which provokes an adaptive immune response in an organism. Antigens can be any immunogenic substance including, in particular, proteins, polypeptides, polysaccharides, nucleic acids, lipids and the like. Exemplary antigens are derived from infectious agents. Such agents can include parts or subunits of infectious agents, for example, coats, coat components, e.g., coat protein or polypeptides, surface components, *e.g.,* surface proteins or polypeptides, capsule components, cell wall components, flagella, fimbrae,

and/or toxins or toxoids) of infectious agents, for example, bacteria, viruses, and other microorganisms. Certain antigens, for example, lipids and/or nucleic acids are antigenic, preferably, when combined with proteins and/or polysaccharides.

[0746] *Antigens of interest or desired antigens:* As used herein, the terms "antigens of interest" or "desired antigens" include those proteins and other biomolecules provided herein that are components of or encoded by polynucleotides which are components of one or more NAVs.

[0747] *Approximately:* As used herein, the term "approximately" or "about," as applied to one or more values of interest, refers to a value that is similar to a stated reference value. In certain embodiments, the term "approximately" or "about" refers to a range of values that fall within 25%, 20%, 19%, 18%, 17%, 16%, 15%, 14%, 13%, 12%, 11%, 10%, 9%, 8%, 7%, 6%, 5%, 4%, 3%, 2%, 1%, or less in either direction (greater than or less than) of the stated reference value unless otherwise stated or otherwise evident from the context (except where such number would exceed 100% of a possible value).

[0748] *Associated with:* As used herein, the terms "associated with," "conjugated," "linked," "attached," and "tethered," when used with respect to two or more moieties, means that the moieties are physically associated or connected with one another, either directly or via one or more additional moieties that serves as a linking agent, to form a structure that is sufficiently stable so that the moieties remain physically associated under the conditions in which the structure is used, *e.g.,* physiological conditions. An "association" need not be strictly through direct covalent chemical bonding. It may also suggest ionic or hydrogen bonding or a hybridization based connectivity sufficiently stable such that the "associated" entities remain physically associated.

[0749] *Bifunctional:* As used herein, the term "bifunctional" refers to any substance, molecule or moiety which is capable of or maintains at least two functions. The functions may effect the same outcome or a different outcome. The structure that produces the function may be the same or different. For example, bifunctional modified RNAs of the present invention may encode a cytotoxic peptide (a first function) while those nucleosides which comprise the encoding RNA are, in and of themselves, cytotoxic (second function). In this example, delivery of the bifunctional modified RNA to a cancer cell would produce not only a peptide or protein molecule which may ameliorate or treat the cancer but would also deliver a cytotoxic payload of nucleosides to the cell should degradation, instead of translation of the modified RNA, occur.

[0750] *Biocompatible:* As used herein, the term "biocompatible" means compatible with living cells, tissues, organs or systems posing little to no risk of injury, toxicity or rejection by the immune system.

[0751] *Biodegradable:* As used herein, the term "biodegradable" means capable of being broken down into innocuous products by the action of living things.

[0752] *Biologically active:* As used herein, the phrase "biologically active" refers to a characteristic of any substance that has activity in a biological system and/or organism. For instance, a substance that, when administered to an organism, has a biological effect on that organism, is considered to be biologically active. In particular embodiments, a polynucleotide of the present invention may be considered biologically active if even a portion of the polynucleotides is biologically active or mimics an activity considered biologically relevant.

[0753] *Cancer stem cells:* As used herein, "cancer stem cells" are cells that can undergo self-renewal and/or abnormal proliferation and differentiation to form a tumor.

[0754] *Chemical terms:* The following provides the definition of various chemical terms from "acyl" to "thiol."

[0755] The term "acyl," as used herein, represents a hydrogen or an alkyl group (e.g., a haloalkyl group), as defined herein, that is attached to the parent molecular group through a carbonyl group, as defined herein, and is exemplified by formyl (i.e., a carboxyaldehyde group), acetyl, trifluoroacetyl, propionyl, butanoyl and the like. Exemplary unsubstituted acyl groups include from 1 to 7, from 1 to 11, or from 1 to 21 carbons. In some embodiments, the alkyl group is further substituted with 1, 2, 3, or 4 substituents as described herein.

[0756] Non-limiting examples of optionally substituted acyl groups include, alkoxycarbonyl, alkoxycarbonylacyl, arylalkoxycarbonyl, aryloyl, carbamoyl, carboxyaldehyde, (heterocyclyl) imino, and (heterocyclyl)oyl:

The "alkoxycarbonyl" group, which as used herein, represents an alkoxy, as defined herein, attached to the parent molecular group through a carbonyl atom (e.g., -C(O)-OR, where R is H or an optionally substituted C_{1-6}, C_{1-10}, or C_{1-20} alkyl group). Exemplary unsubstituted alkoxycarbonyl include from 1 to 21 carbons (e.g., from 1 to 11 or from 1 to 7 carbons). In some embodiments, the alkoxy group is further substituted with 1, 2, 3, or 4 substituents as described herein.

[0757] The "alkoxycarbonylacyl" group, which as used herein, represents an acyl group, as defined herein, that is substituted with an alkoxycarbonyl group, as defined herein (e.g., -C(O) -alkyl-C(O)-OR, where R is an optionally substituted C_{1-6}, C_{1-10}, or C_{1-20} alkyl group). Exemplary unsubstituted alkoxycarbonylacyl include from 3 to 41 carbons (e.g., from 3 to 10, from 3 to 13, from 3 to 17, from 3 to 21, or from 3 to 31 carbons, such as C_{1-6} alkoxycarbonyl-C_{1-6} acyl, C_{1-10} alkoxycarbonyl-C_{1-10} acyl, or C_{1-20} alkoxycarbonyl-C_{1-20} acyl). In some embodiments, each alkoxy and alkyl group is further independently substituted with 1, 2, 3, or 4 substituents, as described herein (e.g., a hydroxy group) for each group.

[0758] The "arylalkoxycarbonyl" group, which as used herein, represents an arylalkoxy group, as defined herein, attached to the parent molecular group through a carbonyl (e.g., -C(O)-O-alkyl-aryl). Exemplary unsubstituted arylalkoxy groups include from 8 to 31 carbons (e.g., from 8 to 17 or from 8 to 21 carbons, such as C_{6-10} aryl-C_{1-6} alkoxy-carbonyl,

C_{6-10} aryl-C_{1-10} alkoxy-carbonyl, or C_{6-10} aryl-C_{1-20} alkoxy-carbonyl). In some embodiments, the arylalkoxycarbonyl group can be substituted with 1, 2, 3, or 4 substituents as defined herein.

[0759] The "aryloyl" group, which as used herein, represents an aryl group, as defined herein, that is attached to the parent molecular group through a carbonyl group. Exemplary unsubstituted aryloyl groups are of 7 to 11 carbons. In some embodiments, the aryl group can be substituted with 1, 2, 3, or 4 substituents as defined herein.

[0760] The "carbamoyl" group, which as used herein, represents -C(O)-N(R^{N1})$_2$, where the meaning of each R^{N1} is found in the definition of "amino" provided herein.

[0761] The "carboxyaldehyde" group, which as used herein, represents an acyl group having the structure -CHO.

[0762] The "(heterocyclyl) imino" group, which as used herein, represents a heterocyclyl group, as defined herein, attached to the parent molecular group through an imino group. In some embodiments, the heterocyclyl group can be substituted with 1, 2, 3, or 4 substituent groups as defined herein.

[0763] The "(heterocyclyl)oyl" group, which as used herein, represents a heterocyclyl group, as defined herein, attached to the parent molecular group through a carbonyl group. In some embodiments, the heterocyclyl group can be substituted with 1, 2, 3, or 4 substituent groups as defined herein.

[0764] The term "alkyl," as used herein, is inclusive of both straight chain and branched chain saturated groups from 1 to 20 carbons (e.g., from 1 to 10 or from 1 to 6), unless otherwise specified. Alkyl groups are exemplified by methyl, ethyl, n- and iso-propyl, n-, sec-, iso- and tert-butyl, neopentyl, and the like, and may be optionally substituted with one, two, three, or, in the case of alkyl groups of two carbons or more, four substituents independently selected from the group consisting of: (1) C_{1-6} alkoxy; (2) C_{1-6} alkylsulfinyl; (3) amino, as defined herein (e.g., unsubstituted amino (i.e., -NH_2) or a substituted amino (i.e., -N(R^{N1})$_2$, where R^{N1} is as defined for amino); (4) C_{6-10} aryl-C_{1-6} alkoxy; (5) azido; (6) halo; (7) (C_{2-9} heterocyclyl)oxy; (8) hydroxy, optionally substituted with an O-protecting group; (9) nitro; (10) oxo (e.g., carboxyaldehyde or acyl); (11) C_{1-7} spirocyclyl; (12) thioalkoxy; (13) thiol; (14) - $CO_2R^{A'}$, optionally substituted with an O-protecting group and where $R^{A'}$ is selected from the group consisting of (a) C_{1-20} alkyl (e.g., C_{1-6} alkyl), (b) C_{2-20} alkenyl (e.g., C_{2-6} alkenyl), (c) C_{6-10} aryl, (d) hydrogen, (e) C_{1-6} alk-C_{6-10} aryl, (f) amino-C_{1-20} alkyl, (g) polyethylene glycol of-$(CH_2)_{s2}(OCH_2CH_2)_{s1}(CH_2)_{s3}OR'$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and R' is H or C_{1-20} alkyl, and (h) amino-polyethylene glycol of - $NR^{N1}(CH_2)_{s2}(CH_2CH_2O)_{s1}(CH_2)_{s3}NR^{N1}$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and each R^{N1} is, independently, hydrogen or optionally substituted C_{1-6} alkyl; (15) -C(O)$NR^{B'}R^{C'}$, where each of $R^{B'}$ and $R^{C'}$ is, independently, selected from the group consisting of (a) hydrogen, (b) C_{1-6} alkyl, (c) C_{6-10} aryl, and (d) C_{1-6} alk-C_{6-10} aryl; (16) -$SO_2R^{D'}$, where $R^{D'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, (c) C_{1-6} alk-C_{6-10} aryl, and (d) hydroxy; (17) -$SO_2NR^{E'}R^{F'}$, where each of $R^{E'}$ and $R^{F'}$ is, independently, selected from the group consisting of (a) hydrogen, (b) C_{1-6} alkyl, (c) C_{6-10} aryl and (d) C_{1-6} alk-C_{6-10} aryl; (18) -C(O)$R^{G'}$, where $R^{G'}$ is selected from the group consisting of (a) C_{1-20} alkyl (e.g., C_{1-6} alkyl), (b) C_{2-20} alkenyl (e.g., C_{2-6} alkenyl), (c) C_{6-10} aryl, (d) hydrogen, (e) C_{1-6} alk-C_{6-10} aryl, (f) amino-C_{1-20} alkyl, (g) polyethylene glycol of -$(CH_2)_{s2}(OCH_2CH_2)_{s1}(CH_2)_{s3}OR'$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and R' is H or C_{1-20} alkyl, and (h) amino-polyethylene glycol of -$NR^{N1}(CH_2)_{s2}(CH_2CH_2O)_{s1}(CH_2)_{s3}NR^{N1}$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and each R^{N1} is, independently, hydrogen or optionally substituted C_{1-6} alkyl; (19) -$NR^{H'}C(O)R^{I'}$, wherein $R^{H'}$ is selected from the group consisting of (a1) hydrogen and (b1) C_{1-6} alkyl, and $R^{I'}$ is selected from the group consisting of (a2) C_{1-20} alkyl (e.g., C_{1-6} alkyl), (b2) C_{2-20} alkenyl (e.g., C_{2-6} alkenyl), (c2) C_{6-10} aryl, (d2) hydrogen, (e2) C_{1-6} alk-C_{6-10} aryl, (f2) amino-C_{1-20} alkyl, (g2) polyethylene glycol of - $(CH_2)_{s2}(OCH_2CH_2)_{s1}(CH_2)_{s3}OR'$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and R' is H or C_{1-20} alkyl, and (h2) amino-polyethylene glycol of -$NR^{N1}(CH_2)_{s2}(CH_2CH_2O)_{s1}(CH_2)_{s3}NR^{N1}$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and each R^{N1} is, independently, hydrogen or optionally substituted C_{1-6} alkyl; (20) -$NR^{J'}C(O)OR^{K'}$, wherein $R^{J'}$ is selected from the group consisting of (a1) hydrogen and (b1) C_{1-6} alkyl, and $R^{K'}$ is selected from the group consisting of (a2) C_{1-20} alkyl (e.g., C_{1-6} alkyl), (b2) C_{2-20} alkenyl (e.g., C_{2-6} alkenyl), (c2) C_{6-10} aryl, (d2) hydrogen, (e2) C_{1-6} alk-C_{6-10} aryl, (f2) amino-C_{1-20} alkyl, (g2) polyethylene glycol of -$(CH_2)_{s2}(OCH_2CH_2)_{s1}(CH_2)_{s3}OR'$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and R' is H or C_{1-20} alkyl, and (h2) amino-polyethylene glycol of -$NR^{N1}(CH_2)_{s2}(CH_2CH_2O)_{s1}(CH_2)_{s3}NR^{N1}$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and each R^{N1} is, independently, hydrogen or optionally substituted C_{1-6} alkyl; and (21) amidine. In some embodiments, each of these groups can be further substituted as described herein. For example, the alkylene group of a

C_1-alkaryl can be further substituted with an oxo group to afford the respective aryloyl substituent.

[0765] The term "alkylene," as used herein, represent a saturated divalent hydrocarbon group derived from a straight or branched chain saturated hydrocarbon by the removal of two hydrogen atoms, and is exemplified by methylene, ethylene, isopropylene, and the like. The term "C_{x-y}alkylene" and the prefix "C_{x-y} alk-" represent alkylene groups having between x and y carbons. Exemplary values for x are 1, 2, 3, 4, 5, and 6, and exemplary values for y are 2, 3, 4, 5, 6, 7, 8, 9, 10, 12, 14, 16, 18, or 20 (e.g., C_{1-6}, C_{1-10}, C_{2-20}, C_{2-6}, C_{2-10}, or C_{2-20} alkylene). In some embodiments, the alkylene can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein for an alkyl group. Similarly, the suffix "-ene" appended to any group indicates that the group is a divalent group.

[0766] Non-limiting examples of optionally substituted alkyl and alkylene groups include acylaminoalkyl, acyloxyalkyl, alkoxyalkyl, alkoxycarbonylalkyl, alkylsulfinyl, alkylsulfinylalkyl, aminoalkyl, carbamoylalkyl, carboxyalkyl, carboxyaminoalkyl, haloalkyl, hydroxyalkyl, perfluoroalkyl, and sulfoalkyl:

The "acylaminoalkyl" group, which as used herein, represents an acyl group, as defined herein, attached to an amino group that is in turn attached to the parent molecular group through an alkylene group, as defined herein (i.e., -alkyl-$N(R^{N1})$-C(O)-R, where R is H or an optionally substituted C_{1-6}, C_{1-10}, or C_{1-20} alkyl group (e.g., haloalkyl) and R^{N1} is as defined herein). Exemplary unsubstituted acylaminoalkyl groups include from 1 to 41 carbons (e.g., from 1 to 7, from 1 to 13, from 1 to 21, from 2 to 7, from 2 to 13, from 2 to 21, or from 2 to 41 carbons). In some embodiments, the alkylene group is further substituted with 1, 2, 3, or 4 substituents as described herein, and/or the amino group is $-NH_2$ or $-NHR^{N1}$, wherein R^{N1} is, independently, OH, NO_2, NH_2, NR^{N2}_2, SO_2OR^{N2}, SO_2R^{N2}, SOR^{N2}, alkyl, aryl, acyl (e.g., acetyl, trifluoroacetyl, or others described herein), or alkoxycarbonylalkyl, and each R^{N2} can be H, alkyl, or aryl.

[0767] The "acyloxyalkyl" group, which as used herein, represents an acyl group, as defined herein, attached to an oxygen atom that in turn is attached to the parent molecular group though an alkylene group (i.e., -alkyl-O-C(O)-R, where R is H or an optionally substituted C_{1-6}, C_{1-10}, or C_{1-20} alkyl group). Exemplary unsubstituted acyloxyalkyl groups include from 1 to 21 carbons (e.g., from 1 to 7 or from 1 to 11 carbons). In some embodiments, the alkylene group is, independently, further substituted with 1, 2, 3, or 4 substituents as described herein.

[0768] The "alkoxyalkyl" group, which as used herein, represents an alkyl group that is substituted with an alkoxy group. Exemplary unsubstituted alkoxyalkyl groups include between 2 to 40 carbons (e.g., from 2 to 12 or from 2 to 20 carbons, such as C_{1-6} alkoxy-C_{1-6} alkyl, C_{1-10} alkoxy-C_{1-10} alkyl, or C_{1-20} alkoxy-C_{1-20} alkyl). In some embodiments, the alkyl and the alkoxy each can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein for the respective group.

[0769] The "alkoxycarbonylalkyl" group, which as used herein, represents an alkyl group, as defined herein, that is substituted with an alkoxycarbonyl group, as defined herein (e.g., -alkyl-C(O)-OR, where R is an optionally substituted C_{1-20}, C_{1-10}, or C_{1-6} alkyl group). Exemplary unsubstituted alkoxycarbonylalkyl include from 3 to 41 carbons (e.g., from 3 to 10, from 3 to 13, from 3 to 17, from 3 to 21, or from 3 to 31 carbons, such as C_{1-6} alkoxycarbonyl-C_{1-6} alkyl, C_{1-10} alkoxycarbonyl-C_{1-10} alkyl, or C_{1-20} alkoxycarbonyl-C_{1-20} alkyl). In some embodiments, each alkyl and alkoxy group is further independently substituted with 1, 2, 3, or 4 substituents as described herein (e.g., a hydroxy group).

[0770] The "alkylsulfinylalkyl" group, which as used herein, represents an alkyl group, as defined herein, substituted with an alkylsulfinyl group. Exemplary unsubstituted alkylsulfinylalkyl groups are from 2 to 12, from 2 to 20, or from 2 to 40 carbons. In some embodiments, each alkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein.

[0771] The "aminoalkyl" group, which as used herein, represents an alkyl group, as defined herein, substituted with an amino group, as defined herein. The alkyl and amino each can be further substituted with 1, 2, 3, or 4 substituent groups as described herein for the respective group (e.g., $CO_2R^{A'}$, where $R^{A'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, (c) hydrogen, and (d) C_{1-6} alk-C_{6-10} aryl, e.g., carboxy, and/or an N-protecting group).

[0772] The "carbamoylalkyl" group, which as used herein, represents an alkyl group, as defined herein, substituted with a carbamoyl group, as defined herein. The alkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein.

[0773] The "carboxyalkyl" group, which as used herein, represents an alkyl group, as defined herein, substituted with a carboxy group, as defined herein. The alkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein, and the carboxy group can be optionally substituted with one or more O-protecting groups.

[0774] The "carboxyaminoalkyl" group, which as used herein, represents an aminoalkyl group, as defined herein, substituted with a carboxy, as defined herein. The carboxy, alkyl, and amino each can be further substituted with 1, 2, 3, or 4 substituent groups as described herein for the respective group (e.g., $CO_2R^{A'}$, where $R^{A'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, (c) hydrogen, and (d) C_{1-6} alk-C_{6-10} aryl, e.g., carboxy, and/or an *N*-protecting group, and/or an O-protecting group).

[0775] The "haloalkyl" group, which as used herein, represents an alkyl group, as defined herein, substituted with a halogen group (i.e., F, Cl, Br, or I). A haloalkyl may be substituted with one, two, three, or, in the case of alkyl groups of two carbons or more, four halogens. Haloalkyl groups include perfluoroalkyls (e.g., $-CF_3$), $-CHF_2$, $-CH_2F$, $-CCl_3$, $-CH_2CH_2Br$, $-CH_2CH(CH_2CH_2Br)CH_3$, and $-CHICH_3$. In some embodiments, the haloalkyl group can be further substi-

tuted with 1, 2, 3, or 4 substituent groups as described herein for alkyl groups.

[0776] The "hydroxyalkyl" group, which as used herein, represents an alkyl group, as defined herein, substituted with one to three hydroxy groups, with the proviso that no more than one hydroxy group may be attached to a single carbon atom of the alkyl group, and is exemplified by hydroxymethyl, dihydroxypropyl, and the like. In some embodiments, the hydroxyalkyl group can be substituted with 1, 2, 3, or 4 substituent groups (e.g., O-protecting groups) as defined herein for an alkyl.

[0777] The "perfluoroalkyl" group, which as used herein, represents an alkyl group, as defined herein, where each hydrogen radical bound to the alkyl group has been replaced by a fluoride radical. Perfluoroalkyl groups are exemplified by trifluoromethyl, pentafluoroethyl, and the like.

[0778] The "sulfoalkyl" group, which as used herein, represents an alkyl group, as defined herein, substituted with a sulfo group of $-SO_3H$. In some embodiments, the alkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein, and the sulfo group can be further substituted with one or more O-protecting groups (e.g., as described herein).

[0779] The term "alkenyl," as used herein, represents monovalent straight or branched chain groups of, unless otherwise specified, from 2 to 20 carbons (e.g., from 2 to 6 or from 2 to 10 carbons) containing one or more carbon-carbon double bonds and is exemplified by ethenyl, 1-propenyl, 2-propenyl, 2-methyl-1-propenyl, 1-butenyl, 2-butenyl, and the like. Alkenyls include both cis and trans isomers. Alkenyl groups may be optionally substituted with 1, 2, 3, or 4 substituent groups that are selected, independently, from amino, aryl, cycloalkyl, or heterocyclyl (e.g., heteroaryl), as defined herein, or any of the exemplary alkyl substituent groups described herein.

[0780] Non-limiting examples of optionally substituted alkenyl groups include, alkoxycarbonylalkenyl, aminoalkenyl, and hydroxyalkenyl:

The "alkoxycarbonylalkenyl" group, which as used herein, represents an alkenyl group, as defined herein, that is substituted with an alkoxycarbonyl group, as defined herein (e.g., -alkenyl-C(O)-OR, where R is an optionally substituted C_{1-20}, C_{1-10}, or C_{1-6} alkyl group). Exemplary unsubstituted alkoxycarbonylalkenyl include from 4 to 41 carbons (e.g., from 4 to 10, from 4 to 13, from 4 to 17, from 4 to 21, or from 4 to 31 carbons, such as C_{1-6} alkoxycarbonyl-C_{2-6} alkenyl, C_{1-10} alkoxycarbonyl-C_{2-10} alkenyl, or C_{1-20} alkoxycarbonyl-C_{2-20} alkenyl). In some embodiments, each alkyl, alkenyl, and alkoxy group is further independently substituted with 1, 2, 3, or 4 substituents as described herein (e.g., a hydroxy group).

[0781] The "aminoalkenyl" group, which as used herein, represents an alkenyl group, as defined herein, substituted with an amino group, as defined herein. The alkenyl and amino each can be further substituted with 1, 2, 3, or 4 substituent groups as described herein for the respective group (e.g., $CO_2R^{A'}$, where $R^{A'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, (c) hydrogen, and (d) C_{1-6} alk-C_{6-10} aryl, e.g., carboxy, and/or an N-protecting group).

[0782] The "hydroxyalkenyl" group, which as used herein, represents an alkenyl group, as defined herein, substituted with one to three hydroxy groups, with the proviso that no more than one hydroxy group may be attached to a single carbon atom of the alkyl group, and is exemplified by dihydroxypropenyl, hydroxyisopentenyl, and the like. In some embodiments, the hydroxyalkenyl group can be substituted with 1, 2, 3, or 4 substituent groups (e.g., O-protecting groups) as defined herein for an alkyl.

[0783] The term "alkynyl," as used herein, represents monovalent straight or branched chain groups from 2 to 20 carbon atoms (e.g., from 2 to 4, from 2 to 6, or from 2 to 10 carbons) containing a carbon-carbon triple bond and is exemplified by ethynyl, 1-propynyl, and the like. Alkynyl groups may be optionally substituted with 1, 2, 3, or 4 substituent groups that are selected, independently, from aryl, cycloalkyl, or heterocyclyl (e.g., heteroaryl), as defined herein, or any of the exemplary alkyl substituent groups described herein.

[0784] Non-limiting examples of optionally substituted alkynyl groups include alkoxycarbonylalkynyl, aminoalkynyl, and hydroxyalkynyl:

The "alkoxycarbonylalkynyl" group, which as used herein, represents an alkynyl group, as defined herein, that is substituted with an alkoxycarbonyl group, as defined herein (e.g., -alkynyl-C(O)-OR, where R is an optionally substituted C_{1-20}, C_{1-10}, or C_{1-6} alkyl group). Exemplary unsubstituted alkoxycarbonylalkynyl include from 4 to 41 carbons (e.g., from 4 to 10, from 4 to 13, from 4 to 17, from 4 to 21, or from 4 to 31 carbons, such as C_{1-6} alkoxycarbonyl-C_{2-6} alkynyl, C_{1-10} alkoxycarbonyl-C_{2-10} alkynyl, or C_{1-20} alkoxycarbonyl-C_{2-20} alkynyl). In some embodiments, each alkyl, alkynyl, and alkoxy group is further independently substituted with 1, 2, 3, or 4 substituents as described herein (e.g., a hydroxy group).

[0785] The "aminoalkynyl" group, which as used herein, represents an alkynyl group, as defined herein, substituted with an amino group, as defined herein. The alkynyl and amino each can be further substituted with 1, 2, 3, or 4 substituent groups as described herein for the respective group (e.g., $CO_2R^{A'}$, where $R^{A'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, (c) hydrogen, and (d) C_{1-6} alk-C_{6-10} aryl, e.g., carboxy, and/or an N-protecting group).

[0786] The "hydroxyalkynyl" group, which as used herein, represents an alkynyl group, as defined herein, substituted with one to three hydroxy groups, with the proviso that no more than one hydroxy group may be attached to a single carbon atom of the alkyl group. In some embodiments, the hydroxyalkynyl group can be substituted with 1, 2, 3, or 4

substituent groups (e.g., O-protecting groups) as defined herein for an alkyl.

[0787] The term "amino," as used herein, represents -N(R^{N1})$_2$, wherein each R^{N1} is, independently, H, OH, NO_2, N(R^{N2})$_2$, SO_2OR^{N2}, SO_2R^{N2}, SOR^{N2}, an N-protecting group, alkyl, alkenyl, alkynyl, alkoxy, aryl, alkaryl, cycloalkyl, alkcycloalkyl, carboxyalkyl (e.g., optionally substituted with an O-protecting group, such as optionally substituted arylalkoxycarbonyl groups or any described herein), sulfoalkyl, acyl (e.g., acetyl, trifluoroacetyl, or others described herein), alkoxycarbonylalkyl (e.g., optionally substituted with an O-protecting group, such as optionally substituted arylalkoxycarbonyl groups or any described herein), heterocyclyl (e.g., heteroaryl), or alkheterocyclyl (e.g., alkheteroaryl), wherein each of these recited R^{N1} groups can be optionally substituted, as defined herein for each group, or two R^{N1} combine to form a heterocyclyl or an N-protecting group, and wherein each R^{N2} is, independently, H, alkyl, or aryl. The amino groups of the invention can be an unsubstituted amino (i.e., -NH_2) or a substituted amino (i.e., -N(R^{N1})$_2$). In a preferred embodiment, amino is -NH_2 or -NHR^{N1}, wherein R^{N1} is, independently, OH, NO_2, NH_2, $NR^{N2}{}_2$, SO_2OR^{N2}, SO_2R^{N2}, SOR^{N2}, alkyl, carboxyalkyl, sulfoalkyl, acyl (e.g., acetyl, trifluoroacetyl, or others described herein), alkoxycarbonylalkyl (e.g., t-butoxycarbonylalkyl) or aryl, and each R^{N2} can be H, C_{1-20} alkyl (e.g., C_{1-6} alkyl), or C_{6-10} aryl.

[0788] Non-limiting examples of optionally substituted amino groups include acylamino and carbamyl:

The "acylamino" group, which as used herein, represents an acyl group, as defined herein, attached to the parent molecular group though an amino group, as defined herein (i.e., -N(R^{N1})-C(O)-R, where R is H or an optionally substituted C_{1-6}, C_{1-10}, or C_{1-20} alkyl group (e.g., haloalkyl) and R^{N1} is as defined herein). Exemplary unsubstituted acylamino groups include from 1 to 41 carbons (e.g., from 1 to 7, from 1 to 13, from 1 to 21, from 2 to 7, from 2 to 13, from 2 to 21, or from 2 to 41 carbons). In some embodiments, the alkyl group is further substituted with 1, 2, 3, or 4 substituents as described herein, and/or the amino group is -NH_2 or -NHR^{N1}, wherein R^{N1} is, independently, OH, NO_2, NH_2, $NR^{N2}{}_2$, SO_2OR^{N2}, SO_2R^{N2}, SOR^{N2}, alkyl, aryl, acyl (e.g., acetyl, trifluoroacetyl, or others described herein), or alkoxycarbonylalkyl, and each R^{N2} can be H, alkyl, or aryl.

[0789] The "carbamyl" group, which as used herein, refers to a carbamate group having the structure -$NR^{N1}C(=O)OR$ or -$OC(=O)N(R^{N1})_2$, where the meaning of each R^{N1} is found in the definition of "amino" provided herein, and R is alkyl, cycloalkyl , alkcycloalkyl, aryl, alkaryl, heterocyclyl (e.g., heteroaryl), or alkheterocyclyl (e.g., alkheteroaryl), as defined herein.

[0790] The term "amino acid," as described herein, refers to a molecule having a side chain, an amino group, and an acid group (e.g., a carboxy group of -CO_2H or a sulfo group of-SO_3H), wherein the amino acid is attached to the parent molecular group by the side chain, amino group, or acid group (e.g., the side chain). In some embodiments, the amino acid is attached to the parent molecular group by a carbonyl group, where the side chain or amino group is attached to the carbonyl group. Exemplary side chains include an optionally substituted alkyl, aryl, heterocyclyl, alkaryl, alkheterocyclyl, aminoalkyl, carbamoylalkyl, and carboxyalkyl. Exemplary amino acids include alanine, arginine, asparagine, aspartic acid, cysteine, glutamic acid, glutamine, glycine, histidine, hydroxynorvaline, isoleucine, leucine, lysine, methionine, norvaline, ornithine, phenylalanine, proline, pyrrolysine, selenocysteine, serine, taurine, threonine, tryptophan, tyrosine, and valine. Amino acid groups may be optionally substituted with one, two, three, or, in the case of amino acid groups of two carbons or more, four substituents independently selected from the group consisting of: (1) C_{1-6} alkoxy; (2) C_{1-6} alkylsulfinyl; (3) amino, as defined herein (e.g., unsubstituted amino (i.e., -NH_2) or a substituted amino (i.e., -N(R^{N1})$_2$, where R^{N1} is as defined for amino); (4) C_{6-10} aryl-C_{1-6} alkoxy; (5) azido; (6) halo; (7) (C_{2-9} heterocyclyl)oxy; (8) hydroxy; (9) nitro; (10) oxo (e.g., carboxyaldehyde or acyl); (11) C_{1-7} spirocyclyl; (12) thioalkoxy; (13) thiol; (14) -$CO_2R^{A'}$, where $R^{A'}$ is selected from the group consisting of (a) C_{1-20} alkyl (e.g., C_{1-6} alkyl), (b) C_{2-20} alkenyl (e.g., C_{2-6} alkenyl), (c) C_{6-10} aryl, (d) hydrogen, (e) C_{1-6} alk-C_{6-10} aryl, (f) amino-C_{1-20} alkyl, (g) polyethylene glycol of -$(CH_2)_{s2}(OCH_2CH_2)_{s1}(CH_2)_{s3}OR'$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and R' is H or C_{1-20} alkyl, and (h) amino-polyethylene glycol of -$NR^{N1}(CH_2)_{s2}(CH_2CH_2O)_{s1}(CH_2)_{s3}NR^{N1}$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and each R^{N1} is, independently, hydrogen or optionally substituted C_{1-6} alkyl; (15) -$C(O)NR^{B'}R^{C'}$, where each of $R^{B'}$ and $R^{C'}$ is, independently, selected from the group consisting of (a) hydrogen, (b) C_{1-6} alkyl, (c) C_{6-10} aryl, and (d) C_{1-6} alk-C_{6-10} aryl; (16) - $SO_2R^{D'}$, where $R^{D'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, (c) C_{1-6} alk-C_{6-10} aryl, and (d) hydroxy; (17) -$SO_2NR^{E'}R^{F'}$, where each of $R^{E'}$ and $R^{F'}$ is, independently, selected from the group consisting of (a) hydrogen, (b) C_{1-6} alkyl, (c) C_{6-10} aryl and (d) C_{1-6} alk-C_{6-10} aryl; (18) - $C(O)R^{G'}$, where $R^{G'}$ is selected from the group consisting of (a) C_{1-20} alkyl (e.g., C_{1-6} alkyl), (b) C_{2-20} alkenyl (e.g., C_{2-6} alkenyl), (c) C_{6-10} aryl, (d) hydrogen, (e) C_{1-6} alk-C_{6-10} aryl, (f) amino-C_{1-20} alkyl, (g) polyethylene glycol of -$(CH_2)_{s2}(OCH_2CH_2)_{s1}(CH_2)_{s3}OR'$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and R' is H or C_{1-20} alkyl, and (h) amino-polyethylene glycol of -$NR^{N1}(CH_2)_{s2}(CH_2CH_2O)_{s1}(CH_2)_{s3}NR^{N1}$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and each R^{N1} is, independently, hydrogen or optionally substituted C_{1-6} alkyl; (19) -$NR^{H'}C(O)R^{I'}$, wherein $R^{H'}$ is selected from the group consisting of

(a1) hydrogen and (b1) C_{1-6} alkyl, and $R^{I'}$ is selected from the group consisting of (a2) C_{1-20} alkyl (e.g., C_{1-6} alkyl), (b2) C_{2-20} alkenyl (e.g., C_{2-6} alkenyl), (c2) C_{6-10} aryl, (d2) hydrogen, (e2) C_{1-6} alk-C_{6-10} aryl, (f2) amino-C_{1-20} alkyl, (g2) polyethylene glycol of-$(CH_2)_{s2}(OCH_2CH_2)_{s1}(CH_2)_{s3}OR'$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and R' is H or C_{1-20} alkyl, and (h2) amino-polyethylene glycol of $-NR^{N1}(CH_2)_{s2}(CH_2CH_2O)_{s1}(CH_2)_{s3}NR^{N1}$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and each R^{N1} is, independently, hydrogen or optionally substituted C_{1-6} alkyl; (20) $-NR^{J'}C(O)OR^{K'}$, wherein $R^{J'}$ is selected from the group consisting of (a1) hydrogen and (b1) C_{1-6} alkyl, and $R^{K'}$ is selected from the group consisting of (a2) C_{1-20} alkyl (e.g., C_{1-6} alkyl), (b2) C_{2-20} alkenyl (e.g., C_{2-6} alkenyl), (c2) C_{6-10} aryl, (d2) hydrogen, (e2) C_{1-6} alk-C_{6-10} aryl, (f2) amino-C_{1-20} alkyl, (g2) polyethylene glycol of-$(CH_2)_{s2}(OCH_2CH_2)_{s1}(CH_2)_{s3}OR'$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and R' is H or C_{1-20} alkyl, and (h2) amino-polyethylene glycol of - $NR^{N1}(CH_2)_{s2}(CH_2CH_2O)_{s1}(CH_2)_{s3}NR^{N1}$, wherein s1 is an integer from 1 to 10 (e.g., from 1 to 6 or from 1 to 4), each of s2 and s3, independently, is an integer from 0 to 10 (e.g., from 0 to 4, from 0 to 6, from 1 to 4, from 1 to 6, or from 1 to 10), and each R^{N1} is, independently, hydrogen or optionally substituted C_{1-6} alkyl; and (21) amidine. In some embodiments, each of these groups can be further substituted as described herein.

[0791] The term "aryl," as used herein, represents a mono-, bicyclic, or multicyclic carbocyclic ring system having one or two aromatic rings and is exemplified by phenyl, naphthyl, 1,2-dihydronaphthyl, 1,2,3,4-tetrahydronaphthyl, anthracenyl, phenanthrenyl, fluorenyl, indanyl, indenyl, and the like, and may be optionally substituted with 1, 2, 3, 4, or 5 substituents independently selected from the group consisting of: (1) C_{1-7} acyl (e.g., carboxyaldehyde); (2) C_{1-20} alkyl (e.g., C_{1-6} alkyl, C_{1-6} alkoxy-C_{1-6} alkyl, C_{1-6} alkylsulfinyl-C_{1-6} alkyl, amino-C_{1-6} alkyl, azido-C_{1-6} alkyl, (carboxyaldehyde)-C_{1-6} alkyl, halo-C_{1-6} alkyl (e.g., perfluoroalkyl), hydroxy-C_{1-6} alkyl, nitro-C_{1-6} alkyl, or C_{1-6} thioalkoxy-C_{1-6} alkyl); (3) C_{1-20} alkoxy (e.g., C_{1-6} alkoxy, such as perfluoroalkoxy); (4) C_{1-6} alkylsulfinyl; (5) C_{6-10} aryl; (6) amino; (7) C_{1-6} alk-C_{6-10} aryl; (8) azido; (9) C_{3-8} cycloalkyl; (10) C_{1-6} alk-C_3-a cycloalkyl; (11) halo; (12) C_{1-12} heterocyclyl (e.g., C_{1-12} heteroaryl); (13) (C_{1-12} heterocyclyl)oxy; (14) hydroxy; (15) nitro; (16) C_{1-20} thioalkoxy (e.g., C_{1-6} thioalkoxy); (17) $-(CH_2)_qCO_2R^{A'}$, where q is an integer from zero to four, and $R^{A'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, (c) hydrogen, and (d) C_{1-6} alk-C_{6-10} aryl; (18) $-(CH_2)_qCONR^{B'}R^{C'}$, where q is an integer from zero to four and where $R^{B'}$ and $R^{C'}$ are independently selected from the group consisting of (a) hydrogen, (b) C_{1-6} alkyl, (c) C_{6-10} aryl, and (d) C_{1-6} alk-C_{6-10} aryl; (19) $-(CH_2)_qSO_2R^{D'}$, where q is an integer from zero to four and where $R^{D'}$ is selected from the group consisting of (a) alkyl, (b) C_{6-10} aryl, and (c) alk-C_{6-10} aryl; (20) $-(CH_2)_qSO_2NR^{E'}R^{F'}$, where q is an integer from zero to four and where each of $R^{E'}$ and $R^{F'}$ is, independently, selected from the group consisting of (a) hydrogen, (b) C_{1-6} alkyl, (c) C_{6-10} aryl, and (d) C_{1-6} alk-C_{6-10} aryl; (21) thiol; (22) C_{6-10} aryloxy; (23) C_{3-8} cycloalkoxy; (24) C_{6-10} aryl-C_{1-6} alkoxy; (25) C_{1-6} alk-C_{1-12} heterocyclyl (e.g., C_{1-6} alk-C_{1-12} heteroaryl); (26) C_{2-20} alkenyl; and (27) C_{2-20} alkynyl. In some embodiments, each of these groups can be further substituted as described herein. For example, the alkylene group of a C_1-alkaryl or a C_1-alkheterocyclyl can be further substituted with an oxo group to afford the respective aryloyl and (heterocyclyl)oyl substituent group.

[0792] The "arylalkyl" group, which as used herein, represents an aryl group, as defined herein, attached to the parent molecular group through an alkylene group, as defined herein. Exemplary unsubstituted arylalkyl groups are from 7 to 30 carbons (e.g., from 7 to 16 or from 7 to 20 carbons, such as C_{1-6} alk-C_{6-10} aryl, C_{1-10} alk-C_{6-10} aryl, or C_{1-20} alk-C_{6-10} aryl). In some embodiments, the alkylene and the aryl each can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein for the respective groups Other groups preceded by the prefix "alk-" are defined in the same manner, where "alk" refers to a C_{1-6} alkylene, unless otherwise noted, and the attached chemical structure is as defined herein.

[0793] The term "azido" represents an $-N_3$ group, which can also be represented as -N=N=N.

[0794] The term "bicyclic," as used herein, refer to a structure having two rings, which may be aromatic or non-aromatic. Bicyclic structures include spirocyclyl groups, as defined herein, and two rings that share one or more bridges, where such bridges can include one atom or a chain including two, three, or more atoms. Exemplary bicyclic groups include a bicyclic carbocyclyl group, where the first and second rings are carbocyclyl groups, as defined herein; a bicyclic aryl groups, where the first and second rings are aryl groups, as defined herein; bicyclic heterocyclyl groups, where the first ring is a heterocyclyl group and the second ring is a carbocyclyl (e.g., aryl) or heterocyclyl (e.g., heteroaryl) group; and bicyclic heteroaryl groups, where the first ring is a heteroaryl group and the second ring is a carbocyclyl (e.g., aryl) or heterocyclyl (e.g., heteroaryl) group. In some embodiments, the bicyclic group can be substituted with 1, 2, 3, or 4 substituents as defined herein for cycloalkyl, heterocyclyl, and aryl groups.

[0795] The term "boranyl," as used herein, represents $-B(R^{B1})_3$, where each R^{B1} is, independently, selected from the group consisting of Hand optionally substituted alkyl. In some embodiments, the boranyl group can be substituted with 1, 2, 3, or 4 substituents as defined herein for alkyl.

[0796] The terms "carbocyclic" and "carbocyclyl," as used herein, refer to an optionally substituted C_{3-12} monocyclic, bicyclic, or tricyclic structure in which the rings, which may be aromatic or non-aromatic, are formed by carbon atoms.

Carbocyclic structures include cycloalkyl, cycloalkenyl, cycloalkynyl, and aryl groups.

[0797] The term "carbonyl," as used herein, represents a C(O) group, which can also be represented as C=O.

[0798] The term "carboxy," as used herein, means $-CO_2H$.

[0799] The term "cyano," as used herein, represents an -CN group.

[0800] The term "cycloalkyl," as used herein represents a monovalent saturated or unsaturated non-aromatic cyclic hydrocarbon group from three to eight carbons, unless otherwise specified, and is exemplified by cyclopropyl, cyclobutyl, cyclopentyl, cyclohexyl, cycloheptyl, bicycle heptyl, and the like. When what would otherwise be a cycloalkyl group includes one or more carbon-carbon double bonds, the group is referred to as a "cycloalkenyl" group. For the purposes of this invention, cycloalkenyl excludes aryl groups. When what would otherwise be a cycloalkyl group includes one or more carbon-carbon triple bonds, the group is referred to as a "cycloalkynyl" group. Exemplary cycloalkenyl groups include cyclopentenyl, cyclohexenyl, and the like. The cycloalkyl groups of this invention can be optionally substituted with: (1) C_{1-7} acyl (e.g., carboxyaldehyde); (2) C_{1-20} alkyl (e.g., C_{1-6} alkyl, C_{1-6} alkoxy-C_{1-6} alkyl, C_{1-6} alkylsulfinyl-C_{1-6} alkyl, amino-C_{1-6} alkyl, azido-C_{1-6} alkyl, (carboxyaldehyde)-C_{1-6} alkyl, halo-C_{1-6} alkyl (e.g., perfluoroalkyl), hydroxy-C_{1-6} alkyl, nitro-C_{1-6} alkyl, or C_{1-6} thioalkoxy-C_{1-6} alkyl); (3) C_{1-20} alkoxy (e.g., C_{1-6} alkoxy, such as perfluoroalkoxy); (4) C_{1-6} alkylsulfinyl; (5) C_{6-10} aryl; (6) amino; (7) C_{1-6} alk-C_{6-10} aryl; (8) azido; (9) C_{3-8} cycloalkyl; (10) C_{1-6} alk-C_{3-8} cycloalkyl; (11) halo; (12) C_{1-12} heterocyclyl (e.g., C_{1-12} heteroaryl); (13) (C_{1-12} heterocyclyl)oxy; (14) hydroxy; (15) nitro; (16) C_{1-20} thioalkoxy (e.g., C_{1-6} thioalkoxy); (17) $-(CH_2)_qCO_2R^{A'}$, where q is an integer from zero to four, and $R^{A'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, (c) hydrogen, and (d) C_{1-6} alk-C_{6-10} aryl; (18) $-(CH_2)_qCONR^{B'}R^{C'}$, where q is an integer from zero to four and where $R^{B'}$ and $R^{C'}$ are independently selected from the group consisting of (a) hydrogen, (b) C_{6-10} alkyl, (c) C_{6-10} aryl, and (d) C_{1-6} alk-C_{6-10} aryl; (19) $-(CH_2)_qSO_2R^{D'}$, where q is an integer from zero to four and where $R^{D'}$ is selected from the group consisting of (a) C_{6-10} alkyl, (b) C_{6-10} aryl, and (c) C_{1-6} alk-C_{6-10} aryl; (20) $-(CH_2)_qSO_2NR^{E'}R^{F'}$, where q is an integer from zero to four and where each of $R^{E'}$ and $R^{F'}$ is, independently, selected from the group consisting of (a) hydrogen, (b) C_{6-10} alkyl, (c) C_{6-10} aryl, and (d) C_{1-6} alk-C_{6-10} aryl; (21) thiol; (22) C_{6-10} aryloxy; (23) C_{3-8} cycloalkoxy; (24) C_{6-10} aryl-C_{1-6} alkoxy; (25) C_{1-6} alk-C_{1-12} heterocyclyl (e.g., C_{1-6} alk-C_{1-12} heteroaryl); (26) oxo; (27) C_{2-20} alkenyl; and (28) C_{2-20} alkynyl. In some embodiments, each of these groups can be further substituted as described herein. For example, the alkylene group of a C_1-alkaryl or a C_1-alkheterocyclyl can be further substituted with an oxo group to afford the respective aryloyl and (heterocyclyl)oyl substituent group.

[0801] The "cycloalkylalkyl" group, which as used herein, represents a cycloalkyl group, as defined herein, attached to the parent molecular group through an alkylene group, as defined herein (e.g., an alkylene group of from 1 to 4, from 1 to 6, from 1 to 10, or form 1 to 20 carbons). In some embodiments, the alkylene and the cycloalkyl each can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein for the respective group.

[0802] The term "halo," as used herein, represents a halogen selected from bromine, chlorine, iodine, or fluorine.

[0803] The term "heteroalkyl," as used herein, refers to an alkyl group, as defined herein, in which one or two of the constituent carbon atoms have each been replaced by nitrogen, oxygen, or sulfur. In some embodiments, the heteroalkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein for alkyl groups. The terms "heteroalkenyl" and heteroalkynyl," as used herein refer to alkenyl and alkynyl groups, as defined herein, respectively, in which one or two of the constituent carbon atoms have each been replaced by nitrogen, oxygen, or sulfur. In some embodiments, the heteroalkenyl and heteroalkynyl groups can be further substituted with 1, 2, 3, or 4 substituent groups as described herein for alkyl groups.

[0804] Non-limiting examples of optionally substituted heteroalkyl, heteroalkenyl, and heteroalkynyl groups include acyloxy, alkenyloxy, alkoxy, alkoxyalkoxy, alkoxycarbonylalkoxy, alkynyloxy, aminoalkoxy, arylalkoxy, carboxyalkoxy, cycloalkoxy, haloalkoxy, (heterocyclyl)oxy, perfluoroalkoxy, thioalkoxy, and thioheterocyclylalkyl:

The "acyloxy" group, which as used herein, represents an acyl group, as defined herein, attached to the parent molecular group though an oxygen atom (i.e., -O-C(O)-R, where R is H or an optionally substituted C_{1-6}, C_{1-10}, or C_{1-20} alkyl group). Exemplary unsubstituted acyloxy groups include from 1 to 21 carbons (e.g., from 1 to 7 or from 1 to 11 carbons). In some embodiments, the alkyl group is further substituted with 1, 2, 3, or 4 substituents as described herein.

[0805] The "alkenyloxy" group, which as used here, represents a chemical substituent of formula -OR, where R is a C_{2-20} alkenyl group (e.g., C_{2-6} or C_{2-10} alkenyl), unless otherwise specified. Exemplary alkenyloxy groups include ethenyloxy, propenyloxy, and the like. In some embodiments, the alkenyl group can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein (e.g., a hydroxy group).

[0806] The "alkoxy" group, which as used herein, represents a chemical substituent of formula-OR, where R is a C_{1-20} alkyl group (e.g., C_{1-6} or C_{1-10} alkyl), unless otherwise specified. Exemplary alkoxy groups include methoxy, ethoxy, propoxy (e.g., n-propoxy and isopropoxy), t-butoxy, and the like. In some embodiments, the alkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein (e.g., hydroxy or alkoxy).

[0807] The "alkoxyalkoxy" group, which as used herein, represents an alkoxy group that is substituted with an alkoxy group. Exemplary unsubstituted alkoxyalkoxy groups include between 2 to 40 carbons (e.g., from 2 to 12 or from 2 to 20 carbons, such as C_{1-6} alkoxy-C_{1-6} alkoxy, C_{1-10} alkoxy-C_{1-10} alkoxy, or C_{1-20} alkoxy-C_{1-20} alkoxy). In some embodiments, the each alkoxy group can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein.

[0808] The "alkoxycarbonylalkoxy" group, which as used herein, represents an alkoxy group, as defined herein, that is substituted with an alkoxycarbonyl group, as defined herein (e.g., -O-alkyl-C(O)-OR, where R is an optionally substituted C_{1-6}, C_{1-10}, or C_{1-20} alkyl group). Exemplary unsubstituted alkoxycarbonylalkoxy include from 3 to 41 carbons (e.g., from 3 to 10, from 3 to 13, from 3 to 17, from 3 to 21, or from 3 to 31 carbons, such as C_{1-6} alkoxycarbonyl-C_{1-6} alkoxy, C_{1-10} alkoxycarbonyl-C_{1-10} alkoxy, or C_{1-20} alkoxycarbonyl-C_{1-20} alkoxy). In some embodiments, each alkoxy group is further independently substituted with 1, 2, 3, or 4 substituents, as described herein (e.g., a hydroxy group).

[0809] The "alkynyloxy" group, which as used herein, represents a chemical substituent of formula -OR, where R is a C_{2-20} alkynyl group (e.g., C_{2-6} or C_{2-10} alkynyl), unless otherwise specified. Exemplary alkynyloxy groups include ethynyloxy, propynyloxy, and the like. In some embodiments, the alkynyl group can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein (e.g., a hydroxy group).

[0810] The "aminoalkoxy" group, which as used herein, represents an alkoxy group, as defined herein, substituted with an amino group, as defined herein. The alkyl and amino each can be further substituted with 1, 2, 3, or 4 substituent groups as described herein for the respective group (e.g., $CO_2R^{A'}$, where $R^{A'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, (c) hydrogen, and (d) C_{1-6} alk-C_{6-10} aryl, e.g., carboxy).

[0811] The "arylalkoxy" group, which as used herein, represents an alkaryl group, as defined herein, attached to the parent molecular group through an oxygen atom. Exemplary unsubstituted arylalkoxy groups include from 7 to 30 carbons (e.g., from 7 to 16 or from 7 to 20 carbons, such as C_{6-10} aryl-C_{1-6} alkoxy, C_{6-10} aryl-C_{1-10} alkoxy, or C_{6-10} aryl-C_{1-20} alkoxy). In some embodiments, the arylalkoxy group can be substituted with 1, 2, 3, or 4 substituents as defined herein.

[0812] The "aryloxy" group, which as used herein, represents a chemical substituent of formula -OR', where R' is an aryl group of 6 to 18 carbons, unless otherwise specified. In some embodiments, the aryl group can be substituted with 1, 2, 3, or 4 substituents as defined herein.

[0813] The "carboxyalkoxy" group, which as used herein, represents an alkoxy group, as defined herein, substituted with a carboxy group, as defined herein. The alkoxy group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein for the alkyl group, and the carboxy group can be optionally substituted with one or more O-protecting groups.

[0814] The "cycloalkoxy" group, which as used herein, represents a chemical substituent of formula -OR, where R is a C_{3-8} cycloalkyl group, as defined herein, unless otherwise specified. The cycloalkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein. Exemplary unsubstituted cycloalkoxy groups are from 3 to 8 carbons. In some embodiment, the cycloalkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein.

[0815] The "haloalkoxy" group, which as used herein, represents an alkoxy group, as defined herein, substituted with a halogen group (i.e., F, Cl, Br, or I). A haloalkoxy may be substituted with one, two, three, or, in the case of alkyl groups of two carbons or more, four halogens. Haloalkoxy groups include perfluoroalkoxys (e.g., $-OCF_3$), $-OCHF_2$, $-OCH_2F$, $-OCCl_3$, $-OCH_2CH_2Br$, $-OCH_2CH(CH_2CH_2Br)CH_3$, and $-OCHICH_3$. In some embodiments, the haloalkoxy group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein for alkyl groups.

[0816] The "(heterocyclyl)oxy" group, which as used herein, represents a heterocyclyl group, as defined herein, attached to the parent molecular group through an oxygen atom. In some embodiments, the heterocyclyl group can be substituted with 1, 2, 3, or 4 substituent groups as defined herein.

[0817] The "perfluoroalkoxy" group, which as used herein, represents an alkoxy group, as defined herein, where each hydrogen radical bound to the alkoxy group has been replaced by a fluoride radical. Perfluoroalkoxy groups are exemplified by trifluoromethoxy, pentafluoroethoxy, and the like.

[0818] The "alkylsulfinyl" group, which as used herein, represents an alkyl group attached to the parent molecular group through an -S(O)-group. Exemplary unsubstituted alkylsulfinyl groups are from 1 to 6, from 1 to 10, or from 1 to 20 carbons. In some embodiments, the alkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein.

[0819] The "thioarylalkyl" group, which as used herein, represents a chemical substituent of formula -SR, where R is an arylalkyl group. In some embodiments, the arylalkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein.

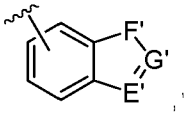
[0820] The "thioalkoxy" group as used herein, represents a chemical substituent of formula -SR, where R is an alkyl group, as defined herein. In some embodiments, the alkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein.

[0821] The "thioheterocyclylalkyl" group, which as used herein, represents a chemical substituent of formula -SR, where R is an heterocyclylalkyl group. In some embodiments, the heterocyclylalkyl group can be further substituted with 1, 2, 3, or 4 substituent groups as described herein.

[0822] The term "heteroaryl," as used herein, represents that subset of heterocyclyls, as defined herein, which are aromatic: i.e., they contain $4n+2$ pi electrons within the mono- or multicyclic ring system. Exemplary unsubstituted heteroaryl groups are of 1 to 12 (e.g., 1 to 11, 1 to 10, 1 to 9, 2 to 12, 2 to 11, 2 to 10, or 2 to 9) carbons. In some embodiment, the heteroaryl is substituted with 1, 2, 3, or 4 substituents groups as defined for a heterocyclyl group.

[0823] The term "heteroarylalkyl" refers to a heteroaryl group, as defined herein, attached to the parent molecular group through an alkylene group, as defined herein. Exemplary unsubstituted heteroarylalkyl groups are from 2 to 32 carbons (e.g., from 2 to 22, from 2 to 18, from 2 to 17, from 2 to 16, from 3 to 15, from 2 to 14, from 2 to 13, or from 2 to 12 carbons, such as C_{1-6} alk-C_{1-12} heteroaryl, C_{1-10} alk-C_{1-12} heteroaryl, or C_{1-20} alk-C_{1-12} heteroaryl). In some embodiments, the alkylene and the heteroaryl each can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein for the respective group. Heteroarylalkyl groups are a subset of heterocyclylalkyl groups.

[0824] The term "heterocyclyl," as used herein represents a 5-, 6- or 7-membered ring, unless otherwise specified, containing one, two, three, or four heteroatoms independently selected from the group consisting of nitrogen, oxygen, and sulfur. The 5-membered ring has zero to two double bonds, and the 6- and 7-membered rings have zero to three double bonds. Exemplary unsubstituted heterocyclyl groups are of 1 to 12 (e.g., 1 to 11, 1 to 10, 1 to 9, 2 to 12, 2 to 11, 2 to 10, or 2 to 9) carbons. The term "heterocyclyl" also represents a heterocyclic compound having a bridged multicyclic structure in which one or more carbons and/or heteroatoms bridges two non-adjacent members of a monocyclic ring, e.g., a quinuclidinyl group. The term "heterocyclyl" includes bicyclic, tricyclic, and tetracyclic groups in which any of the above heterocyclic rings is fused to one, two, or three carbocyclic rings, e.g., an aryl ring, a cyclohexane ring, a cyclohexene ring, a cyclopentane ring, a cyclopentene ring, or another monocyclic heterocyclic ring, such as indolyl, quinolyl, isoquinolyl, tetrahydroquinolyl, benzofuryl, benzothienyl and the like. Examples of fused heterocyclyls include tropanes and 1,2,3,5,8,8a-hexahydroindolizine. Heterocyclics include pyrrolyl, pyrrolinyl, pyrrolidinyl, pyrazolyl, pyrazolinyl, pyrazolidinyl, imidazolyl, imidazolinyl, imidazolidinyl, pyridyl, piperidinyl, homopiperidinyl, pyrazinyl, piperazinyl, pyrimidinyl, pyridazinyl, oxazolyl, oxazolidinyl, isoxazolyl, isoxazolidiniyl, morpholinyl, thiomorpholinyl, thiazolyl, thiazolidinyl, isothiazolyl, isothiazolidinyl, indolyl, indazolyl, quinolyl, isoquinolyl, quinoxalinyl, dihydroquinoxalinyl, quinazolinyl, cinnolinyl, phthalazinyl, benzimidazolyl, benzothiazolyl, benzoxazolyl, benzothiadiazolyl, furyl, thienyl, thiazolidinyl, isothiazolyl, triazolyl, tetrazolyl, oxadiazolyl (e.g., 1,2,3-oxadiazolyl), purinyl, thiadiazolyl (e.g., 1,2,3-thiadiazolyl), tetrahydrofuranyl, dihydrofuranyl, tetrahydrothienyl, dihydrothienyl, dihydroindolyl, dihydroquinolyl, tetrahydroquinolyl, tetrahydroisoquinolyl, dihydroisoquinolyl, pyranyl, dihydropyranyl, dithiazolyl, benzofuranyl, isobenzofuranyl, benzothienyl, and the like, including dihydro and tetrahydro forms thereof, where one or more double bonds are reduced and replaced with hydrogens. Still other exemplary heterocyclyls include: 2,3,4,5-tetrahydro-2-oxo-oxazolyl; 2,3-dihydro-2-oxo-1H-imidazolyl; 2,3,4,5-tetrahydro-5-oxo-1H-pyrazolyl (e.g., 2,3,4,5-tetrahydro-2-phenyl-5-oxo-1H-pyrazolyl); 2,3,4,5-tetrahydro-2,4-dioxo-1H-imidazolyl (e.g., 2,3,4,5-tetrahydro-2,4-dioxo-5-methyl-5-phenyl-1H-imidazolyl); 2,3-dihydro-2-thioxo-1,3,4-oxadiazolyl (e.g., 2,3-dihydro-2-thioxo-5-phenyl-1,3,4-oxadiazolyl); 4,5-dihydro-5-oxo-1H-triazolyl (e.g., 4,5-dihydro-3-methyl-4-amino 5-oxo-1H-triazolyl); 1,2,3,4-tetrahydro-2,4-dioxopyridinyl (e.g., 1,2,3,4-tetrahydro-2,4-dioxo-3,3-diethylpyridinyl); 2,6-dioxo-piperidinyl (e.g., 2,6-dioxo-3-ethyl-3-phenylpiperidinyl); 1,6-dihydro-6-oxopyridiminyl; 1,6-dihydro-4-oxopyrimidinyl (e.g., 2-(methylthio)-1,6-dihydro-4-oxo-5-methylpyrimidin-1-yl); 1,2,3,4-tetrahydro-2,4-dioxopyrimidinyl (e.g., 1,2,3,4-tetrahydro-2,4-dioxo-3-ethylpyrimidinyl); 1,6-dihydro-6-oxo-pyridazinyl (e.g., 1,6-dihydro-6-oxo-3-ethylpyridazinyl); 1,6-dihydro-6-oxo-1,2,4-triazinyl (e.g., 1,6-dihydro-5-isopropyl-6-oxo-1,2,4-triazinyl); 2,3-dihydro-2-oxo-1*H*-indolyl (e.g., 3,3-dimefhyl-2,3-dihydro-2-oxo-1*H*-indolyl and 2,3-dihydro-2-oxo-3,3'-spiropropane-1*H*-indol-1-yl); 1,3-dihydro-1-oxo-2*H*-iso-indolyl; 1,3-dihydro-1,3-dioxo-2*H*-iso-indolyl; 1*H*-benzopyrazolyl (e.g., 1-(ethoxycarbonyl)- 1*H*-benzopyrazolyl); 2,3-dihydro-2-oxo-1*H*-benzimidazolyl (e.g., 3-ethyl-2,3-dihydro-2-oxo-1*H*-benzimidazolyl); 2,3-dihydro-2-oxo-benzoxazolyl (e.g., 5-chloro-2,3-dihydro-2-oxo-benzoxazolyl); 2,3-dihydro-2-oxo-benzoxazolyl; 2-oxo-2H-benzopyranyl; 1,4-benzodioxanyl; 1,3-benzodioxanyl; 2,3-dihydro-3-oxo,4*H*-1,3-benzothiazinyl; 3,4-dihydro-4-oxo-3*H*-quinazolinyl (e.g., 2-methyl-3,4-dihydro-4-oxo-3*H*-quinazolinyl); 1,2,3,4-tetrahydro-2,4-dioxo-3*H*-quinazolyl (e.g., 1-ethyl-1,2,3,4-tetrahydro-2,4-dioxo-3*H*-quinazolyl); 1,2,3,6-tetrahydro-2,6-dioxo-7*H*-purinyl (e.g., 1,2,3,6-tetrahydro-1,3-dimethyl-2,6-dioxo-7 *H* -purinyl); 1,2,3,6-tetrahydro-2,6-dioxo-1 *H*-purinyl (e.g., 1,2,3,6-tetrahydro-3,7-dimethyl-2,6-dioxo-1 *H* -purinyl); 2-oxobenz[*c,d*]indolyl; 1,1-dioxo-2H-naphth[1,8-*c,d*]isothiazolyl; and 1,8-naphthylenedicarboxamido. Additional heterocyclics include 3,3a,4,5,6,6a-hexahydro-pyrrolo[3,4-b]pyrrol-(2H)-yl, and 2,5-diazabicyclo[2.2.1]heptan-2-yl, homopiperazinyl (or diazepanyl), tetrahydropyranyl, dithiazolyl, benzofuranyl, benzothienyl, oxepanyl, thiepanyl, azocanyl, oxecanyl, and thiocanyl. Heterocyclic groups also include groups of the formula



where

E' is selected from the group consisting of -N- and -CH-; F' is selected from the group consisting of -N=CH-, -NH-CH_2-, -NH-C(O)-, -NH-, -CH=N-, -CH_2-NH-, -C(O)-NH-, -CH=CH-, -CH_2-, -CH_2CH_2-, -CH_2O-, -OCH_2-, -O-, and -S-; and G' is selected from the group consisting of -CH- and -N-. Any of the heterocyclyl groups mentioned herein may be optionally substituted with one, two, three, four or five substituents independently selected from the group consisting of: (1) C_{1-7} acyl (e.g., carboxyaldehyde); (2) C_{1-20} alkyl (e.g., C_{1-6} alkyl, C_{1-6} alkoxy-C_{1-6} alkyl, C_{1-6} alkylsulfinyl-C_{1-6} alkyl, amino-

C_{1-6} alkyl, azido-C_{1-6} alkyl, (carboxyaldehyde)-C_{1-6} alkyl, halo-C_{1-6} alkyl (e.g., perfluoroalkyl), hydroxy-C_{1-6} alkyl, nitro-C_{1-6} alkyl, or C_{1-6} thioalkoxy-C_{1-6} alkyl); (3) C_{1-20} alkoxy (e.g., C_{1-6} alkoxy, such as perfluoroalkoxy); (4) C_{1-6} alkylsulfinyl; (5) C_{6-10} aryl; (6) amino; (7) C_{1-6} alk-C_{6-10} aryl; (8) azido; (9) C_{3-8} cycloalkyl; (10) C_{1-6} alk-C_{3-8} cycloalkyl; (11) halo; (12) C_{1-12} heterocyclyl (e.g., C_{2-12} heteroaryl); (13) (C_{1-12} heterocyclyl)oxy; (14) hydroxy; (15) nitro; (16) C_{1-20} thioalkoxy (e.g., C_{1-6} thioalkoxy); (17) -$(CH_2)_qCO_2R^{A'}$, where q is an integer from zero to four, and $R^{A'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, (c) hydrogen, and (d) C_{1-6} alk-C_{6-10} aryl; (18) -$(CH_2)_qCONR^{B'}R^{C'}$, where q is an integer from zero to four and where $R^{B'}$ and $R^{C'}$ are independently selected from the group consisting of (a) hydrogen, (b) C_{1-6} alkyl, (c) C_{6-10} aryl, and (d) C_{1-6} alk-C_{6-10} aryl; (19) -$(CH_2)_qSO_2R^{D'}$, where q is an integer from zero to four and where $R^{D'}$ is selected from the group consisting of (a) C_{1-6} alkyl, (b) C_{6-10} aryl, and (c) C_{1-6} alk-C_{6-10} aryl; (20) -$(CH_2)_qSO_2NR^{E'}R^{F'}$, where q is an integer from zero to four and where each of $R^{E'}$ and $R^{F'}$ is, independently, selected from the group consisting of (a) hydrogen, (b) C_{1-6} alkyl, (c) C_{6-10} aryl, and (d) C_{1-6} alk-C_{6-10} aryl; (21) thiol; (22) C_{6-10} aryloxy; (23) C_{3-8} cycloalkoxy; (24) arylalkoxy; (25) C_{1-6} alk-C_{1-12} heterocyclyl (e.g., C_{1-6} alk-C_{1-12} heteroaryl); (26) oxo; (27) (C_{1-12} heterocyclyl)imino; (28) C_{2-20} alkenyl; and (29) C_{2-20} alkynyl. In some embodiments, each of these groups can be further substituted as described herein. For example, the alkylene group of a C_1-alkaryl or a C_1-alkheterocyclyl can be further substituted with an oxo group to afford the respective aryloyl and (heterocyclyl)oyl substituent group.

[0825] The "heterocyclylalkyl" group, which as used herein, represents a heterocyclyl group, as defined herein, attached to the parent molecular group through an alkylene group, as defined herein. Exemplary unsubstituted heterocyclylalkyl groups are from 2 to 32 carbons (e.g., from 2 to 22, from 2 to 18, from 2 to 17, from 2 to 16, from 3 to 15, from 2 to 14, from 2 to 13, or from 2 to 12 carbons, such as C_{1-6} alk-C_{1-12} heterocyclyl, C_{1-10} alk-C_{1-12} heterocyclyl, or C_{1-20} alk-C_{1-12} heterocyclyl). In some embodiments, the alkylene and the heterocyclyl each can be further substituted with 1, 2, 3, or 4 substituent groups as defined herein for the respective group.

[0826] The term "hydrocarbon," as used herein, represents a group consisting only of carbon and hydrogen atoms.

[0827] The term "hydroxy," as used herein, represents an -OH group.

[0828] The term "*N*-protected amino," as used herein, refers to an amino group, as defined herein, to which is attached one or two N-protecting groups, as defined herein.

[0829] The term "*N*-protecting group," as used herein, represents those groups intended to protect an amino group against undesirable reactions during synthetic procedures. Commonly used *N*-protecting groups are disclosed in Greene, "Protective Groups in Organic Synthesis," 3rd Edition (John Wiley & Sons, New York, 1999). *N*-protecting groups include acyl, aryloyl, or carbamyl groups such as formyl, acetyl, propionyl, pivaloyl, t-butylacetyl, 2-chloroacetyl, 2-bromoacetyl, trifluoroacetyl, trichloroacetyl, phthalyl, o-nitrophenoxyacetyl, α-chlorobutyryl, benzoyl, 4-chlorobenzoyl, 4-bromobenzoyl, 4-nitrobenzoyl, and chiral auxiliaries such as protected or unprotected D, L or D, L-amino acids such as alanine, leucine, phenylalanine, and the like; sulfonyl-containing groups such as benzenesulfonyl, p-toluenesulfonyl, and the like; carbamate forming groups such as benzyloxycarbonyl, p-chlorobenzyloxycarbonyl, p-methoxybenzyloxycarbonyl, p-nitrobenzyloxycarbonyl, 2-nitrobenzyloxycarbonyl, p-bromobenzyloxycarbonyl, 3,4-dimethoxybenzyloxycarbonyl, 3,5-dimethoxybenzyloxycarbonyl, 2,4-dimethoxybenzyloxycarbonyl, 4-methoxybenzyloxycarbonyl, 2-nitro-4,5-dimethoxybenzyloxycarbonyl, 3,4,5-trimethoxybenzyloxycarbonyl, 1-(p-biphenylyl)-1-methylethoxycarbonyl, o,o-dimethyl-3,5-dimethoxybenzyloxycarbonyl, benzhydryloxy carbonyl, t-butyloxycarbonyl, diisopropylmethoxycarbonyl, isopropyloxycarbonyl, ethoxycarbonyl, methoxycarbonyl, allyloxycarbonyl, 2,2,2,-trichloroethoxycarbonyl, phenoxycarbonyl, 4-nitrophenoxy carbonyl, fluorenyl-9-methoxycarbonyl, cyclopentyloxycarbonyl, adamantyloxycarbonyl, cyclohexyloxycarbonyl, phenylthiocarbonyl, and the like, alkaryl groups such as benzyl, triphenylmethyl, benzyloxymethyl, and the like and silyl groups, such as trimethylsilyl, and the like. Preferred N-protecting groups are formyl, acetyl, benzoyl, pivaloyl, t-butylacetyl, alanyl, phenylsulfonyl, benzyl, t-butyloxycarbonyl (Boc), and benzyloxycarbonyl (Cbz).

[0830] The term "nitro," as used herein, represents an -NO_2 group.

[0831] The term "O-protecting group," as used herein, represents those groups intended to protect an oxygen containing (e.g., phenol, hydroxyl, or carbonyl) group against undesirable reactions during synthetic procedures. Commonly used O-protecting groups are disclosed in Greene, "Protective Groups in Organic Synthesis," 3rd Edition (John Wiley & Sons, New York, 1999). Exemplary O-protecting groups include acyl, aryloyl, or carbamyl groups, such as formyl, acetyl, propionyl, pivaloyl, t-butylacetyl, 2-chloroacetyl, 2-bromoacetyl, trifluoroacetyl, trichloroacetyl, phthalyl, o-nitrophenoxyacetyl, α-chlorobutyryl, benzoyl, 4-chlorobenzoyl, 4-bromobenzoyl, t-butyldimethylsilyl, tri-*iso*-propylsilyloxymethyl, 4,4'-dimethoxytrityl, isobutyryl, phenoxyacetyl, 4-isopropylpehenoxyacetyl, dimethylformamidino, and 4-nitrobenzoyl; alkylcarbonyl groups, such as acyl, acetyl, propionyl, pivaloyl, and the like; optionally substituted arylcarbonyl groups, such as benzoyl; silyl groups, such as trimethylsilyl (TMS), tert-butyldimethylsilyl (TBDMS), tri-iso-propylsilyloxymethyl (TOM), triisopropylsilyl (TIPS), and the like; ether-forming groups with the hydroxyl, such methyl, methoxymethyl, tetrahydropyranyl, benzyl, p-methoxybenzyl, trityl, and the like; alkoxycarbonyls, such as methoxycarbonyl, ethoxycarbonyl, isopropoxycarbonyl, n-isopropoxycarbonyl, n-butyloxycarbonyl, isobutyloxycarbonyl, sec-butyloxycarbonyl, t-butyloxycarbonyl, 2-ethylhexyloxycarbonyl, cyclohexyloxycarbonyl, methyloxycarbonyl, and the like; alkoxyalkoxycarbonyl groups, such as methoxymethoxycarbonyl, ethoxymethoxycarbonyl, 2-methoxyethoxycarbonyl, 2-ethoxyethoxycarbonyl, 2-butoxyethoxycarbonyl, 2-methoxyethoxymethoxycarbonyl, allyloxycarbonyl, propargyloxycarbonyl, 2-butenoxycarbonyl,

3-methyl-2-butenoxycarbonyl, and the like; haloalkoxycarbonyls, such as 2-chloroethoxycarbonyl, 2-chloroethoxycarbonyl, 2,2,2-trichloroethoxycarbonyl, and the like; optionally substituted arylalkoxycarbonyl groups, such as benzyloxycarbonyl, p-methylbenzyloxycarbonyl, p-methoxybenzyloxycarbonyl, p-nitrobenzyloxycarbonyl, 2,4-dinitrobenzyloxycarbonyl, 3,5-dimethylbenzyloxycarbonyl, p-chlorobenzyloxycarbonyl, p-bromobenzyloxy-carbonyl, fluorenylmethyloxycarbonyl, and the like; and optionally substituted aryloxycarbonyl groups, such as phenoxycarbonyl, p-nitrophenoxycarbonyl, o-nitrophenoxycarbonyl, 2,4-dinitrophenoxycarbonyl, p-methyl-phenoxycarbonyl, m-methylphenoxycarbonyl, o-bromophenoxycarbonyl, 3,5-dimethylphenoxycarbonyl, p-chlorophenoxycarbonyl, 2-chloro-4-nitrophenoxy-carbonyl, and the like); substituted alkyl, aryl, and alkaryl ethers (e.g., trityl; methylthiomethyl; methoxymethyl; benzyloxymethyl; siloxymethyl; 2,2,2,-trichloroethoxymethyl; tetrahydropyranyl; tetrahydrofuranyl; ethoxyethyl; 1-[2-(trimethylsilyl)ethoxy]ethyl; 2-trimethylsilylethyl; t-butyl ether, p-chlorophenyl, p-methoxyphenyl, p-nitrophenyl, benzyl, p-methoxybenzyl, and nitrobenzyl); silyl ethers (e.g., trimethylsilyl; triethylsilyl; triisopropylsilyl; dimethylisopropylsilyl; t-butyldimethylsilyl; t-butyldiphenylsilyl; tribenzylsilyl; triphenylsilyl; and diphenymethylsilyl); carbonates (e.g., methyl, methoxymethyl, 9-fluorenylmethyl; ethyl; 2,2,2-trichloroethyl; 2-(trimethylsilyl)ethyl; vinyl, allyl, nitrophenyl; benzyl; methoxybenzyl; 3,4-dimethoxybenzyl; and nitrobenzyl); carbonyl-protecting groups (e.g., acetal and ketal groups, such as dimethyl acetal, 1,3-dioxolane, and the like; acylal groups, and dithiane groups, such as 1,3-dithianes, 1,3-dithiolane, and the like); carboxylic acid-protecting groups (e.g., ester groups, such as methyl ester, benzyl ester, t-butyl ester, orthoesters, and the like; and oxazoline groups.

[0832] The term "oxo" as used herein, represents =O.

[0833] The prefix "perfluoro," as used herein, represents anyl group, as defined herein, where each hydrogen radical bound to the alkyl group has been replaced by a fluoride radical. For example, perfluoroalkyl groups are exemplified by trifluoromethyl, pentafluoroethyl, and the like.

[0834] The term "protected hydroxyl," as used herein, refers to an oxygen atom bound to an O-protecting group.

[0835] The term "spirocyclyl," as used herein, represents a C_{2-7} alkylene diradical, both ends of which are bonded to the same carbon atom of the parent group to form a spirocyclic group, and also a C_{1-6} heteroalkylene diradical, both ends of which are bonded to the same atom. The heteroalkylene radical forming the spirocyclyl group can containing one, two, three, or four heteroatoms independently selected from the group consisting of nitrogen, oxygen, and sulfur. In some embodiments, the spirocyclyl group includes one to seven carbons, excluding the carbon atom to which the diradical is attached. The spirocyclyl groups of the invention may be optionally substituted with 1, 2, 3, or 4 substituents provided herein as optional substituents for cycloalkyl and/or heterocyclyl groups.

[0836] The term "sulfonyl," as used herein, represents an $-S(O)_2-$ group.

[0837] The term "thiol," as used herein represents an -SH group.

[0838] *Chimera:* As used herein, "chimera" is an entity having two or more incongruous or heterogeneous parts or regions.

[0839] *Chimeric polynucleotide:* As used herein, "chimeric polynucleotides" are those nucleic acid polymers having portions or regions which differ in size and/or chemical modification pattern, chemical modification position, chemical modification percent or chemical modification population and combinations of the foregoing.

[0840] *Compound:* As used herein, the term "compound," is meant to include all stereoisomers, geometric isomers, tautomers, and isotopes of the structures depicted.

[0841] The compounds described herein can be asymmetric (*e.g.,* having one or more stereocenters). All stereoisomers, such as enantiomers and diastereomers, are intended unless otherwise indicated. Compounds of the present disclosure that contain asymmetrically substituted carbon atoms can be isolated in optically active or racemic forms. Methods on how to prepare optically active forms from optically active starting materials are known in the art, such as by resolution of racemic mixtures or by stereoselective synthesis. Many geometric isomers of olefins, C=N double bonds, and the like can also be present in the compounds described herein, and all such stable isomers are contemplated in the present disclosure. Cis and trans geometric isomers of the compounds of the present disclosure are described and may be isolated as a mixture of isomers or as separated isomeric forms.

[0842] Compounds of the present disclosure also include tautomeric forms. Tautomeric forms result from the swapping of a single bond with an adjacent double bond and the concomitant migration of a proton. Tautomeric forms include prototropic tautomers which are isomeric protonation states having the same empirical formula and total charge. Examples prototropic tautomers include ketone - enol pairs, amide - imidic acid pairs, lactam - lactim pairs, amide - imidic acid pairs, enamine - imine pairs, and annular forms where a proton can occupy two or more positions of a heterocyclic system, such as, 1H- and 3H-imidazole, 1H-, 2H- and 4H-1,2,4-triazole, 1H- and 2H- isoindole, and 1H- and 2H-pyrazole. Tautomeric forms can be in equilibrium or sterically locked into one form by appropriate substitution.

[0843] Compounds of the present disclosure also include all of the isotopes of the atoms occurring in the intermediate or final compounds. "Isotopes" refers to atoms having the same atomic number but different mass numbers resulting from a different number of neutrons in the nuclei. For example, isotopes of hydrogen include tritium and deuterium.

[0844] The compounds and salts of the present disclosure can be prepared in combination with solvent or water molecules to form solvates and hydrates by routine methods.

[0845] *Conserved:* As used herein, the term "conserved" refers to nucleotides or amino acid residues of a polynucleotide sequence or polypeptide sequence, respectively, that are those that occur unaltered in the same position of two or more sequences being compared. Nucleotides or amino acids that are relatively conserved are those that are conserved amongst more related sequences than nucleotides or amino acids appearing elsewhere in the sequences.

[0846] In some embodiments, two or more sequences are said to be "completely conserved" if they are 100% identical to one another. In some embodiments, two or more sequences are said to be "highly conserved" if they are at least 70% identical, at least 80% identical, at least 90% identical, or at least 95% identical to one another. In some embodiments, two or more sequences are said to be "highly conserved" if they are about 70% identical, about 80% identical, about 90% identical, about 95%, about 98%, or about 99% identical to one another. In some embodiments, two or more sequences are said to be "conserved" if they are at least 30% identical, at least 40% identical, at least 50% identical, at least 60% identical, at least 70% identical, at least 80% identical, at least 90% identical, or at least 95% identical to one another. In some embodiments, two or more sequences are said to be "conserved" if they are about 30% identical, about 40% identical, about 50% identical, about 60% identical, about 70% identical, about 80% identical, about 90% identical, about 95% identical, about 98% identical, or about 99% identical to one another. Conservation of sequence may apply to the entire length of an polynucleotide or polypeptide or may apply to a portion, region or feature thereof.

[0847] *Controlled Release:* As used herein, the term "controlled release" refers to a pharmaceutical composition or compound release profile that conforms to a particular pattern of release to effect a therapeutic outcome.

[0848] *Cyclic or Cyclized*: As used herein, the term "cyclic" refers to the presence of a continuous loop. Cyclic molecules need not be circular, only joined to form an unbroken chain of subunits. Cyclic molecules such as the engineered RNA or mRNA of the present invention may be single units or multimers or comprise one or more components of a complex or higher order structure.

[0849] *Cytostatic:* As used herein, "cytostatic" refers to inhibiting, reducing, suppressing the growth, division, or multiplication of a cell *(e.g.,* a mammalian cell *(e.g.,* a human cell)), bacterium, virus, fungus, protozoan, parasite, prion, or a combination thereof.

[0850] *Cytotoxic*: As used herein, "cytotoxic" refers to killing or causing injurious, toxic, or deadly effect on a cell *(e.g.,* a mammalian cell *(e.g.,* a human cell)), bacterium, virus, fungus, protozoan, parasite, prion, or a combination thereof.

[0851] *Delivery:* As used herein, "delivery" refers to the act or manner of delivering a compound, substance, entity, moiety, cargo or payload.

[0852] *Delivery Agent:* As used herein, "delivery agent" refers to any substance which facilitates, at least in part, *the in vivo* delivery of a polynucleotide to targeted cells.

[0853] *Destabilized:* As used herein, the term "destable," "destabilize," or "destabilizing region" means a region or molecule that is less stable than a starting, wild-type or native form of the same region or molecule.

[0854] *Detectable label:* As used herein, "detectable label" refers to one or more markers, signals, or moieties which are attached, incorporated or associated with another entity that is readily detected by methods known in the art including radiography, fluorescence, chemiluminescence, enzymatic activity, absorbance and the like. Detectable labels include radioisotopes, fluorophores, chromophores, enzymes, dyes, metal ions, ligands such as biotin, avidin, streptavidin and haptens, quantum dots, and the like. Detectable labels may be located at any position in the peptides or proteins disclosed herein. They may be within the amino acids, the peptides, or proteins, or located at the N- or C- termini.

[0855] *Diastereomer* As used herein, the term "diastereomer," means stereoisomers that are not mirror images of one another and are non-superimposable on one another.

[0856] *Digest:* As used herein, the term "digest" means to break apart into smaller pieces or components. When referring to polypeptides or proteins, digestion results in the production of peptides.

[0857] *Differentiated cell:* As used herein, the term "differentiated cell" refers to any somatic cell that is not, in its native form, pluripotent. Differentiated cell also encompasses cells that are partially differentiated.

[0858] *Differentiation:* As used herein, the term "differentiation factor" refers to a developmental potential altering factor such as a protein, RNA or small molecule that can induce a cell to differentiate to a desired cell-type.

[0859] *Differentiate:* As used herein, "differentiate" refers to the process where an uncommitted or less committed cell acquires the features of a committed cell.

[0860] *Distal:* As used herein, the term "distal" means situated away from the center or away from a point or region of interest.

[0861] *Dosing regimen:* As used herein, a "dosing regimen" is a schedule of administration or physician determined regimen of treatment, prophylaxis, or palliative care.

[0862] *Dose splitting factor (DSF)*-ratio of PUD of dose split treatment divided by PUD of total daily dose or single unit dose. The value is derived from comparison of dosing regimens groups.

[0863] *Enantiomer* As used herein, the term "enantiomer" means each individual optically active form of a compound of the invention, having an optical purity or enantiomeric excess (as determined by methods standard in the art) of at least 80% (i.e., at least 90% of one enantiomer and at most 10% of the other enantiomer), preferably at least 90% and more preferably at least 98%.

[0864] *Encapsulate:* As used herein, the term "encapsulate" means to enclose, surround or encase.

[0865] *Encoded protein cleavage signal:* As used herein, "encoded protein cleavage signal" refers to the nucleotide sequence which encodes a protein cleavage signal.

[0866] *Engineered:* As used herein, embodiments of the invention are "engineered" when they are designed to have a feature or property, whether structural or chemical, that varies from a starting point, wild type or native molecule.

[0867] *Effective Amount:* As used herein, the term "effective amount" of an agentis that amount sufficient to effect beneficial or desired results, for example, clinical results, and, as such, an "effective amount" depends upon the context in which it is being applied. For example, in the context of administering an agent that treats cancer, an effective amount of an agent is, for example, an amount sufficient to achieve treatment, as defined herein, of cancer, as compared to the response obtained without administration of the agent.

[0868] *Exosome:* As used herein, "exosome" is a vesicle secreted by mammalian cells or a complex involved in RNA degradation.

[0869] *Expression:* As used herein, "expression" of a nucleic acid sequence refers to one or more of the following events: (1) production of an RNA template from a DNA sequence *(e.g.,* by transcription); (2) processing of an RNA transcript *(e.g.,* by splicing, editing, 5' cap formation, and/or 3' end processing); (3) translation of an RNA into a polypeptide or protein; and (4) post-translational modification of a polypeptide or protein.

[0870] *Feature:* As used herein, a "feature" refers to a characteristic, a property, or a distinctive element.

[0871] *Formulation*: As used herein, a "formulation" includes at least a polynucleotide of a NAV and a delivery agent.

[0872] *Fragment:* A "fragment," as used herein, refers to a portion. For example, fragments of proteins may comprise polypeptides obtained by digesting full-length protein isolated from cultured cells.

[0873] *Functional:* As used herein, a "functional" biological molecule is a biological molecule in a form in which it exhibits a property and/or activity by which it is characterized.

[0874] *Homology:* As used herein, the term "homology" refers to the overall relatedness between polymeric molecules, *e.g.* between nucleic acid molecules *(e.g.* DNA molecules and/or RNA molecules) and/or between polypeptide molecules. In some embodiments, polymeric molecules are considered to be "homologous" to one another if their sequences are at least 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95%, or 99% identical or similar. The term "homologous" necessarily refers to a comparison between at least two sequences (polynucleotide or polypeptide sequences). In accordance with the invention, two polynucleotide sequences are considered to be homologous if the polypeptides they encode are at least about 50%, 60%, 70%, 80%, 90%, 95%, or even 99% for at least one stretch of at least about 20 amino acids. In some embodiments, homologous polynucleotide sequences are characterized by the ability to encode a stretch of at least 4-5 uniquely specified amino acids. For polynucleotide sequences less than 60 nucleotides in length, homology is determined by the ability to encode a stretch of at least 4-5 uniquely specified amino acids. In accordance with the invention, two protein sequences are considered to be homologous if the proteins are at least about 50%, 60%, 70%, 80%, or 90% identical for at least one stretch of at least about 20 amino acids.

[0875] *Identity:* As used herein, the term "identity" refers to the overall relatedness between polymeric molecules, *e.g.,* between polynucleotide molecules (*e.g.* DNA molecules and/or RNA molecules) and/or between polypeptide molecules. Calculation of the percent identity of two polynucleotide sequences, for example, can be performed by aligning the two sequences for optimal comparison purposes (*e.g.,* gaps can be introduced in one or both of a first and a second nucleic acid sequences for optimal alignment and non-identical sequences can be disregarded for comparison purposes). In certain embodiments, the length of a sequence aligned for comparison purposes is at least 30%, at least 40%, at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, or 100% of the length of the reference sequence. The nucleotides at corresponding nucleotide positions are then compared. When a position in the first sequence is occupied by the same nucleotide as the corresponding position in the second sequence, then the molecules are identical at that position. The percent identity between the two sequences is a function of the number of identical positions shared by the sequences, taking into account the number of gaps, and the length of each gap, which needs to be introduced for optimal alignment of the two sequences The comparison of sequences and determination of percent identity between two sequences can be accomplished using a mathematical algorithm. For example, the percent identity between two nucleotide sequences can be determined using methods such as those described in Computational Molecular Biology, Lesk, A. M., ed., Oxford University Press, New York, 1988; Biocomputing: Informatics and Genome Projects, Smith, D. W., ed., Academic Press, New York, 1993; Sequence Analysis in Molecular Biology, von Heinje, G., Academic Press, 1987; Computer Analysis of Sequence Data, Part I, Griffin, A. M., and Griffin, H. G., eds, Humana Press, New Jersey, 1994; and Sequence Analysis Primer, Gribskov, M. and Devereux, J., eds., M Stockton Press, New York, 1991. For example, the percent identity between two nucleotide sequences can be determined using the algorithm of Meyers and Miller (CABIOS, 1989, 4:11-17), which has been incorporated into the ALIGN program (version 2.0) using a PAM120 weight residue table, a gap length penalty of 12 and a gap penalty of 4 The percent identity between two nucleotide sequences can, alternatively, be determined using the GAP program in the GCG software package using an NWSgapdna.CMP matrix. Methods commonly employed to determine percent identity between sequences include, but are not limited to those disclosed in Carillo, H., and Lipman, D., SIAM J Applied Math, 48:1073 (1988). Techniques for

determining identity are codified in publicly available computer programs. Exemplary computer software to determine homology between two sequences include, but are not limited to, GCG program package, Devereux, J., et al., Nucleic Acids Research, 12(1), 387 (1984)), BLASTP, BLASTN, and FASTA Altschul, S. F. et al., J. Molec. Biol., 215, 403 (1990)).

[0876] *Infectious Agent:* As used herein, the phrase "infectious agent" means an agent capable of producing an infection in an organism, for example, in an animal. An infectious agent may refer to any microorganism, virus, infectious substance, or biological product that may be engineered as a result of biotechnology, or any naturally occurring or bioengineered component of any such microorganism, virus, infectious substance, or biological product, can cause emerging and contagious disease, death or other biological malfunction in a human, an animal, a plant or another living organism.

[0877] *Inhibit expression of a gene:* As used herein, the phrase "inhibit expression of a gene" means to cause a reduction in the amount of an expression product of the gene. The expression product can be an RNA transcribed from the gene (*e.g.,* an mRNA) or a polypeptide translated from an mRNA transcribed from the gene. Typically a reduction in the level of an mRNA results in a reduction in the level of a polypeptide translated therefrom. The level of expression may be determined using standard techniques for measuring mRNA or protein.

[0878] *Influenza:* As used herein, "influenza" or "flu" is an infectious disease of birds and mammals caused by RNA viruses of the family Orthomyxoviridae, the influenza viruses.

[0879] Isomer As used herein, the term "isomer" means any tautomer, stereoisomer, enantiomer, or diastereomer of any compound of the invention. It is recognized that the compounds of the invention can have one or more chiral centers and/or double bonds and, therefore, exist as stereoisomers, such as double-bond isomers (i.e., geometric E/Z isomers) or diastereomers (e.g., enantiomers (i.e., (+) or (-)) or cis/trans isomers). According to the invention, the chemical structures depicted herein, and therefore the compounds of the invention, encompass all of the corresponding stereoisomers, that is, both the stereomerically pure form (e.g., geometrically pure, enantiomerically pure, or diastereomerically pure) and enantiomeric and stereoisomeric mixtures, e.g., racemates. Enantiomeric and stereoisomeric mixtures of compounds of the invention can typically be resolved into their component enantiomers or stereoisomers by well-known methods, such as chiral-phase gas chromatography, chiral-phase high performance liquid chromatography, crystallizing the compound as a chiral salt complex, or crystallizing the compound in a chiral solvent. Enantiomers and stereoisomers can also be obtained from stereomerically or enantiomerically pure intermediates, reagents, and catalysts by well-known asymmetric synthetic methods.

[0880] *In vitro:* As used herein, the term *"in vitro"* refers to events that occur in an artificial environment, *e.g.,* in a test tube or reaction vessel, in cell culture, in a Petri dish, *etc.,* rather than within an organism (*e.g.,* animal, plant, or microbe).

[0881] *In vivo:* As used herein, the term *"in vivo"* refers to events that occur within an organism (*e.g.,* animal, plant, or microbe or cell or tissue thereof).

[0882] *Isolated:* As used herein, the term "isolated" refers to a substance or entity that has been separated from at least some of the components with which it was associated (whether in nature or in an experimental setting). Isolated substances may have varying levels of purity in reference to the substances from which they have been associated. Isolated substances and/or entities may be separated from at least about 10%, about 20%, about 30%, about 40%, about 50%, about 60%, about 70%, about 80%, about 90%, or more of the other components with which they were initially associated In some embodiments, isolated agents are more than about 80%, about 85%, about 90%, about 91%, about 92%, about 93%, about 94%, about 95%, about 96%, about 97%, about 98%, about 99%, or more than about 99% pure. As used herein, a substance is "pure" if it is substantially free of other components. *Substantially isolated:* By "substantially isolated" is meant that the compound is substantially separated from the environment in which if was formed or detected. Partial separation can include, for example, a composition enriched in the compound of the present disclosure. Substantial separation can include compositions containing at least about 50%, at least about 60%, at least about 70%, at least about 80%, at least about 90%, at least about 95%, at least about 97%, or at least about 99% by weight of the compound of the present disclosure, or salt thereof. Methods for isolating compounds and their salts are routine in the art.

[0883] *IVT Polynucleotide:* As used herein, an "IVT polynucleotide" is a linear polynucleotide which may be made using *in vitro* transcription (IVT) enzymatic synthesis methods.

[0884] *Linker* As used herein, a "linker" refers to a group of atoms, e.g., 10-1,000 atoms, and can be comprised of the atoms or groups such as, but not limited to, carbon, amino, alkylamino, oxygen, sulfur, sulfoxide, sulfonyl, carbonyl, and imine. The linker can be attached to a modified nucleoside or nucleotide on the nucleobase or sugar moiety at a first end, and to a payload, e.g., a detectable or therapeutic agent, at a second end. The linker may be of sufficient length as to not interfere with incorporation into a nucleic acid sequence. The linker can be used for any useful purpose, such as to form polynucleotide multimers (e.g., through linkage of two or more chimeric polynucleotides molecules or IVT polynucleoties) or polynucleotides conjugates, as well as to administer a payload, as described herein. Examples of chemical groups that can be incorporated into the linker include, but are not limited to, alkyl, alkenyl, alkynyl, amido, amino, ether, thioether, ester, alkylene, heteroalkylene, aryl, or heterocyclyl, each of which can be optionally substituted, as described herein. Examples of linkers include, but are not limited to, unsaturated alkanes, polyethylene glycols (e.g.,

ethylene or propylene glycol monomeric units, e.g., diethylene glycol, dipropylene glycol, triethylene glycol, tripropylene glycol, tetraethylene glycol, or tetraethylene glycol), and dextran polymers and derivatives thereof, Other examples include, but are not limited to, cleavable moieties within the linker, such as, for example, a disulfide bond (-S-S-) or an azo bond (-N=N-), which can be cleaved using a reducing agent or photolysis. Non-limiting examples of a selectively cleavable bond include an amido bond can be cleaved for example by the use of tris(2-carboxyethyl)phosphine (TCEP), or other reducing agents, and/or photolysis, as well as an ester bond can be cleaved for example by acidic or basic hydrolysis.

[0885] *MicroRNA (miRNA) binding site:* As used herein, a microRNA (miRNA) binding site represents a nucleotide location or region of a nucleic acid transcript to which at least the "seed" region of a miRNA binds.

[0886] *Modified:* As used herein "modified" refers to a changed state or structure of a molecule of the invention. Molecules may be modified in many ways including chemically, structurally, and functionally. In one embodiment, the polynucleotide molecules of the present invention are modified by the introduction of non-natural nucleosides and/or nucleotides, e.g., as it relates to the natural ribonucleotides A, U, G, and C. Noncanonical nucleotides such as the cap structures are not considered "modified" although they differ from the chemical structure of the A, C, G, U ribonucleotides.

[0887] *Mucus:* As used herein, "mucus" refers to the natural substance that is viscous and comprises mucin glycoproteins.

[0888] *Naturally occurring:* As used herein, "naturally occurring" means existing in nature without artificial aid.

[0889] *Neutralizing antibody:* As used herein, a "neutralizing antibody" refers to an antibody which binds to its antigen and defends a cell from an antigen or infectious agent by neutralizing or abolishing any biological activity it has

[0890] *Non-human vertebrate:* As used herein, a "non human vertebrate" includes all vertebrates except *Homo sapiens,* including wild and domesticated species. Examples of non-human vertebrates include, but are not limited to, mammals, such as alpaca, banteng, bison, camel, cat, cattle, deer, dog, donkey, gayal, goat, guinea pig, horse, llama, mule, pig, rabbit, reindeer, sheep wafer buffalo, and yak.

[0891] *Nucleic Acid Vaccine:* As used herein, "nucleic acid vaccine" or "NAV" or "NAV composition" refers to a vaccine or vaccine composition which includes a nucleic acid or nucleic acid molecule (e.g., a polynucleotide) encoding an antigen (e.g., an antigenic protein or polypeptide) In exemplary instances, a nucleic acid vaccine or NAV includes a ribonucleic ("RNA") polynucleotide, ribonucleic acid ("RNA") or ribonucleic acid ("RNA") molecule. Such instancescan be referred to as ribonucleic acid ("RNA") vaccines (RNAVs). In preferred instances, a nucleic acid vaccine or NAV includes a messenger RNA ("mRNA") polynucleotide, messenger RNA ("mRNA") or messenger RNA ("mRNA") molecule as described in detail herein. Such instancescan be referred to as messenger RNA ("mRNA") vaccines (mRNAVs). References herein to "NAVs of the invention" relate to those as set out in the appended claims.

[0892] *Off-target:* As used herein, "off target" refers to any unintended effect on any one or more target, gene, or cellular transcript.

[0893] *Open reading frame:* As used herein, the term "open reading frame" or "ORF" refers to a continuous polynucleotide sequence, for example, a DNA sequence or RNA sequence (*e.g.,* an mRNA sequence), comprising a start codon, a subsequent region comprising a plurality of amino acid-encoding codons, and a terminal stop codon, wherein the region comprising the plurality of amino acid-encoding codons contains no stop codons.

[0894] *Operably linked:* As used herein, the phrase "operably linked" refers to a functional connection between two or more molecules, constructs, transcripts, entities, moieties or the like.

[0895] *Optionally substituted:* Herein a phrase of the form "optionally substituted X" (*e.g.,* optionally substituted alkyl) is intended to be equivalent to "X, wherein X is optionally substituted" (*e.g.,* "alkyl, wherein said alkyl is optionally substituted"). It is not intended to mean that the feature "X" (*e.g.* alkyl) *per se* is optional.

[0896] *Part:* As used herein, a "part" or "region" of a polynucleotide is defined as any portion of the polynucleotide which is less than the entire length of the polynucleotide.

[0897] *Peptide:* As used herein, "peptide" is less than or equal to 50 amino acids long, e.g., about 5, 10, 15, 20, 25, 30, 35, 40, 45, or 50 amino acids long.

[0898] *Paratope:* As used herein, a "paratope" refers to the antigen-binding site of an antibody.

[0899] *Patient:* As used herein, "patient" refers to a subject who may seek or be in need of treatment, requires treatment, is receiving treatment, will receive treatment, or a subject who is under care by a trained professional for a particular disease or condition.

[0900] *Pharmaceutically acceptable:* The phrase "pharmaceutically acceptable" is employed herein to refer to those compounds, materials, compositions, and/or dosage forms which are, within the scope of sound medical judgment, suitable for use in contact with the tissues of human beings and animals without excessive toxicity, irritation, allergic response, or other problem or complication, commensurate with a reasonable benefit/risk ratio.

[0901] *Pharmaceutically acceptable excipients:* The phrase "pharmaceutically acceptable excipient," as used herein, refers any ingredient other than the compounds described herein (for example, a vehicle capable of suspending or dissolving the active compound) and having the properties of being substantially nontoxic and non-inflammatory in a patient. Excipients may include, for example: antiadherents, antioxidants, binders, coatings, compression aids, disinte-

grants, dyes (colors), emollients, emulsifiers, fillers (diluents), film formers or coatings, flavors, fragrances, glidants (flow enhancers), lubricants, preservatives, printing inks, sorbents, suspensing or dispersing agents, sweeteners, and wafers of hydration. Exemplary excipients include, but are not limited to: butylated hydroxytoluene (BHT), calcium carbonate, calcium phosphate (dibasic), calcium stearate, croscarmellose, crosslinked polyvinyl pyrrolidone, citric acid, crospovidone, cysteine, ethylcellulose, gelatin, hydroxypropyl cellulose, hydroxypropyl methylcellulose, lactose, magnesium stearate, maltitol, mannitol, methionine, methylcellulose, methyl paraben, microcrystalline cellulose, polyethylene glycol, polyvinyl pyrrolidone, povidone, pregelatinized starch, propyl paraben, retinyl palmitate, shellac, silicon dioxide, sodium carboxymethyl cellulose, sodium citrate, sodium starch glycolate, sorbitol, starch (corn), stearic acid, sucrose, talc, titanium dioxide, vitamin A, vitamin E, vitamin C, and xylitol.

[0902] *Pharmaceutically acceptable salts:* The present disclosure also includes pharmaceutically acceptable salts of the compounds described herein. As used herein, "pharmaceutically acceptable salts" refers to derivatives of the disclosed compounds wherein the parent compound is modified by converting an existing acid or base moiety to its salt form (e.g., by reacting the free base group with a suitable organic acid). Examples of pharmaceutically acceptable salts include, but are not limited to, mineral or organic acid salts of basic residues such as amines; alkali or organic salts of acidic residues such as carboxylic acids; and the like. Representative acid addition salts include acetate, acetic acid, adipate, alginate, ascorbate, aspartate, benzenesulfonate, benzene sulfonic acid, benzoate, bisulfate, borate, butyrate, camphorate, camphorsulfonate, citrate, cyclopentanepropionate, digluconate, dodecylsulfate, ethanesulfonate, fumarate, glucoheptonate, glycerophosphate, hemisulfate, heptonate, hexanoate, hydrobromide, hydrochloride, hydroiodide, 2-hydroxy-ethanesulfonate, lactobionate, lactate, laurate, lauryl sulfate, malate, maleate, malonate, methanesulfonate, 2-naphthalenesulfonate, nicotinate, nitrate, oleate, oxalate, palmitate, pamoate, pectinate, persulfate, 3-phenylpropionate, phosphate, picrate, pivalate, propionate, stearate, succinate, sulfate, tartrate, thiocyanate, toluenesulfonate, undecanoate, valerate salts, and the like. Representative alkali or alkaline earth metal salts include sodium, lithium, potassium, calcium, magnesium, and the like, as well as nontoxic ammonium, quaternary ammonium, and amine cations, including, but not limited to ammonium, tetramethylammonium, tetraethylammonium, methylamine, dimethylamine, trimethylamine, triethylamine, ethylamine, and the like. The pharmaceutically acceptable salts of the present disclosure include the conventional non-toxic salts of the parent compound formed, for example, from non-toxic inorganic or organic acids. The pharmaceutically acceptable salts of the present disclosure can be synthesized from the parent compound which contains a basic or acidic moiety by conventional chemical methods. Generally, such salts can be prepared by reacting the free acid or base forms of these compounds with a stoichiometric amount of the appropriate base or acid in wafer or in an organic solvent, or in a mixture of the two; generally, nonaqueous media like ether, ethyl acetate, ethanol, isopropanol, or acetonitrile are preferred. Lists of suitable salts are found in Remington's Pharmaceutical Sciences, 17th ed., Mack Publishing Company, Easton, Pa., 1985, p. 1418, Pharmaceutical Salts: Properties, Selection, and Use, P.H. Stahl and C.G. Wermuth (eds), Wiley-VCH, 2008, and Berge et al., Journal of Pharmaceutical Science, 66,1-19 (1977).

[0903] *Pharmaceutically acceptable solvate:* The term "pharmaceutically acceptable solvate," as used herein, means a compound of the disclosure wherein molecules of a suitable solvent are incorporated in the crystal lattice. A suitable solvent is physiologically tolerable at the dosage administered. For example, solvates may be prepared by crystallization, recrystallization, or precipitation from a solution that includes organic solvents, water, or a mixture thereof. Examples of suitable solvents are ethanol, wafer (for example, mono-, di-, and tri-hydrates), *N*-methylpyrrolidinone (NMP), dimethyl sulfoxide (DMSO), *N,N*'-dimethylformamide (DMF), *N,N*'-dimethylacetamide (DMAC), 1,3-dimethyl-2-imidazolidinone (DMEU), 1,3-dimethyl-3,4,5,6-tetrahydro-2-(1H)-pyrimidinone (DMPU), acetonitrile (ACN), propylene glycol, ethyl acetate, benzyl alcohol, 2-pyrrolidone, benzyl benzoate, and the like. When wafer is the solvent, the solvate is referred to as a "hydrate."

[0904] *Pharmacokinetic:* As used herein, "pharmacokinetic" refers to any one or more properties of a molecule or compound as it relates to the determination of the fate of substances administered to a living organism. Pharmacokinetics is divided into several areas including the extent and rate of absorption, distribution, metabolism and excretion. This is commonly referred to as ADME where: (A) Absorption is the process of a substance entering the blood circulation; (D) Distribution is the dispersion or dissemination of substances throughout the fluids and tissues of the body; (M) Metabolism (or Biotransformation) is the irreversible transformation of parent compounds into daughter metabolites; and (E) Excretion (or Elimination) refers to the elimination of the substances from the body In rare cases, some drugs irreversibly accumulate in body tissue.

[0905] *Physicochemical:* As used herein, "physicochemical" means of or relating to a physical and/or chemical property.

[0906] *Polypeptide per unit drug (PUD):* As used herein, a PUD or product per unit drug, is defined as a subdivided portion of total daily dose, usually 1 mg, pg, kg, etc., of a product (such as a polypeptide) as measured in body fluid or tissue, usually defined in concentration such as pmol/mL, mmol/mL, etc divided by the measure in the body fluid.

[0907] *Preventing:* As used herein, the term "preventing" refers to partially or completely delaying onset of an infection, disease, disorder and/or condition; partially or completely delaying onset of one or more symptoms, features, or clinical manifestations of a particular infection, disease, disorder, and/or condition; partially or completely delaying onset of one or more symptoms, features, or manifestations of a particular infection, disease, disorder, and/or condition; partially or

completely delaying progression from an infection, a particular disease, disorder and/or condition; and/or decreasing the risk of developing pathology associated with the infection, the disease, disorder, and/or condition.

[0908] *Prodrug:* The present disclosure also includes prodrugs of the compounds described herein. As used herein, "prodrugs" refer to any substance, molecule or entity which is in a form predicate for that substance, molecule or entity to act as a therapeutic upon chemical or physical alteration. Prodrugs may by covalently bonded or sequestered in some way and which release or are converted into the active drug moiety prior to, upon or after administered to a mammalian subject. Prodrugs can be prepared by modifying functional groups present in the compounds in such a way that the modifications are cleaved, either in routine manipulation or *in vivo,* to the parent compounds. Prodrugs include compounds wherein hydroxyl, amino, sulfhydryl, or carboxyl groups are bonded to any group that, when administered to a mammalian subject, cleaves to form a free hydroxyl, amino, sulfhydryl, or carboxyl group respectively. Preparation and use of prodrugs is discussed in T. Higuchi and V. Stella, "Pro-drugs as Novel Delivery Systems," Vol. 14 of the A.C.S. Symposium Series, and in Bioreversible Carriers in Drug Design, ed. Edward B. Roche, American Pharmaceutical Association and Pergamon Press, 1987.

[0909] *Proliferate:* As used herein, the term "proliferate" means to grow, expand or increase or cause to grow, expand or increase rapidly. "Proliferative" means having the ability to proliferate. "Anti-proliferative" means having properties counter to or inapposite to proliferative properties.

[0910] *Prophylactic:* As used herein, "prophylactic" refers to a therapeutic or course of action used to prevent the spread of disease.

[0911] *Prophylaxis:* As used herein, a "prophylaxis" refers to a measure taken to maintain health and prevent the spread of disease. An "immune phrophylaxis" refers to a measure to produce active or passive immunity to prevent the spread of disease.

[0912] *Protein cleavage site:* As used herein, "protein cleavage site" refers to a site where controlled cleavage of the amino acid chain can be accomplished by chemical, enzymatic or photochemical means.

[0913] *Protein cleavage signal:* As used herein "protein cleavage signal" refers to at least one amino acid that flags or marks a polypeptide for cleavage.

[0914] *Protein of interest:* As used herein, the terms "proteins of interest" or "desired proteins" include those provided herein and fragments, mutants, variants, and alterations thereof.

[0915] *Proximal:* As used herein, the term "proximal" means situated nearer to the center or to a point or region of interest.

[0916] *Pseudouridine:* As used herein, pseudouridine refers to the C-glycoside isomer of the nucleoside uridine. A "pseudouridine analog" is any modification, variant, isoform or derivative of pseudouridine. For example, pseudouridine analogs include but are not limited to 1-carboxymethyl-pseudouridine, 1-propynyl-pseudouridine, 1-taurinomethyl-pseudouridine, 1-taurinomethyl-4-thio-pseudouridine, 1-methylpseudouridine ($m^1\psi$), 1-methyl-4-thio-pseudouridine ($m^1s^4\psi$), 4-thio-1-methyl-pseudouridine, 3-methyl-pseudouridine ($m^3\psi$), 2-thio-1-methyl-pseudouridine, 1-methyl-1-deazapseudouridine, 2-thio-1-methyl-1-deaza-pseudouridine, dihydropseudouridine, 2-thio-dihydropseudouridine, 2-methoxyuridine, 2-methoxy-4-thio-uridine, 4-methoxy-pseudouridine, 4-methoxy-2-thio-pseudouridine, N 1-methyl-pseudouridine, 1-methyl-3-(3-amino-3-carboxypropyl)pseudouridine ($acp^3\ \psi$), and 2'-O-methyl-pseudouridine (ψm).

[0917] *Purified:* As used herein, "purify," "purified," "purification" means to make substantially pure or clear from unwanted components, material defilement, admixture or imperfection.

[0918] *Repeated transfection:* As used herein, the term "repeated transfection" refers to transfection of the same cell culture with a polynucleotide a plurality of times. The cell culture can be transfected at least twice, at least 3 times, at least 4 times, at least 5 times, at least 6 times, at least 7 times, at least 8 times, at least 9 times, at least 10 times, at least 11 times, at least 12 times, at least 13 times, at least 14 times, at least 15 times, at least 16 times, at least 17 times at least 18 times, at least 19 times, at least 20 times, at least 25 times, at least 30 times, at least 35 times, at least 40 times, at least 45 times, at least 50 times or more.

[0919] *Sample:* As used herein, the term "sample" or "biological sample" refers to a subset of its tissues, cells or component parts (e.g. body fluids, including but not limited to blood, mucus, lymphatic fluid, synovial fluid, cerebrospinal fluid, saliva, amniotic fluid, amniotic cord blood, urine, vaginal fluid and semen). A sample further may include a homogenate, lysate or extract prepared from a whole organism or a subset of its tissues, cells or component parts, or a fraction or portion thereof, including but not limited to, for example, plasma, serum, spinal fluid, lymph fluid, the external sections of the skin, respiratory, intestinal, and genitourinary tracts, fears, saliva, milk, blood cells, tumors, organs. A sample further refers to a medium, such as a nutrient broth or gel, which may contain cellular components, such as proteins or nucleic acid molecule.

[0920] *Signal Sequences:* As used herein, the phrase "signal sequences" refers to a sequence which can direct the transport or localization of a protein.

[0921] *Single unit dose:* As used herein, a "single unit dose" is a dose of any therapeutic administed in one dose/at one time/single route/single point of contact, i.e., single administration event.

[0922] *Similarity*: As used herein, the term "similarity" refers to the overall relatedness between polymeric molecules,

e.g. between polynucleotide molecules (e.g. DNA molecules and/or RNA molecules) and/or between polypeptide molecules. Calculation of percent similarity of polymeric molecules to one another can be performed in the same manner as a calculation of percent identity, except that calculation of percent similarity takes into account conservative substitutions as is understood in the art.

[0923] *Split dose:* As used herein, a "split dose" is the division of single unit dose or total daily dose into two or more doses.

[0924] *Stable:* As used herein "stable" refers to a compound that is sufficiently robust to survive isolation to a useful degree of purity from a reaction mixture, and preferably capable of formulation into an efficacious therapeutic agent.

[0925] *Stabilized:* As used herein, the term "stabilize", "stabilized," "stabilized region" means to make or become stable.

[0926] *Stereoisomer:* As used herein, the term "stereoisomer" refers to all possible different isomeric as well as conformational forms which a compound may possess (e.g., a compound of any formula described herein), in particular all possible stereochemically and conformationally isomeric forms, all diastereomers, enantiomers and/or conformers of the basic molecular structure. Some compounds of the present disclosure may exist in different tautomeric forms, all of the latter being included within the scope of the present disclosure.

[0927] *Subject:* As used herein, the term "subject" or "patient" refers to any organism to which a composition in accordance with the disclosure may be administered, *e.g.,* for experimental, diagnostic, prophylactic, and/or therapeutic purposes. Typical subjects include animals (*e.g.,* mammals such as mice, rats, rabbits, non-human primates, and humans) and/or plants.

[0928] *Substantially:* As used herein, the term "substantially" refers to the qualitative condition of exhibiting total or near-total extent or degree of a characteristic or property of interest. One of ordinary skill in the biological arts will understand that biological and chemical phenomena rarely, if ever, go to completion and/or proceed to completeness or achieve or avoid an absolute result. The term "substantially" is therefore used herein to capture the potential lack of completeness inherent in many biological and chemical phenomena.

[0929] *Substantially equal:* As used herein as it relates to time differences between doses, the term means plus/minus 2%.

[0930] *Substantially simultaneously:* As used herein and as it relates to plurality of doses, the term means within 2 seconds.

[0931] *Suffering from:* An individual who is "suffering from" a disease, disorder, and/or condition has been diagnosed with or displays one or more symptoms of a disease, disorder, and/or condition.

[0932] *Susceptible to:* An individual who is "susceptible to" a disease, disorder, and/or condition has not been diagnosed with and/or may not exhibit symptoms of the disease, disorder, and/or condition but harbors a propensity to develop a disease or its symptoms. In some embodiments, an individual who is susceptible to a disease, disorder, and/or condition (for example, cancer) may be characterized by one or more of the following: (1) a genetic mutation associated with development of the disease, disorder, and/or condition; (2) a genetic polymorphism associated with development of the disease, disorder, and/or condition; (3) increased and/or decreased expression and/or activity of a protein and/or nucleic acid associated with the disease, disorder, and/or condition; (4) habits and/or lifestyles associated with development of the disease, disorder, and/or condition; (5) a family history of the disease, disorder, and/or condition; and (6) exposure to and/or infection with a microbe associated with development of the disease, disorder, and/or condition. In some embodiments, an individual who is susceptible to a disease, disorder, and/or condition will develop the disease, disorder, and/or condition. In some embodiments, an individual who is susceptible to a disease, disorder, and/or condition will not develop the disease, disorder, and/or condition.

[0933] *Sustained release:* As used herein, the term "sustained release" refers to a pharmaceutical composition or compound release profile that conforms to a release rate over a specific period of time.

[0934] *Synthetic:* The term "synthetic" means produced, prepared, and/or manufactured by the hand of man. Synthesis of polynucleotides or polypeptides or other molecules of the present disclosure may be chemical or enzymatic.

[0935] *Vaccine:* As used herein, a vaccine is a compound or composition which comprises at least one polynucleotide encoding at least one antigen.

[0936] *Targeted Cells:* As used herein, "targeted cells" refers to any one or more cells of interest. The cells may be found *in vitro, in vivo, in* situ or in the tissue or organ of an organism. The organism may be an animal, preferably a mammal, more preferably a human and most preferably a patient.

[0937] *Therapeutic Agent:* The term "therapeutic agent" refers to any agent that, when administered to a subject, has a therapeutic, diagnostic, and/or prophylactic effect and/or elicits a desired biological and/or pharmacological effect.

[0938] *Therapeutically effective amount:* As used herein, the term "therapeutically effective amount" means an amount of an agent to be delivered (*e.g.,* nucleic acid, drug, therapeutic agent, diagnostic agent, prophylactic agent, *etc.*) that is sufficient, when administered to a subject suffering from or susceptible to an infection, disease, disorder, and/or condition, to treat, improve symptoms of, diagnose, prevent, and/or delay the onset of the infection, disease, disorder, and/or condition.

[0939] *Therapeutically effective outcome:* As used herein, the term "therapeutically effective outcome" means an

outcome that is sufficient in a subject suffering from or susceptible to an infection, disease, disorder, and/or condition, to treat, improve symptoms of, diagnose, prevent, and/or delay the onset of the infection, disease, disorder, and/or condition.

[0940] *Total daily dose:* As used herein, a "total daily dose" is an amount given or prescribed in 24 hr period. It may be administered as a single unit dose.

[0941] *Transcription* factor As used herein, the term "transcription factor" refers to a DNA-binding protein that regulates transcription of DNA into RNA, for example, by activation or repression of transcription. Some transcription factors effect regulation of transcription alone, while others act in concert with other proteins. Some transcription factor can both activate and repress transcription under certain conditions. In general, transcription factors bind a specific target sequence or sequences highly similar to a specific consensus sequence in a regulatory region of a target gene. Transcription factors may regulate transcription of a target gene alone or in a complex with other molecules.

[0942] *Transfection:* As used herein, the term "transfection" refers to methods to introduce exogenous nucleic acids into a cell. Methods of transfection include, but are not limited to, chemical methods, plysical treatments and cationic lipids or mixtures.

[0943] *Translation:* As used herein "translation" is the process by which a polynucleotide molecule is processed by a ribosome or ribosomal-like machinery, e.g., cellular or artificial, to produce a peptide or polypeptide.

[0944] *Transcription:* As used herein "transcription" is the process by which a polynucleotide molecule is processed by a polymerase or other enzyme to produce a polynucleotide, e.g., an RNA polynucleotide.

[0945] *Treating:* As used herein, the term "treating" refers to partially or completely alleviating, ameliorating, improving, relieving, delaying onset of, inhibiting progression of, reducing severity of, and/or reducing incidence of one or more symptoms or features of a particular infection, disease, disorder, and/or condition. Treatment may be administered to a subject who does not exhibit signs of a disease, infection, disorder, and/or condition and/or to a subject who exhibits only early signs of a disease, infection, disorder, and/or condition for the purpose of decreasing the risk of developing pathology associated with the disease, infection, disorder, and/or condition.

[0946] *Unmodified:* As used herein, "unmodified" refers to any substance, compound or molecule prior to being changed in any way. Unmodified may, but does not always, refer to the wild type or native form of a biomolecule. Molecules may undergo a series of modifications whereby each modified molecule may serve as the "unmodified" starting molecule for a subsequent modification.

[0947] *Vaccine:* As used herein, the phrase "vaccine" refers to a biological preparation that improves immunity in the context of a particular disease, disorder or condition.

[0948] *Viral protein:* As used herein, the pharse "viral protein" means any protein originating from a virus.

Equivalents and Scope

[0949] Those skilled in the art will recognize, or be able to ascertain using no more than routine experimentation, many equivalents to the specific embodiments in accordance with the invention described herein. The scope of the present invention is not intended to be limited to the above Description, but rather is as set forth in the appended claims.

[0950] In the claims, articles such as "a," "an," and "the" may mean one or more than one unless indicated to the contrary or otherwise evident from the context. Claims or descriptions that include "or" between one or more members of a group are considered satisfied if one, more than one, or all of the group members are present in, employed in, or otherwise relevant to a given product or process unless indicated to the contrary or otherwise evident from the context. The invention includes embodiments in which exactly one member of the group is present in, employed in, or otherwise relevant to a given product or process. The invention includes embodiments in which more than one, or all of the group members are present in, employed in, or otherwise relevant to a given product or process.

[0951] It is also noted that the term "comprising" is intended to be open and permits but does not require the inclusion of additional elements or steps. When the term "comprising" is used herein, the term "consisting of" is thus also encompassed and disclosed.

[0952] Where ranges are given, endpoints are included. Furthermore, it is to be understood that unless otherwise indicated or otherwise evident from the context and understanding of one of ordinary skill in the art, values that are expressed as ranges can assume any specific value or subrange within the stated ranges in different embodiments of the invention, to the tenth of the unit of the lower limit of the range, unless the context clearly dictates otherwise.

[0953] Section and table headings are not intended to be limiting.

EXAMPLES

Example 1. Manufacture of Polynucleotides

[0954] According to the present invention, the manufacture of polynucleotides and or parts or regions thereof may be

accomplished utilizing the methods taught in USSN 61/800,049 filed March 15, 2013 entitled "Manufacturing Methods for Production of RNA Transcripts".

[0955] Purification methods may include those taught in US Provisional Patent Application No. 61/799,872, US Provisional Patent Application No. 61/794,842, US Provisional Patent Application 61/800,326.

[0956] Detection and characterization methods of the polynucleotides may be performed as taught in US Provisional Patent Application No. 61/799,780 and US Provisional Patent Application No. 61/798,945.

[0957] Characterization of the polynucleotides of the invention may be accomplished using a procedure selected from the group consisting of polynucleotide mapping, reverse transcriptase sequencing, charge distribution analysis, and detection of RNA impurities, wherein characterizing comprises determining the RNA transcript sequence, determining the purity of the RNA transcript, or determining the charge heterogeneity of the RNA transcript. Such methods are taught in, for example, US Provisional Patent Application No. 61/799,905 and US Provisional Patent Application No. 61/800,110.

Example 2. Chimeric polynucleotide synthesis

Introduction

[0958] According to the present invention, two regions or parts of a chimeric polynucleotide may be joined or ligated using triphosphate chemistry.

[0959] According to this method, a first region or part of 100 nucleotides or less is chemically synthesized with a 5' monophosphate and terminal 3'desOH or blocked OH. If the region is longer than 80 nucleotides, it may be synthesized as two strands for ligation.

[0960] If the first region or part is synthesized as a non-positionally modified region or part using in vitro transcription (IVT), conversion the 5'monophosphate with subsequent capping of the 3' terminus may follow.

[0961] Monophosphate protecting groups may be selected from any of those known in the art.

[0962] The second region or part of the chimeric polynucleotide may be synthesized using either chemical synthesis or IVT methods. IVT methods may include an RNA polymerase that can utilize a primer with a modified cap. Alternatively, a cap of up to 130 nucleotides may be chemically synthesized and coupled to the IVT region or part.

[0963] It is noted that for ligation methods, ligation with DNA T4 ligase, followed by treatment with DNAse should readily avoid concatenation.

[0964] The entire chimeric polynucleotide need not be manufactured with a phosphate-sugar backbone. If one of the regions or parts encodes a polypeptide, then it is preferable that such region or part comprise a phosphate-sugar backbone.

[0965] Ligation is then performed using any known click chemistry, orthoclick chemistry, solulink, or other bioconjugate chemistries known to those in the art.

Synthetic route

[0966] The chimeric polynucleotide is made using a series of starting segments Such segments include:

(a) Capped and protected 5' segment comprising a normal 3'OH (SEG. 1)
(b) 5' triphosphate segment which may include the coding region of a polypeptide and comprising a normal 3'OH (SEG. 2)
(c) 5' monophosphate segment for the 3' end of the chimeric polynucleotide (e.g., the tail) comprising cordycepin or no 3'OH (SEG. 3)

[0967] After synthesis (chemical or IVT), segment 3 (SEG. 3) is treated with cordycepin and then with pyrophosphatase to create the 5'monophosphate.

[0968] Segment 2 (SEG. 2) is then ligated to SEG. 3 using RNA ligase. The ligated polynucleotide is then purified and treated with pyrophosphatase to cleave the diphosphate. The treated SEG.2-SEG. 3 construct is then purified and SEG. 1 is ligated to the 5' terminus. A further purification step of the chimeric polynucleotide may be performed.

[0969] Where the chimeric polynucleotide encodes a polypeptide, the ligated or joined segments may be represented as: 5'UTR (SEG. 1), open reading frame or ORF (SEG. 2) and 3'UTR+PolyA (SEG. 3).

[0970] The yields of each step may be as much as 90-95%.

Example 3: PCR for cDNA Production

[0971] PCR procedures for the preparation of cDNA are performed using 2x KAPA HIFI™ HotStart ReadyMix by Kapa Biosystems (Woburn, MA). This system includes 2x KAPA ReadyMix12.5 μl; Forward Primer (10 uM) 0.75 μl; Reverse

Primer (10 uM) 0.75 μl; Template cDNA -100 ng; and dH_20 diluted to 25.0 μl. The reaction conditions are at 95° C for 5 min. and 25 cycles of 98° C for 20 sec, then 58° C for 15 sec, then 72° C for 45 sec, then 72° C for 5 min. then 4° C to termination.

[0972] The reverse primer of the instant disclosure incorporates a poly-T_{120} (SEQ ID NO: 2283) for a poly-A_{120} (SEQ ID NO: 2282) in the mRNA. Other reverse primers with longer or shorter poly(T) tracts can be used to adjust the length of the poly(A) tail in the polynucleotide mRNA.

[0973] The reaction is cleaned up using Invitrogen's PURELINK™ PCR Micro Kit (Carlsbad, CA) per manufacturer's instructions (up to 5 μg). Larger reactions will require a cleanup using a product with a larger capacity. Following the cleanup, the cDNA is quantified using the NANODROP™ and analyzed by agarose gel electrophoresis to confirm the cDNA is the expected size. The cDNA is then submitted for sequencing analysis before proceeding to the *in vitro* transcription reaction.

Example 4. ***In vitro* Transcription (IVT)**

[0974] The *in vitro* transcription reaction generates polynucletodies containing uniformly modified polynucleotides. Such uniformly modified polynucleotides may comprise a region or part of the polynucleotides of the invention. The input nucleotide triphosphate (NTP) mix is made in-house using natural and un-natural NTPs.

[0975] A typical *in vitro* transcription reaction includes the following:

1	Template cDNA	1.0 μg	
2	10x transcription buffer (400 mM Tris-HCl pH 8.0, 190 mM $MgCl_2$, 50 mM DTT, 10 mM Spermidine)		2.0 μl
3	Custom NTPs (25mM each)	7.2 μl	
4	RNase Inhibitor	20 U	
5	T7 RNA polymerase	3000 U	
6	dH_20		Up to 20.0 μl. and
7	Incubation at 37° C for 3 hr-5 hrs.		

[0976] The crude IVT mix may be stored at 4° C overnight for cleanup the next day. 1 U of RNase-free DNase is then used to digest the original template. After 15 minutes of incubation at 37° C, the mRNA is purified using Ambion's MEGACLEAR™ Kit (Austin, TX) following the manufacturer's instructions. This kit can purify up to 500 pg of RNA. Following the cleanup, the RNA is quantified using the NanoDrop and analyzed by agarose gel electrophoresis to confirm the RNA is the proper size and that no degradation of the RNA has occurred.

Example 5. Enzymatic Capping

[0977] Capping of a polynucleotide is performed as follows where the mixture includes: IVT RNA 60 μg-180μg and dH_20 up to 72 μl. The mixture is incubated at 65° C for 5 minutes to denature RNA, and then is transferred immediately to ice.

[0978] The protocol then involves the mixing of 10x Capping Buffer (0.5 M Tris-HCl (pH 8.0), 60 mM KCl, 12.5 mM $MgCl_2$) (10.0 μl); 20 mM GTP (5.0 μl); 20 mM S-Adenosyl Methionine (2.5 μl); RNase Inhibitor (100 U); 2'-O-Methyl-transferase (400U); Vaccinia capping enzyme (Guanylyl transferase) (40 U); dH_20 (Up to 28 μl); and incubation at 37° C for 30 minutes for 60 pg RNA or up to 2 hours for 180 pg of RNA.

[0979] The polynucleotide is then purified using Ambion's MEGACLEAR™ Kit (Austin, TX) following the manufacturer's instructions. Following the cleanup, the RNA is quantified using the NANODROP™ (ThermoFisher, Waltham, MA) and analyzed by agarose gel electrophoresis to confirm the RNA is the proper size and that no degradation of the RNA has occurred. The RNA product may also be sequenced by running a reverse-transcription-PCR to generate the cDNA for sequencing.

Example 6. PolyA Tailing Reaction

[0980] Without a poly-T in the cDNA, a poly-A failing reaction must be performed before cleaning the final product. This is done by mixing Capped IVT RNA (100 μl); RNase Inhibitor (20 U); 10x Tailing Buffer (0.5 M Tris-HCl (pH 8.0), 2.5 M NaCl, 100 mM $MgCl_2$)(12.0 μl), 20 mM ATP (6.0 μl); Poly-A Polymerase (20 U); dH_20 up to 123.5 μl and incubation at 37° C for 30 min. If the poly-A tail is already in the transcript, then the failing reaction may be skipped and proceed

directly to cleanup with Ambion's MEGACLEAR™ kit (Austin, TX) (up to 500 μg). Poly-A Polymerase is preferably a recombinant enzyme expressed in yeast.

[0981] It should be understood that the processivity or integrity of the polyA failing reaction may not always result in an exact size polyA fail. Hence polyA tails of approximately between 40-200 nucleotides (SEQ ID NO: 2284), e.g, about 40, 50, 60, 70, 80, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, 100, 101, 102, 103, 104, 105, 106, 107, 108, 109, 110, 150-165, 155, 156, 157, 158, 159, 160, 161, 162, 163, 164 or 165 are within the scope of the invention.

Example 7. Natural 5' Caps and 5' Cap Analogues

[0982] 5'-capping of polynucleotides may be completed concomitantly during the in vitro-transcription reaction using the following chemical RNA cap analogs to generate the 5'-guanosine cap structure according to manufacturer protocols: 3'-O-Me-m7G(5')ppp(5') G [the ARCA cap];G(5')ppp(5')A; G(5')ppp(5')G; m7G(5')ppp(5')A; m7G(5')ppp(5')G (New England BioLabs, Ipswich, MA). 5'-capping of modified RNA may be completed post-transcriptionally using a Vaccinia Virus Capping Enzyme to generate the "Cap 0" structure: m7G(5')ppp(5')G (New England BioLabs, Ipswich, MA). Cap 1 structure may be generated using both Vaccinia Virus Capping Enzyme and a 2'-O methyl-transferase to generate: m7G(5')ppp(5')G-2'-O-methyl. Cap 2 structure may be generated from the Cap 1 structure followed by the 2'-O-methylation of the 5'-antepenultimate nucleotide using a 2'-O methyl-transferase. Cap 3 structure may be generated from the Cap 2 structure followed by the 2'-O-methylation of the 5'-preantepenultimate nucleotide using a 2'-O methyl-transferase. Enzymes are preferably derived from a recombinant source.

[0983] When transfected into mammalian cells, the modified mRNAs have a stability of between 12-18 hours or more than 18 hours, e.g., 24, 36, 48, 60, 72 or greater than 72 hours.

Example 8. Capping Assays

A. Protein Expression Assay

[0984] Polynucleotides encoding a polypeptide, containing any of the caps taught herein can be transfected into cells at equal concentrations. 6, 12, 24 and 36 hours post-transfection the amount of protein secreted into the culture medium can be assayed by ELISA. Synthetic polynucleotides that secrete higher levels of protein into the medium would correspond to a synthetic polynucleotide with a higher translationally-competent Cap structure.

B. Purity Analysis Synthesis

[0985] Polynucleotides encoding a polypeptide, containing any of the caps taught herein can be compared for purity using denaturing Agarose-Urea gel electrophoresis or HPLC analysis. Polynucleotides with a single, consolidated band by electrophoresis correspond to the higher purity product compared to polynucleotides with multiple bands or streaking bands. Synthetic polynucleotides with a single HPLC peak would also correspond to a higher purity product. The capping reaction with a higher efficiency would provide a more pure polynucleotide population.

C. Cytokine Analysis

[0986] Polynucleotides encoding a polypeptide, containing any of the caps taught herein can be transfected into cells at multiple concentrations. 6, 12, 24 and 36 hours post-transfection the amount of pro-inflammatory cytokines such as TNF-alpha and IFN-beta secreted into the culture medium can be assayed by ELISA. Polynucleotides resulting in the secretion of higher levels of pro-inflammatory cytokines into the medium would correspond to a polynucleotides containing an immune-activating cap structure.

D. Capping Reaction Efficiency

[0987] Polynucleotides encoding a polypeptide, containing any of the caps taught herein can be analyzed for capping reaction efficiency by LC-MS after nuclease treatment. Nuclease treatment of capped polynucleotides would yield a mixture of free nucleotides and the capped 5'-5-triphosphate cap structure detectable by LC-MS. The amount of capped product on the LC-MS spectra can be expressed as a percent of total polynucleotide from the reaction and would correspond to capping reaction efficiency. The cap structure with higher capping reaction efficiency would have a higher amount of capped product by LC-MS.

Example 9. Agarose Gel Electrophoresis of Modified RNA or RT PCR Products

[0988] Individual polynucleotides (200-400 ng in a 20 pl volume) or reverse transcribed PCR products (200-400 ng) are loaded into a well on a non-denaturing 1.2% Agarose E-Gel (Invitrogen, Carlsbad, CA) and run for 12-15 minutes according to the manufacturer protocol.

Example 10. Nanodrop Modified RNA Quantification and UV Spectral Data

[0989] Modified polynucleotides in TE buffer (1 pl) are used for Nanodrop UV absorbance readings to quantitate the yield of each polynucleotide from an chemical synthesis or *in vitro* transcription reaction.

Example 11. Formulation of Modified mRNA Using Lipidoids

[0990] Polynucleotides are formulated for *in vitro* experiments by mixing the polynucleotides with the lipidoid at a set ratio prior to addition to cells. *In vivo* formulation may require the addition of extra ingredients to facilitate circulation throughout the body To test the ability of these lipidoids to form particles suitable for *in vivo* work, a standard formulation process used for siRNA-lipidoid formulations may used as a starting point. After formation of the particle, polynucleotide is added and allowed to integrate with the complex. The encapsulation efficiency is determined using a standard dye exclusion assays.

Example 12. Method of Screening for Protein Expression

A. Electrospray Ionization

[0991] A biological sample which may contain proteins encoded by a polynucleotide administered to the subject is prepared and analyzed according to the manufacturer protocol for electrospray ionization (ESI) using 1, 2, 3 or 4 mass analyzers. A biologic sample may also be analyzed using a tandem ESI mass spectrometry system.

[0992] Patterns of protein fragments, or whole proteins, are compared to known controls for a given protein and identity is determined by comparison.

B. Matrix-Assisted Laser Desorption/Ionization

[0993] A biological sample which may contain proteins encoded by one or more polynucleotides administered to the subject is prepared and analyzed according to the manufacturer protocol for matrix-assisted laser desorption/ionization (MALDI).

[0994] Patterns of protein fragments, or whole proteins, are compared to known controls for a given protein and identity is determined by comparison.

C. Liquid Chromatography-Mass spectrometry-Mass spectrometry

[0995] A biological sample, which may contain proteins encoded by one or more polynucleotides, may be treated with a trypsin enzyme to digest the proteins contained within. The resulting peptides are analyzed by liquid chromatography-mass spectrometry-mass spectrometry (LC/MS/MS). The peptides are fragmented in the mass spectrometer to yield diagnostic patterns that can be matched to protein sequence databases via computer algorithms. The digested sample may be diluted to achieve 1 ng or less starting material for a given protein. Biological samples containing a simple buffer background (e.g. water or volatile salts) are amenable to direct in-solution digest; more complex backgrounds (e.g. detergent, non-volatile salts, glycerol) require an additional clean-up step to facilitate the sample analysis.

[0996] Patterns of protein fragments, or whole proteins, are compared to known controls for a given protein and identity is determined by comparison.

Example 13. Cyclization and/or concatemerization

[0997] According to the present invention, a polynucleotide may be cyclized, or concatemerized, to generate a translation competent molecule to assist interactions between poly-A binding proteins and 5'-end binding proteins. The mechanism of cyclization or concatemerization may occur through at least 3 different routes: 1) chemical, 2) enzymatic, and 3) ribozyme catalyzed. The newly formed 5'-/3'-linkage may be intramolecular or intermolecular.

[0998] In the first route, the 5'-end and the 3'-end of the nucleic acid contain chemically reactive groups that, when close together, form a new covalent linkage between the 5'-end and the 3'-end of the molecule. The 5'-end may contain

an NHS-ester reactive group and the 3'-end may contain a 3'-amino-terminated nucleotide such that in an organic solvent the 3'-amino-terminated nucleotide on the 3'-end of a synthetic mRNA molecule will undergo a nucleophilic attack on the 5'-NHS-ester moiety forming a new 5'-/3'-amide bond.

[0999] In the second route, T4 RNA ligase may be used to enzymatically link a 5'-phosphorylated nucleic acid molecule to the 3'-hydroxyl group of a nucleic acid forming a new phosphorodiester linkage. In an example reaction, 1μg of a nucleic acid molecule is incubated at 37°C for 1 hour with 1-10 units of T4 RNA ligase (New England Biolabs, Ipswich, MA) according to the manufacturer's protocol. The ligation reaction may occur in the presence of a split polynucleotide capable of base-pairing with both the 5'- and 3'- region in juxtaposition to assist the enzymatic ligation reaction.

[1000] In the third route, either the 5'-or 3'-end of the cDNA template encodes a ligase ribozyme sequence such that during *in vitro* transcription, the resultant nucleic acid molecule can contain an active ribozyme sequence capable of ligating the 5'-end of a nucleic acid molecule to the 3'-end of a nucleic acid molecule. The ligase ribozyme may be derived from the Group I Intron, Group I Intron, Hepatitis Delta Virus, Hairpin ribozyme or may be selected by SELEX (systematic evolution of ligands by exponential enrichment). The ribozyme ligase reaction may take 1 to 24 hours at temperatures between 0 and 37°C.

Example 14. Antigen Polynucleotides

[1001] Polynucleotides used in the studies hereing which encode certain infectious agent antigens orvariants thereof are shown in Table 28.

Table 28. Antigen polynucleotides (RNA vaccine)

Construct Number	Gene ID	Description	ORF SEQ ID NO	mRNA SEQ ID NO	Protein SEQ ID NO	Construct
1	120552	Flu PR8 HA ferritin	2042	2115	2188	HAPR8_Ferritin_Hs3

2	120553	flu 1999NC HA ferritin	2043	2116	2189	HA1999NC_Ferritin_Hs3
3	125387	H5N1 HA antigen	2044	2117	2190	H5N1_HA_NIBRG-14
4	125388	H1N1 PR8 HA antigen with MT 5U/3U	2045	2118	2191	CurevacPR8HA_MT5U_Hs3U
5	135929	Influenza HA antigen	2046	2119	2192	HA_A/Mallard/Netherlands/12/2000_H7N3_Hs3U
6	135930	Influenza HA antigen	2047	2120	2193	HA_A/Puerto Rico/8/1934_H1N1_Hs3U
7	135931	Influenza HA antigen	2048	2121	2194	HA_A/Viet Nam/850/2009_H1N1_Hs3U
8	135932	Influenza HA antigen	2049	2122	2195	HA_A/New Caledonia/20/99_H1N1_Hs3U
9	135933	Influenza HA antigen	2050	2123	2196	HA_A/Vietnam/1203/2004_H5N1_Hs3U
10	135934	Influenza HA antigen	2051	2124	2197	HA_A/Hong Kong/1073/99_H9N2_Hs3U
11	135935	Influenza HA antigen	2052	2125	2198	HA_A/Wisconsin/67/2005_H3N2_Hs3U
12	135936	Influenza HA antigen	2053	2126	2199	HA_A/Hong Kong/1/1968_H3N2_Hs3U
13	136473	flu HA antigen	2054	2127	2200	A/duck/Anhui/SC702/2013(H7N9)_Hs3U
14	136754	C diff antigen	2055	2128		Cdiff_toxinA.truc_nMyc.cFLAG_Hs3U
15	136755	C diff antigen	2056	2129		Cdiff_toxinB.truc_nMyc.cFLAG_Hs3U
16	136756	C diff antigen	2057	2130		Cdiff_cdtA.truc_nMyc.cFLAG_Hs3U
17	136757	C diff antigen	2058	2131		Cdiff_cdtB.truc_nMyc.cFLAG_Hs3U
18	120554	Dengue strain 2 domain 3 ferritin	2059	2132		DEN2_DIII_Ferritin_Hs3
19	120555	Dengue strain 3 domain 3 ferritin	2060	2133		DEN3_DIII_Ferritin_Hs3
20	120556	Dengue strain 4 domain 3 ferritin	2061	2134		DEN4_DIII_Ferritin_Hs3
21	120557	dengue strain 1-4 Domain 3	2062	2135		Dengue_TDIII_Hs3
22	121542	Dengue strain 1 domain 3 ferritin	2063	2136		DEN1_DIII_Ferritin_Corr_Hs3
23	131502	Dengue 2, D2Y98P strain, PrME transmembrane antigen	2064	2137	2201	DEN2_D2Y98P_PrME_Hs3
24	131503	Dengue 2, D2Y98P strain, PrME secreted antigen	2065	2138	2202	DEN2_D2Y98P_PrME80_Hs3
25	131507	Dengue 2, D2Y98P strain, PrME secreted antigen with dendritic targeting ScFv against mouse DEC205	2066	2139	2203	DEN2_D2Y98P_PrME80_ScFv.aDEC205.FLAG_Hs3
26	136358	Staph aureus antigen	2067	2140	2204	SpA_sec.D.wt.nXpress.cHA_VKsp_Hs3U

27	136359	Staph aureus antigen	2068	2141	2205	SpA_intra.D.kkaa.nFLAG.cMyc_Hs3U
28	136360	Staph aureus antigen	2069	2142	2206	SpA_mem.D.kkaa.nFLAG.cMyc _CD28mem_Hs3U
29	136361	Staph aureus antigen	2070	2143	2207	SpA_sec.D.kkaa.nFLAG.cMyc_VKsp_Hs3U
30	136362	Staph aureus antigen	2071	2144	2208	SpA_intra.D.wt.nXpress.cHA_Hs3U
31	136363	Staph aureus antigen	2072	2145	2209	SpA_mem.D.wt.nXpress.cHA_CD28mem_Hs3U
32	136364	Staph aureus antigen	2073	2146	2210	SpA_sec.D.wt_VKsp
33	136758	MRSA IsdA antigen	2074	2147	2211	MRSA_IsdA.truc_nMyc.cFLAG_Hs3U
34	136759	MRSA IsdB antigen	2075	2148	2212	MRSA_IsdB.truc_nMyc.cFLAG_Hs3U
35	136763	MRSA SDRD antigen	2076	2149	2213	MRSA_sdrd.truc_nMyc.cFLAG_Hs3U
36	136764	MRSA SDRE antigen	2077	2150	2214	MRSA_sdre.truc_nMyc.cFLAG_Hs3U
37	136765	MRSA MECA antigen	2078	2151	2215	MRSA_mecA.truc_nMyc.cFLAG_Hs3U
38	139325	MRSA SDRD antigen	2079	2152	2216	MRSA_sdrd.contig1._FLAG_Hs3U
39	139326	MRSA SDRD antigen	2080	2153	2217	MRSA_sdrd.Contig2._FLAG_Hs3U
40	139327	MRSA SDRE antigen	2081	2154	2218	MRSA_sdre.Contig1._FLAG_Hs3U
41	139328	MRSA SDRE antigen	2082	2155	2219	MRSA_sdre.Contig2._FLAG_Hs3U
42	140470	MRSA ISDA antigen	2083	2156	2220	MRSA_IsdA.fl_FLAG_Hs3U
43	139277	MRSA MECA antigen	2084	2157	2221	MRSA_mecA.fl._FLAG_Hs3U
44	136747	ETEC antigen	2085	2158		ETEC.eltA(S63K/R192G/L211A)_nMyc.cFLAG_Hs3U
45	136748	ETEC antigen	2086	2159		ETEC.eltB_nMyc.cFLAG_Hs3U
46	136749	ETEC antigen	2087	2160		ETEC.Sta3(A14Q)_nMyc.cFLAG_Hs3U
47	136750	ETEC antigen	2088	2161		ETEC.etpA._nMyc.cFLAG_Hs3U
48	136751	ETEC antigen	2089	2162		ETEC.etpB._nMyc.cFLAG_Hs3U
49	136752	ETEC antigen	2090	2163		ETEC.EatA_nMyc.cFLAG_Hs3U
50	136753	ETEC antigen	2091	2164		ETEC_cssA_nMyc.cFLAG_Hs3U
51	136747	ETEC antigen	2092	2165		ETEC.eltA(S63K/R192G/L211A)_nMyc.cFLAG_Hs3U
52	142544	ETEC antigen	2093	2166		ETEC.etpB.fl_FLAG_Hs3U
53	136749	ETEC antigen	2094	2167		ETEC.Sta3(A14Q)_nMyc.cFLAG_Hs3U
54	136753	ETEC CSSA antigen	2095	2168		ETEC_cssA_nMyc.cFLAG_Hs3U
55	139323	ETEC EATA antigen	2096	2169		ETEC_EatA.contig1._FLAG_Hs3U

56	139324	ETEC EATA antigen	2097	2170		ETEC_EatA.contig2._FLAG_Hs3U
57	139321	ETEC EPTA antigen	2098	2171		ETEC_eptA.contig1._FLAG_Hs3U
58	139322	ETEC ETPA antigen	2099	2172		ETEC_etpA.contig2._FLAG_Hs3U
59	136760	Tuberculosis antigen	2100	2173	2222	TB_Ag85A_nMyc.cFLAG_Hs3U
60	136761	Tuberculosis antigen	2101	2174	2223	TB_Ag85B_nMyc.cFLAG_Hs3U
61	136762	Tuberculosis antigen	2102	2175	2224	TB_TB10.4_nMyc.cFLAG_Hs3U
62	136335	Cholera toxin B	2103	2176		CholeraToxB.H78A.FLAG_VKsp_Hs3U
63	144097	vaccine Influenza HA antigen (nanoscaffold)	2104	2177	2225	PR8HA.Stalk.pscaffold_Hs3U
64	144094	vaccine Influenza HA antigen (splitcore)	2105	2178	2226	PR8HA.stalk.HbCore.monoRNA_Hs3U
65	144095	vaccine Influenza HA antigen (splitcore)	2106	2179	2227	PR8HA.stalk.HbCoreC.RKR_Hs3U
66	144096	vaccine Influenza HA antigen (splitcore)	2107	2180	2228	PR8HA.stalk.HbCoreN.RKR_Hs3U
67	144098	vaccine Influenza HA antigen; aglycosylated	2108	2181	2229	PR8HA.headless.Stalk.ugly(N76D/N270D)_Hs3U
68	144099	vaccine Influenza HA antigen; aglycosylated	2109	2182	2230	PR8HA.headless.Stalk.ugly(N27D/N28D/N40D/N76D/N270D)_Hs3U
69	150088	Vaccine MERS CoV spike protein (V5 epitope tagged)	2110	2183	2231	MERS-CoV_ProteinS/EMC/2012.V5_Hs3U
70	139276	vaccine MRSA IsdB antigen	2111	2184	2232	MRSA_IsdB.fl._FLAG_Hs3U
71	149385	MERS-CoV passive immunity	2112	2185		DPPIV.Fc_HS3U
72	150092	Vaccine HEV71 (polycistronic antigen containing VP1 VP2 & VP3) each separated by a furin cleavage site RKR	2113	2186	2233	HEV71.VP1.2.3.VP5_HS3U
73	144093	vaccine Influenza HA antigen	2114	2187	2234	PR8HA.Stalk_Hs3U
74		Vaccine Klebsiella pneumoniae: cefotaxime, ceftazidime and other broad spectrum cephalosporins.			2235	Beta-lactamase SHV-2
75		Vaccine PsaA SP Streptococcus pneumoniae surface adhesin A Fragment (22–309)			2236	Pneumococcal surface adhesin A
76		Vaccine Pseudomonas aeruginosa ampC beta-lactamase			2237	Pseudomonas type 2 Cephalosporinase

77		Vaccine Pseudomonas aeruginosa VIM-2, a carbapenem-hydrolyzing metallo-beta-lactamase			2238	Beta-lactamase class B VIM-2
78		Vaccine Pseudomonas aeruginosa, LCR-1 hydrolyzes methicillin.			2239	Beta-lactamase LCR-1
79		Vaccine Staph aureus Toxic shock syndrome toxin-1			2240	Staph aureus TSST1
80		Vaccine Staph aureus PVL toxin			2241	Staph aureus PVL toxin
81		vaccine Streptococcus pneumoniae Pneumolysin toxoid			2242	Pneumolysin toxoid
82		Full length with mutations of W433F, D385N, and C428G			2243	Streptococcus pneumoniae ppaC

[1002] Additional antigens including wild type and engineered antigens are taught in Tables 29 and 30.

Pan-flu NAVs

[1003] In one embodiment the "HA head" region of one or more influenza virus strains is removed leaving only the stem or transmembrane region. This region or multiple regions if selected, is then used as the immunogen to screen for optimal response to a viral challenge. As such broad neutralization could be achieved against multiple strains or a multi-response to one strain. The resultant vaccine would represent a pan-influenza vaccine.

[1004] Further, a pan-influenza vaccine could also be combined with any immune potentiator disclosed herein.

[1005] Polynucleotides used in the studies herein which encode certain infectious agent antigens orvariants thereof may be formulated in any of the formulations described herein including LNPs and may be administered intradermally (ID) or intramuscularly (IM) or by any suitable route.

[1006] Disclosed in Table 29 are the HA regions, transmembrane and cytoplasmic regions of several influenza strains to be used in the stem vaccination generation protocol just described.

Table 29. Wild type antigens encoded by NAV polynucleotides

Virus	Strain	Full length HA AA Sequence	HA1	HA2	TM	CY
H1N1	A/PR8/34	(SEQ ID NO: 2244) MKANLLVLLCALAAADADTICIGY HANNSTDTVDTVLEKNVTVTHSV NLLEDSHNGKLCRLKGIAPLQLG KCNIAGWLLGNPECDPLLPVRS WSYIVETPNSENGICYPGDFIDY EELREQLSSVSSFERFEIFPKES SWPNHNTNGVTAACSHEGKSSF YRNLLWLTEKEGSYPKLKNSYV NKKGKEVLVLWGIHHPPNSKEQ QNLYQNENAYVSVVTSNYNRRF TPEIAERPKVRDQAGRMNYYWT LLKPGDTIIFEANGNLIAPMYAFA LSRGFGSGIITSNASMHECNTKC QTPLGAINSSLPYQNIHPVTIGEC PKYVRSAKLRMVTGLRNNPSIQS RGLFGAIAGFIEGGWTGMIDGW YGYHHQNEQGSGYAADQKSTQ NAINGITNKVNTVIEKMNIQFTAV GKEFNKLEKRMENLNKKVDDGF LDIWTYNAELLVLLENERTLDFH DSNVKNLYEKVKSQLKNNAKEIG NGCFEFYHKCDNECMESVRNGT YDYPKYSEESKLNREKVDGVKL ESMGIYQILAIYSTVASSLVLLVSL GAISFWMCSNGSLQCRICI	(SEQ ID NO: 2245) MKANLLVLLCALAA ADADTICIGYHANNS TDTVDTVLEKNVTV THSVNLLEDSHNGK LCRLKGIAPLQLGK CNIAGWLLGNPECD PLLPVRSWSYIVET PNSENGICYPGDFI DYEELREQLSSVSS FERFEIFPKESSWP NHNTNGVTAACSH EGKSSFYRNLLWLT EKEGSYPKLKNSYV NKKGKEVLVLWGIH HPPNSKEQQNLYQ NENAYVSVVTSNYN RRFTPEIAERPKVR DQAGRMNYYWTLL KPGDTIIFEANGNLI APMYAFALSRGFGS GIITSNASMHECNTK CQTPLGAINSSLPY QNIHPVTIGECPKYV RSAKLRMVTGLRNN PSIQSR	(SEQ ID NO: 2246) GLFGAIAGFIEG GWTGMIDGWYG YHHQNEQGSGY AADQKSTQNAIN GITNKVNTVIEKM NIQFTAVGKEFN KLEKRMENLNKK VDDGFLDIWTYN AELLVLLENERTL DFHDSNVKNLYE KVKSQLKNNAKE IGNGCFEFYHKC DNECMESVRNG TYDYPKYSEESK LNREKVDGVKLE SMGIYQ	(SEQ ID NO: 2247) ILAIYS TVASS LVLLV SLGAI	(SEQ ID NO: 2248) SFWMC SNGSLQ CRICI

H1N1	A/New Caledonia/20/1999	(SEQ ID NO: 2249) MKAKLLVLLCTFTATYADTICIGYHANNSTDTVDTVLEKNVTVTHSVNLLEDSHNGKLCLLKGIAPLQLGNCSVAGWILGNPECELLISKESWSYIVETPNPENGTCYPGYFADYEELREQLSSVSSFERFEIFPKESSWPNHTVTGVSASCSHNGKSSFYRNLLWLTGKNGLYPNLSKSYVNNKEKEVLVLWGVHHPPNIGNQRALYHTENAYVSVVSSHYSRRFTPEIAKRPKVRDQEGRINYYWTLLEPGDTIIFEANGNLIAPWYAFALSRGFGSGIITSNAPMDECDAKCQTPQGAINSSLPFQNVHPVTIGECPKYVRSAKLRMVTGLRNIPSIQSRGLFGAIAGFIEGGWTGMVDGWYGYHHQNEQGSGYAADQKSTQNAINGITNKVNSVIEKMNTQFTAVGKEFNKLERRMENLNKKVDDGFLDIWTYNAELLVLLENERTLDFHDSNVKNLYEKVKSQLKNNAKEIGNGCFEFYHKCNNECMESVKNGTYDYPKYSEESKLNREKIDGVKLESMGVYQILAIYSTVASSLVLLVSLGAISFWMCSNGSLQCRICI	(SEQ ID NO: 2250) MKAKLLVLLCTFTATYADTICIGYHANNSTDTVDTVLEKNVTVTHSVNLLEDSHNGKLCLLKGIAPLQLGNCSVAGWILGNPECELLISKESWSYIVETPNPENGTCYPGYFADYEELREQLSSVSSFERFEIFPKESSWPNHTVTGVSASCSHNGKSSFYRNLLWLTGKNGLYPNLSKSYVNNKEKEVLVLWGVHHPPNIGNQRALYHTENAYVSVVSSHYSRRFTPEIAKRPKVRDQEGRINYYWTLLEPGDTIIFEANGNLIAPWYAFALSRGFGSGIITSNAPMDECDAKCQTPQGAINSSLPFQNVHPVTIGECPKYVRSAKLRMVTGLRNIPSIQSR	(SEQ ID NO: 2251) GLFGAIAGFIEGGWTGMVDGWYGYHHQNEQGSGYAADQKSTQNAINGITNKVNSVIEKMNTQFTAVGKEFNKLERRMENLNKKVDDGFLDIWTYNAELLVLLENERTLDFHDSNVKNLYEKVKSQLKNNAKEIGNGCFEFYHKCNNECMESVKNGTYDYPKYSEESKLNREKIDGVKLESMGVYQ	(SEQ ID NO: 2252) ILAIYSTVASSLVLLVSLGAI	(SEQ ID NO: 2253) SFWMCSNGSLQCRICI
H10N8	A/JX346/2013	(SEQ ID NO: 2254) MYKIVVIIALLGAVKGLDKICLGHHAVANGTIVKTLTNEQEEVTNATETVESTGINRLCMKGRKHKDLGNCHPIGMLIGTPACDLHLTGMWDTLIERENAIAYCYPGATVNVEALRQKIMESGGINKISTGFTYGSSINSAGTTRACMRNGGNSFYAELKWLVSKSKGQNFPQTTNTYRNTDTAEHLIMWGIHHPSSTQEKNDLYGTQSLSISVGSSTYRNNFVPVVGARPQVNGQSGRIDFHWTLVQPGDNITFSHNGGLIAPSRVSKLIGRGLGIQSDAPIDNNCESKCFWRGGSINTRLPFQNLSPRTVGQCPKYVNRRSLMLATGMRNVPELIQGGRGLFGAIAGFLENGWEGMVDGWYGFRHQNAQGTGQAADYKSTQAAIDQITGKLNRLVEKTNTEFESIESEFSEIEHQIGNVINWTKDSITDIWTYQAELLVAMENQHTIDMADSEMLNLYERVRKQLRQNAEEDGKGCFEIEIYHACDDSCMESIRNNTYDHSQYREEALLNRLNINPVTLSSGYKDIILILWFSFGASCFVLLAVVMGLFFFCLKNGNMRCTICI	(SEQ ID NO: 2255) MYKIVVIIALLGAVKGLDKICLGHHAVANGTIVKTLTNEQEEVTNATETVESTGINRLCMKGRKHKDLGNCHPIGMLIGTPACDLHLTGMWDTLIERENAIAYCYPGATVNVEALRQKIMESGGINKISTGFTYGSSINSAGTTRACMRNGGNSFYAELKWLVSKSKGQNFPQTTNTYRNTDTAEHLIMWGIHHPSSTQEKNDLYGTQSLSISVGSSTYRNNFVPVVGARPQVNGQSGRIDFHWTLVQPGDNITFSHNGGLIAPSRVSKLIGRGLGIQSDAPIDNNCESKCFWRGGSINTRLPFQNLSPRTVGQCPKYVNRRSLMLATGMRNVPELIQG	(SEQ ID NO: 2256) RGLFGAIAGFLENGWEGMVDGWYGFRHQNAQGTGQAADYKSTQAAIDQITGKLNRLVEKTNTEFESIESEFSEIEHQIGNVINWTKDSITDIWTYQAELLVAMENQHTIDMADSEMLNLYERVRKQLRQNAEEDGKGCFEIYHACDDSCMESIRNNTYDHSQYREEALLNRLNINPVTLSSGYKDIIL	(SEQ ID NO: 2257) WFSFGASCFVLLAVVMGLFFF	(SEQ ID NO: 2258) CLKNGNMRCTICI

H7N9	A/Anhui/1/2013	(SEQ ID NO: 2259) MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI	(SEQ ID NO: 2260) MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKKTVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGQSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEIPKG	(SEQ ID NO: 2261) RGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVIL	(SEQ ID NO: 2262) WFSFGASCFILLAIVMGLVFI	(SEQ ID NO: 2263) CVKNGNMRCTICI
H7N9	.A/Jiangsu/2/2013(H242)	(SEQ ID NO: 2264) MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMTVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYReEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI	(SEQ ID NO: 2265) MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRICSKGKMTVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQILRESGGIDKEAMGFTYSGIRTNGATSACRRSGSSFYAEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRIDFHWLMLNPNDTVTFSFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEIPKG	(SEQ ID NO: 2266) RGLFGAIAGFIENGWEGLIDGWYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYReEAMQNRIQIDPVKLSSGYKDVIL	(SEQ ID NO: 2267) WFSFGASCFILLAIVMGLVFI	(SEQ ID NO: 2268) CVKNGNMRCTICI

In the Table, TM stands for transmembrane and CY stands for cytoplasmic.

[1007] Following the vaccine generation strategy above, engineered antigens may also be used to develop pan-flu vaccines. Such constructs are shown in Table 30.

Table 30. Engineered antigens encoded by NAV polynucleotides

Virus	Strain	Configuration	Protein engineering	Full length Sequence	SEQ ID NO.
H1N1	PR8/34	HA(PR8)-Ferritin	fusion protein of HA antigen with ferritin (H pylori) to form nanoscaffold, secreted	MKANLLVLLCALAAADADTICIGYHANNSTDTVDTV LEKNVTVTHSVNLLEDSHNGKLCRLKGIAPLQLGK CNIAGWLLGNPECDPLLPVRSWSYIVETPNSENGI CYPGDFIDYEELREQLSSVSSFERFEIFPKESSWP NHNTNGVTAACSHEGKSSFYRNLLWLTEKEGSYP KLKNSYVNKKGKEVLVLWGIHHPPNSKEQQNLYQ NENAYVSVVTSNYNRRFTPEIAERPKVRDQAGRM NYYWTLLKPGDTIIFEANGNLIAPMYAFALSRGFGS GIITSNASMHECNTKCQTPLGAINSSLPYQNIHPVT IGECPKYVRSAKLRMVTGLRNNPSIQSRGLFGAIA GFIEGGWTGMIDGWYGYHHQNEQGSGYAADQK STQNAINGITNKVNTVIEKMNIQFTAVGKEFNKLEK RMENLNKKVDDGFLDIWTYNAELLVLLENERTLDF HDSNVKNLYEKVKSQLKNNAKEIGNGCFEFYHKC DNECMESVRNGTYDYPKYSEESKLNREKVDSGG DIIKLLNEQVNKEMQSSNLYMSMSSWCYTHSLDG AGLFLFDHAAEEYEHAKKLIIFLNENNVPVQLTSISA PEHKFEGLTQIFQKAYEHEQHISESINNIVDHAIKSK DHATFNFLQWYVAEQHEEEVLFKDILDKIELIGNEN HGLYLADQYVKGIAKSRKS	2269
H1N1	A/New Caledonia/20/1999 (1999 NC)	HA(NC1999)-Ferritin	fusion protein of HA antigen with ferritin (H pylori) to form nanoscaffold, secreted	MKAKLLVLLCTFTATYADTICIGYHANNSTDTVDTV LEKNVTVTHSVNLLEDSHNGKLCLLKGIAPLQLGN CSVAGWILGNPECELLISKESWSYIVETPNPENGT CYPGYFADYEELREQLSSVSSFERFEIFPKESSWP NHTVTGVSASCSHNGKSSFYRNLLWLTGKNGLYP NLSKSYVNNKEKEVLVLWGVHHPPNIGNQRALYH TENAYVSVVSSHYSRRFTPEIAKRPKVRDQEGRIN YYWTLLEPGDTIIFEANGNLIAPWYAFALSRGFGS GIITSNAPMDECDAKCQTPQGAINSSLPFQNVHPV TIGECPKYVRSAKLRMVTGLRNIPQRETRGLFGAI AGFIEGGWTGMVDGWYGYHHQNEQGSGYAADQ KSTQNAINGITNKVNSVIEKMNTQFTAVGKEFNKL ERRMENLNKKVDDGFLDIWTYNAELLVLLENERTL DFHDSNVKNLYEKVKSQLKNNAKEIGNGCFEFYH KCNNECMESVKNGTYDYPKYSEESKLNREKIDSG GDIIKLLNEQVNKEMQSSNLYMSMSSWCYTHSLD GAGLFLFDHAAEEYEHAKKLIIFLNENNVPVQLTSI SAPEHKFEGLTQIFQKAYEHEQHISESINNIVDHAIK SKDHATFNFLQWYVAEQHEEEVLFKDILDKIELIGN ENHGLYLADQYVKGIAKSRKSGS	2270

H1N1	PR8/34	HA1(-) stem only HA antigen	Headless HA antigen (Stem only), transmembrane,	MKANLLVLLCALAAADADTICIGYHANNSTDTVDTV LEKNVTVTHSVNLLEDSHNGKLCGGGGCNTKCQT PLGAINSSLPYQNIHPVTIGECPKYVRSAKLRMVT GLRNIPSIQSRGLFGAIAGFIEGGWTGMIDGWYGY HHQNEQGSGYAADQKSTQNAINGITNKVNTVIEK MNIQFTAVGKEFNKLEKRMENLNKKVDDGFLDIW TYNAELLVLLENERTLDFHDSNVKNLYEKVKSQLK NNAKEIGNGCFEFYHKCDNECMESVRNGTYDYPK YSEESKLNREKVDGVKLESMGIYQILAIYSTVASSL VLLVSLGAISFWMCSNGSLQCRICI	2271
H1N1	PR8/34	Monoglycosylated HA1(-) stem only HA antigen	Headless HA antigen (Stem only), transmembrane, monglycosylated by N76D, N270D	MKANLLVLLCALAAADADTICIGYHANNSTDTVDTV LEKNVTVTHSVNLLEDSHNGKLCGGGGCNTKCQT PLGAIDSSLPYQNIHPVTIGECPKYVRSAKLRMVT GLRNIPSIQSRGLFGAIAGFIEGGWTGMIDGWYGY HHQNEQGSGYAADQKSTQNAINGITNKVNTVIEK MNIQFTAVGKEFNKLEKRMENLNKKVDDGFLDIW TYNAELLVLLENERTLDFHDSNVKNLYEKVKSQLK NNAKEIGNGCFEFYHKCDNECMESVRDGTYDYPK YSEESKLNREKVDGVKLESMGIYQILAIYSTVASSL VLLVSLGAISFWMCSNGSLQCRICI	2272
H1N1	PR8/34	aglycosylated HA1(-) Stem only HA antigen	Headless HA antigen (Stem only), transmembrane, aglycosylated by N27D,N28D, N40D, N76D, N279D	MKANLLVLLCALAAADADTICIGYHADDSTDTVDTV LEKDVTVTHSVNLLEDSHNGKLCGGGGCNTKCQT PLGAIDSSLPYQNIHPVTIGECPKYVRSAKLRMVT GLRNIPSIQSRGLFGAIAGFIEGGWTGMIDGWYGY HHQNEQGSGYAADQKSTQNAINGITNKVNTVIEK MNIQFTAVGKEFNKLEKRMENLNKKVDDGFLDIW TYNAELLVLLENERTLDFHDSNVKNLYEKVKSQLK NNAKEIGNGCFEFYHKCDNECMESVRDGTYDYPK YSEESKLNREKVDGVKLESMGIYQILAIYSTVASSL VLLVSLGAISFWMCSNGSLQCRICI	2273
H1N1	PR8/34	HA1(-) stem HA -ferritin	Headless HA antigen (stem only) fused to ferritin nanoscaffold, secreted	MKANLLVLLCALAAADADTICIGYHANNSTDTVDTV LEKNVTVTHSVNLLEDSHNGKLCGGGGCNTKCQT PLGAINSSLPYQNIHPVTIGECPKYVRSAKLRMVT GLRNIPSIQSRGLFGAIAGFIEGGWTGMIDGWYGY HHQNEQGSGYAADQKSTQNAINGITNKVNTVIEK MNIQFTAVGKEFNKLEKRMENLNKKVDDGFLDIW TYNAELLVLLENERTLDFHDSNVKNLYEKVKSQLK NNAKEIGNGCFEFYHKCDNECMESVRNGTYDYPK YSEESKLNREKVDSGGDIIKLLNEQVNKEMQSSNL Y MSMSSWCYTHSLDGAGLFLFDHAAEEYEHAKKLII FLNENNVPVQLTSISAPEHKFEGLT QIFQKAYEHEQHISESINNIVDHAIKSKDHATFNFL QWYVAEQHEEEVLFKDILDKIELI GNENHGLYLADQYVKGIAKSRKSGS	2274

Example 15. Influenza study - H1N1 Hemagglutinin (HA) Antigen

[1008] The instant study was designed in two phases, first to test the the immunogenicity of mRNA vaccines encoding HA from Influenza AIPR18 (H1N1) in mice and second to test the efficacy of candidate influenza vaccines in mice against a lethal challenge with influenza (INFV) A/ PR/8/34 (H1N1). The study design is outlined in Table 31. The NAV polynucleotide used in the study was construct 4 from Table 28 (ORF SEQ ID NO: 2045, mRNA SEQ ID NO: 2118).

[1009] In phase I of this study, mice were vaccinated on week 0 and 3 via intravenous (IV), intramuscular (IM), or intradermal (ID) routes. One group remained unvaccinated and one was administered inactivated PR8 antigen. Serum was collected from each mouse on weeks 1, 3 (pre-dose), and 5. Individual bleeds were tested for anti-HA activity via virus neutralization assay and HA inhibition (HAI) from all three time points and pooled samples from week 5 only were

tested by Western blot using AIPR18134 (H1N1).

Phase II

[1010] Since the desired immune response was achieved (HAI titer> 40), phase II of the study was performed. In phase II, mice were challenged with a lethal dose ($10\times LD_{90}$; ~100 plaque-forming units; PFU) of A/PR/8/34 (H1N1) via intranasal (IN) instillation. Mice were monitored for weights, health, and survival for 14 days.

[1011] This study tested the immunogenicity of mRNA vaccines encoding HA from Influenza A/PR/8 (H1N1) in mice. The study utilized 12 groups of five BALB/c female mice. Mice were vaccinated on week 0 and 3 via intranasal (positive control), intravenous, intramuscular, or intradermal routes. One group was unvaccinated and one administered inactivated PR8 antigen via intranasal vaccination.

[1012] This study tested whether candidate ribonucleic acid vaccines and formulations could protect mice from lethal influenza A/PR/8/34 (H1N1) infection. The mice used were 6-8 week old female BALBlc mice in groups of 10. Mice were vaccinated on weeks 0 and 3via IM, ID, or IV route. Mouse serum was tested for microneutralization and HAI. Mice were then challenged with ~1 LD_{90} of influenza A/PR/8/34 (H1N1) on week 7 administered intranasally (IN). Endpoint was day 13 post infection, death, or euthanasia. Animals displaying severe illness as determined by >30% weight loss, extreme lethargy, or paralysis were euthanized. Temperature and weights were taken daily.

[1013] The LNP formulation consisted of a cationic lipid, non-cationic lipid, PEG lipid and structural lipid in the ratios 50:10:1.5:38.5. The cationic lipid was DLin-KC2-DMA (50 mol%), the non-cationic lipid was DSPC (10 mol%), the PEG lipid was PEG-DOMG (1.5 mol%) and the structural lipid was cholesterol (38.5 mol%).

[1014] Serum was collected from each mouse on weeks 1, 3 (pre-dose), and 5. Individual bleeds were tested for anti-HA activity via virus neutralization assay and HA inhibition (HAI) from all three time points (individual animals) and pooled samples from week 5 only were tested by Western blot using inactivated Influenza AIPR18 (H1N1).

Standard Protocol for intranasal infection of mice

[1015] Female 6-8 week old BALB/c mice were housed in groups of 5 mice. Mice were quarantined at the study site (Noble Life Sciences, Gaithersburg, MD) for at least 3 days prior to the start of the study. Food and water was provided ad libitum.

[1016] The groups of mice challenged with INFV were infected via intranasal (IN) inoculation with ~10xLD90 in 100 pL of INFV in PBS under light anesthesia (Isoflurane). After the infection, mice were placed back into their cages for observation and subsequent dosing.

Microneutralization

[1017] The microneutralization protocol was the standard protocol for microneutralization of influenza as published by the World Health Organzation, 6 December, 2010 and outlined by the WHO Collaborating Center for Surveillance, Epidemiology and Control of Influenza, Centers for Disease Control and Prevention, Atlanta, USA.

Serologic diagnosis of influenza virus infections by hemagglutination inhibition.

[1018] The standard protocol for inhibition of hemagglutination can be found in publication WHO Influenza Manual, WHO/CDS/CSR/NCS/2002.5 Rev. 1, at page 37.

[1019] Briefly, the protocol is as follows.

Procedure

[1020]

I. Treatment of sera

1. Identification of Influenza Isolates by Hemagglutination.
2. Adsorb sera to remove nonspecific agglutinins as described above.

II. Standardization of RBCs

Standardize RBCs according to the procedure in art recognized procedures.

III. HA Titration of control antigens

Following the procedure for titration of the control antigens given above.

IV. Preparation of standardized antigens for HAI test and "Back Titration"

Each control antigen must be standardized to contain 4 HA units/25 μl, 8 HA units/50 μl.
Note: Four HA units are added to the test in 25 μl because the HA unit calculations are based on a volume of 50 μl.

V. HAI Test for serologic diagnosis

1. Label appropriate microtiter plates
2. Add 25 μl of PBS to wells B through H (B1 - H12) of each numbered column.
3. Add 50 μl of each treated serum (1:10) to the appropriate first well (A1-A12) of the numbered column.
4. Prepare serial twofold dilutions of the treated sera by transferring 25 μl from the first well of the numbered columns 1-12 to successive wells. Discard the final 25 μl after row H.
5. Add 25 μl of standardized antigen to all wells (A1 - H12) in a set of treated sera.
6. Add 25 μl of PBS instead of antigen to the set of treated sera for serum controls (A1 - H12).
7. Mix the contents of the plates by shaking on a mechanical vibrator for 10 see or by agitating the plates manually.
8. Cover the plates and incubate at room temperature (22 to 25oC.) for 30-45 min.
9. Add 50 μl of standardized RBCs to all wells. Mix as before.
10. Cover the plates and allow the RBCs to settle at room temperature (22 to 25C) for the appropriate time according to RBCs being used.
11. Record the HAI titers.

[1021] The positive control antigens and corresponding antisera should give consistent results when compared with previous tests. A fourfold titer increase between the acute and convalescent serum is considered diagnostically positive for that influenza type/subtype.

Western Blots

[1022] Serum from bleeds performed on vaccinated mice on week 5 was pooled and diluted 1:1000 for analysis in Western blots. Inactivated PR8 antigen was separated via SDS-PAGE and probed with the pooled sera. Positive signals were observed in groups 2-7 and 12, with the highest signals observed in groups 4-7 and 12.
[1023] 500 ng per lane of inactivated PR8 antigen was separated via SDS- on 10% bis-tris gels. Protein was transferred to Polyvinylidene Difluoride (PVDF) membranes and membranes were blocked with milk and probed with a 1:1000 dilution of pooled sera from each group from week 5 bleeds.
[1024] Antibody was detected with 1:3000 dilution of goat anti-mouse alkaline phosphatase (AP)

Observation of mice

[1025] Mice were observed through 13 days post infection (14 days total, 0-13 days post infection).
[1026] Mice were weighed daily on an Ohause scale and the weights were recorded.
[1027] All animals had chips implanted at least 3 days prior to virus challenge that monitored the body temperature. The temperatures were recorded daily.
[1028] Survival and health of each mouse was evaluated once time a day.

Table 31. Study Design

Group	Mouse Strain	Vaccine (n=5 mice/group) Delivered week 0, 3	Dosage/ Route	Readouts
1	Female BALB/c, 6-8 weeks of age	N/A	N/A	Animals displaying severe illness will be euthanized. Serum samples collected on weeks 1, 3, and 5. Serum analyzed via Western blot (week 5 pooled samples), virus neutralization assay (all individuals), and HAI (all individuals).
2		Unmodified	LNP IV, 0.4 mg/kg	
3		N1-methyl pseudouridine/5-methyl cytosine		
4		Unmodified	IM, LNP 0.4 mg/kg	
5		N1-methyl pseudouridine /5-methyl cytosine		
6		Unmodified	ID, LNP 0.4 mg/kg	
7		N1-methyl pseudouridine /5-methyl cytosine		
8		Unmodified	ID, 80 ug w/ lipoplex (Lipofectamine 2000)	
9		N1-methyl pseudouridine /5-methyl cytosine		
10		Unmodified	ID, 80 ug naked mRNA	
11		N1-methyl pseudouridine /5-methyl cytosine		
12		Control	IN, Inactivated PR8 virus	

[1029] In the Table the following abbreviations apply; IM, intramuscular; ID, intradermal; IN, intranasal; IV, intraveneous; LNP, lipid nanoparticle

Table 32. Mean HAI Titers

Sample ID	Week 1	Week 3	Week 5
Naïve	8	7	7
Unmodified; LNP IV, 0.4 mg/kg	8	28	635
N1-methyl pseudouridine/5-methyl cytosine; LNP IV, 0.4 mg/kg	25	139	2004
Unmodified; LNP IM, 0.4 mg/kg	22	63	2560
N1-methyl pseudouridine/5-methyl cytosine; LNP IM, 0.4 mg/kg	60	482	2803
Unmodified; LNP ID, 0.4 mg/kg	39	279	3796
N1-methyl pseudouridine/5-methyl cytosine; LNP ID, 0.4 mg/kg	114	965	10152
Unmodified; 80 ug w/ lipoplex (Lipofectamine 2000)	7	17	197
N1-methyl pseudouridine/5-methyl cytosine; 80 ug w/ lipoplex (Lipofectamine 2000)	7	20	28
Unmodified; ID, 80 ug naked mRNA	7	7	61
N1-methyl pseudouridine/5-methyl cytosine; ID, 80 ug naked mRNA	7	16	133
Control PR8 Antigen	14	14	441

Inhibition of Hemagglutination

[1030] inhibition of hemagglutination (HAI) was measured in mouse sera samples on weeks 1, 3, and 5 post-vaccination.

After week 1, both groups 5 and 7 displayed HAI titers over 40, at 60 and 114, respectively. On week 3, groups 3-7 displayed HAI activity over 40, with the highest being group 7 at 965. On week 5, all groups except 1 (naive) and 9 displayed HAI activity over 40, with group 7 >10,000.

[1031] It is noted that 1:40 is predictive of efficacy.An HAI titer of >40 is deemed necessary to protect from a lethal challenge of influenza.

[1032] The data showed that there was 100% rescue from lethal influenza challenge with rapid onset of protective antibody titers after 1 week and high antibody titers, i.e., 50 fold over unmodified mRNA and 20 fold over the protein vaccine. Furthermore, it was shown that for ribonucleic acid vaccines of the invention a much lower effective mRNA dose is required, i.e., ten fold less than unmodified mRNA. (Figure 10).

Microneutralization

[1033] Two-fold dilutions of mouse sera were added to 100 TCID50/ml of virus in 96-well plates. After 24-hours of incubation, 1.5x104 Madin-Darby Canine-Kidney cells were added to each well. After a ~20 hour incubation at 37◦C, virus was detected and scored with an anti-NP antibody and read at 490nm. No neutralization activity was detected with week 1 samples (signal <50; lower limit of detection). By week 3, mice in groups 5 and 7 displayed neutralizing activity between 79 and 250 (group 5) and 250 (group 7). Neither of the other groups displayed any neutralizing activity On week 5, groups 2-4 showed high neutralizing activity between 789 and 2493, with group 7 displaying neutralizing activity 2494 and -25,000. The control group of mice, vaccinated with inactivated PR8, displayed neturalizing activity in 3 of 5 mice and ranged between 79 and 250. The data are shown in Tables 33-35.

Table 33. Week 1 Microneutralization

Sample ID	Mouse 1	Mouse 2	Mouse 3	Mouse 4	Mouse 5
Naïve	<50*	<50	<50	<50	<50
Unmodified; LNP IV, 0.4 mg/kg	<50	<50	<50	<50	<50
N1-methyl pseudouridine/5-methyl cytosine; LNP IV, 0.4 mg/kg	<50	<50	<50	<50	<50
Unmodified; LNP IM, 0.4 mg/kg	<50	<50	<50	<50	<50
N1-methyl pseudouridine/5-methyl cytosine; LNP IM, 0.4 mg/kg	<50	<50	<50	<50	<50
Unmodified; LNP ID, 0.4 mg/kg	<50	<50	<50	<50	<50
N1-methyl pseudouridine/5-methyl cytosine; LNP ID, 0.4 mg/kg	<50	<50	<50	<50	<50
Unmodified; 80 ug w/ lipoplex (Lipofectamine 2000)	<50	<50	<50	<50	<50
N1-methyl pseudouridine /5-methyl cytosine; 80 ug w/ lipoplex (Lipofectamine 2000)	<50	<50	<50	<50	<50
Unmodified; ID, 80 ug naked mRNA	<50	<50	<50	<50	<50
N1-methyl pseudouridine/5-methyl cytosine; ID, 80 ug naked mRNA	<50	<50	<50	<50	<50
Control PR8 Antigen	<50	<50	<50	<50	<50

*Titers of '<50' indicate titers below Limit of detection (1:50)

Table 34. Week 3 Microneutralization

Sample ID	Mouse 1	Mouse 2	Mouse 3	Mouse 4	Mouse 5
Naïve	<50*	<50	<50	<50	<50
Unmodified; LNP IV, 0.4 mg/kg	<50	<50	<50	<50	<50
N1-methyl pseudouridine/5-methyl cytosine; LNP IV, 0.4 mg/kg	<50	<50	<50	<50	<50
Unmodified; LNP IM, 0.4 mg/kg	<50	<50	<50	<50	<50
N1-methyl pseudouridine/5-methyl cytosine; LNP IM, 0.4 mg/kg	79	79	79	250	250
Unmodified; LNP ID, 0.4 mg/kg	<50	<50	<50	<50	<50
N1-methyl pseudouridine/5-methyl cytosine; LNP ID, 0.4 mg/kg	250	250	250	250	250
Unmodified; 80 ug w/ lipoplex (Lipofectamine 2000)	<50	<50	<50	<50	<50
N1-methyl pseudouridine /5-methyl cytosine; 80 ug w/ lipoplex (Lipofectamine 2000)	<50	<50	<50	<50	<50
Unmodified; ID, 80 ug naked mRNA	<50	<50	<50	<50	<50
N1-methyl pseudouridine/5-methyl cytosine; ID, 80 ug naked mRNA	<50	<50	<50	<50	<50
Control PR8 Antigen	<50	<50	<50	<50	<50

*Titers of '<50' indicate titers below Limit of detection (1:50)

Table 35. Week 5 Microneutralization

Sample ID	Mouse 1	Mouse 2	Mouse 3	Mouse 4	Mouse 5
Naïve	<50*	<50	<50	<50	<50
Unmodified; LNP IV, 0.4 mg/kg	2493	789	789	<50	250
N1-methyl pseudouridine/5-methyl cytosine; LNP IV, 0.4 mg/kg	789	789	2493	789	2493
Unmodified; LNP IM, 0.4 mg/kg	789	789	2493	789	2493
N1-methyl pseudouridine/5-methyl cytosine; LNP IM, 0.4 mg/kg	789	2493	2493	789	789
Unmodified; LNP ID, 0.4 mg/kg	<50	<50	<50	<50	<50
N1-methyl pseudouridine/5-methyl cytosine; LNP ID, 0.4 mg/kg	7877	7877	24892	2493	7877
Unmodified; 80 ug w/ lipoplex (Lipofectamine 2000)	<50	<50	<50	<50	79
N1-methyl pseudouridine /5-methyl cytosine; 80 ug w/ lipoplex (Lipofectamine 2000)	<50	<50	<50	<50	<50
Unmodified; ID, 80 ug naked mRNA	<50	79	<50	<50	<50
N1-methyl pseudouridine/5-methyl cytosine; ID, 80 ug naked mRNA	<50	<50	<50	<50	<50
Control PR8 Antigen	250	79	<50	<50	250

*Titers of '<50' indicate titers below Limit of detection (1:50)

Survival

[1034] All mice were challenged with a lethal dose (10xLD90) of INFV AIPR18134 on week 7 post-vaccination. Mice were observed for morbidity and mortality for up to 14 days. All vaccinated mice displayed 100% survival, compared to the naive group (group 1), which was not vaccinated. 12 groups of 5 mice were challenged via IN instillation with -100 PFU of INFV A/PR/8/34 (H1N1). Mice were observed daily for 14 days for health, morbidity, and mortality. All animals except for unvaccinated, which died at approximately day 7, survived the 14 day study.

Weight loss data

[1035] Weight loss and health of mice challenged with AIPRI8134. 12 groups of 5 mice were challenged via IN instillation with -100 PFU of INFV A/PR/8/34 (H1N1). Mice were observed daily for 14 days for health, morbidity, and mortality.

[1036] All vaccinated mice displayed 100% survival, although some groups displayed weight loss. The unvaccinated group displayed 0% survival and died on days 6 and 7, post-infection. While differences in weight-loss were observed in challenged vaccinated mice, no conclusion can be drawn as to its significance. The group vaccinated with NAVs having N1 methylpseudouridine/5-methyl cytosine with naked mRNA displayed health scores of '2' on days 2 and 3 post-infection, but made a recovery to 'healthy' for the rest of the study. The groups vaccinated with NAVs having N1 methylpseudouridine/5-methyl cytosine ID 80 ug w/ lipofecatmine 2000 (LF2000) and N1 methylpseudouridine and 5-methylcytosine LNP ID both displayed health scores of '2' on days 5 and 6 post-infection which continued through the duration of the study. The LNP formulation consisted of a cationic lipid, non-cationic lipid, PEG lipid and structural lipid in the ratios 50:10:1.5:38.5. The cationic lipid was DLin-KC2-DMA (50 mol%), the non-cationic lipid was DSPC (10 mol%), the PEG lipid was PEG-DOMG (1.5 mol%) and the structural lipid was cholesterol (38.5 mol%).

[1037] All groups generally maintained weight between 90-110% of original weight with the exception of the unvaccinated group. These animals were at approximately 70% of their original weight by day 6. Health scoring was per Table 36.

Table 36. Health score chart

Score	Initial	Description	Appearance	Mobility	Attitude
1	H	Healthy	Smooth coat, bright eyes	Active, scurrying, burrowing	Alert
2	SR	Slighly Ruffled	Slightly ruffled coat (usually only around heand and neck	Active, scurrying, burrowing	Alert
3	R	Ruffled	Ruffled coat throughout body. A "wet" appearance	Active, scurrying, burrowing	Alert
4	S	Sick	Verr ruffled coat. Slightly closed, inset eyes	Walking, but not scurrying	Mildly lethargic
5	VS	Very Sick (euthanized)	Very ruffled coato, closed, inset eyes	Slow to no movement; will return upright position if put on side	Extremely lethargic
6	E	Euthanize	Very ruffled coat, closed inset eyes; moribund requires human euthenasia	No movement or uncontrollable; spastic movement. Will not return to upright	Completely unaware or in noticeable distress
7	D	Deceased	--	--	--

Example 16. MRSA study (Reference Example)

[1038] Antigens of MRSA which may be encoded by the ribonucleic acid vaccines include, SpA, SpAKKAA, IsdA, IsdB, SDRD, SDRE, TSST-1, PVL, a-HL, NMD-1 and SCCmec. The study design is shown in Table 37. In these studies, construct number 27 from Table 28 was used.

A. Test for efficacy of modified ribonucleic acid vaccines in *Staph, aureus* pneumonia challenge model in mice

[1039] This study will test the efficacy of ribonucleic acid vaccine encoding MRSA Ag808 in BALB/c mice. The study utilizes 15 groups of 15 BALB/c female mice (225 total. Mice are vaccinated on week 0 and 3 via intradermal (ID) or intramuscular (IM) route with either an LNP Formulation comprising DLin-KC2-DMA ("KC2") or DLin-MC3-DMA ("MC3"). The KC2 LNP formulation consisted of a cationic lipid (DLin-KC2-DMA, 50 mol%), non-cationic lipid (DSPC, 10 mol%), PEG lipid (PEG-DOMG 1.5 mol%) and a structural lipid (cholesterol, 38.5 mol%). The MC3 LNP formulation consisted of a cationic lipid (DLin-MC3-DMA, 50 mol%), non-cationic lipid (DSPC, 10 mol%), PEG lipid (PEG-DOMG 1.5 mol%) and a structural lipid (cholesterol, 38.5 mol%). One group is unvaccinated and one administered a positive control antigen. Prior to challenge, mice will be bled by tail vein on weeks 1, 3 and 5 and serum samples will be retained for later analysis. The mice will be challenged with MRSA (strain Newman) at ~1xLD90 via intranasal (IN) inoculation on week 5. Mice will be monitored for morbidity using a health score assigned based on a standard scoring system, weight loss, and mortality and the endpoint is day 14 post infection, death, or euthanasia. Animals displaying severe illness as determined by >30% weight loss, extreme lethargy, or paralysis will be euthanized. Mice will be held until week 7, to perform additional work with the animals (ex. further vaccinations, bleed or infectious challenge).

B. Test for Efficacy of N1-methylpseudouridine modified ribonucleic acid Vaccines in Staphylococcus aureus Peritonitis Challenge

[1040] This study tested the efficacy of a ribonucleic acid vaccine encoding MRSA Ag808 in BALB/c mice. The study utilized 15 groups of 15 BALB/c female mice (270 total. Mice were vaccinated on week 0 and 3 via intradermal (ID) or intramuscular (IM) route with either an LNP Formulation comprising DLin-KC2-DMA ("KC2") or DLin-MC3-DMA ("MC3"). The KC2 LNP formulation consisted of a cationic lipid (DLin-KC2-DMA, 50 mol%), non-cationic lipid (DSPC, 10 mol%), PEG lipid (PEG-DOMG 1.5 mol%) and a structural lipid (cholesterol, 38.5 mol%). The MC3 LNP formulation consisted of a cationic lipid (DLin-MC3-DMA, 50 mol%), non-cationic lipid (DSPC, 10 mol%), PEG lipid (PEG-DOMG 1.5 mol%) and a structural lipid (cholesterol, 38.5 mol%). One group was unvaccinated and one administered a positive control antigen. Prior to challenge, mice were bled by tail vein on weeks 1, 3 and 5 and serum samples were retained for later analysis. The mice that were challenged via intranasal instillation received a predicted challenge dose of 2e8 CFU/mouse (actual upon back-titration was 3.3e81mouse) and the mice that were challenged via IP infection received a predicted 1e7 CFU/mouse (actual upon back-titration 6.7e6 CFU/mouse). Mice were monitored for morbidity using a health score assigned based on a standard scoring system, weight loss, and mortality and the endpoint is day 14 post infection, death, or euthanasia. Animals displaying severe illness as determined by >30% weight loss, extreme lethargy, or paralysis

were euthanized.

Table 37. Study Design

Group (n=15)	Antigen	Dosage	Route	Chemistry	Formulation	Administration	Samples
1	N/A	N/A	N/A	N/A	NaCl	Vaccination on weeks 0 and 3	Bled by tail vein on weeks 1, 3, and 5; retained for future analysis
2	MRSA Ag808	0.4mg/kg	ID	N1methyl-pseudoruridine	LNP; KC2		
3		0.08 mg/kg					
4		0.016mg/kg					
5		0.4mg/kg	IM				
6		0.08 mg/kg					
7		0.016mg/kg					
8		0.4mg/kg	ID		LNP; MC3		
9		0.08 mg/kg					
10		0.016mg/kg					
11		0.4mg/kg	IM				
12		0.08 mg/kg					
13		0.016mg/kg					
14	Inactivated bacteria	TBD	TBD	N/A	TBD		
15	Inactivated bacteria	TBD	TBD	N/A	TBD		

[1041] Mice that were vaccinated and challenged via IP route with 3% hog mucin all displayed 0% survival within 24 hours of challenge. Mice that were vaccinated and challenged via IN instillation displayed between 6 and 33% survival and median survival times between 2 and 3 days. Efficacy was not shown with either the tested RNA vaccine construct or the controls (inactivated bacteria and protein control) in either challenge model, suggesting that the model was not adequate for testing the constructs. The vehicle-vaccinated group displayed 20% survival and a median survival time of 3 days. Without being bound in theory, is is believed that the severity of the MRSA infection model precluded detection of vaccine efficacy. Other models can be tested to determine efficacy.

Example 17. Dengue study: Dengue Virus RNA Vaccine Immunogenicity in Mice

[1042] This study provides a preliminary analysis of the immunogenicity of a nucleic acid mRNA vaccine using a dengue virus (DENV) serotype 2 antigen in BALB/c mice. The study utilizes 44 groups of 10 BALBIc female (5) and male (5) mice (440 total, 6-8 weeks of age at study initiation, see Table38 for design summary). In this study, construct numbers used are referenced and found in Table 28.

[1043] Mice were vaccinated on weeks 0 and 3 via intramuscular (IM) or intradermal (ID) routes. One group remained unvaccinated and one was administered 10^5 plaque-forming units (PFU) live DENV2, D2Y98P isolate via intravenous (IV) injection as a positive control. Serum was collected from each mouse on weeks 1, 3, and 5; bleeds on weeks 1 and 3 were in-life samples (tail vein or submandibular bleeds) and week 5 will be a terminal (intracardiac) bleed. Individual serum samples were stored at -80°C until analysis by neutralization or microneutralization assay. Pooled samples from each group at the week 5 time points were tested by Western blot for reactivity with viral lysate.

Table 38. Detailed experimental design (treatment, readouts)

Group	Mouse Strain	Vaccine (n=10, female) mice/group) Delivered week 0 and 3	Chemistry	Formulation/ Route	Dose	Readouts
1	Female BALB/c, 6-8 weeks of age	N/A		N/A	N/A	Serum samples collected on weeks 1, 3, and 5. Serum analyzed via Western blot
2		DEN2Y98-PrME (construct 23 from Table 28)	N1-methylpseudouridine/5-methyl cytosine	ID	0.4 mg/kg in LNP	
3				IM		
4				ID	0.08 mg/kg in LNP	
5				IM		
6				ID	0.016 mg/kg in LNP	
7				IM		
8			N1-methylpseudouridine	ID	0.4 mg/kg in LNP	
9				IM		
10				ID	0.08 mg/kg in LNP	
11				IM		
11				ID	0.016 mg/kg in LNP	
12				IM		
13		DEN2Y98-PrME80 (construct 24 from Table 28)	N1-methylpseudouridine/5-methyl cytosine	ID	0.4 mg/kg in LNP	
14				IM		
15				ID	0.08 mg/kg in LNP	
16				IM		

17				ID	0.016 mg/kg in LNP	
18				IM		
19			N1-methylpseudouridine	ID	0.4 mg/kg in LNP	
20				IM		
21				ID	0.08 mg/kg in LNP	
22				IM		
23				ID	0.016 mg/kg in LNP	
24				IM		
25		DEN2Y98-PrME80-DC (construct 25 from Table 28)	N1-methylpseudouridine/5-methyl cytosine	ID	0.4 mg/kg in LNP	
26				IM		
27				ID	0.08 mg/kg in LNP	
28				IM		
29				ID	0.016 mg/kg in LNP	
30				IM		
31			N1-methylpseudouridine	ID	0.4 mg/kg in LNP	
32				IM		
33				ID	0.08 mg/kg in LNP	
34				IM		
35				ID	0.016 mg/kg in LNP	
36				IM		
37		DEN2-DIII-Ferritin (construct 18 from Table 28)	N1-methylpseudouridine	ID	0.4 mg/kg in LNP	
38				IM		
39				ID	0.08 mg/kg in LNP	
40				IM		
41				ID	0.016 mg/kg in LNP	
42				IM		
43		Control, D2Y98P live virus	--	IV	10^5 PFU	

[1044] Signal was detected in groups 5, 15, 39, and 44 (live virus control) by a band that appeared between 50 and 60 kDa in the Western blot data. The data suggests that a mRNA vaccine to a single dengue viral antigen can produce antibody in preliminary studies.

Example 18. Tuberculosis ribonucleic acid vaccine: Combinatorial approach of adjuvants and antigens (Reference Example)

[1045] The objective of the study is to identify a multi-valent, multi-adjuvant vaccine effective in different diseases stages of tuberculosis. Initial experiment assesses 12 antigens with 8 cytokine adjuvants in three disease stage models. The antigens encoded by the polynucleotides of the invention include Ag85A (Rv3804c), Ag85B (Rv1886c), TB10.4 (Rv0288), ESAT6 (Rv3785), Rv2660L, Rv3619, Rv1813c, Rv3620c, Rv2608, Rv1196, Rv0125 and/or MT401. Target cytokine adjuvants include GM-CSF, IL-17, IFNg, IL-15, IL-2, IL-21, Anti-PD1/2 and/or Lactoferrin.

Example 19. Human enterovirus (HEV68 and HEV71) study.

[1046] Several key antigens for use in generating vaccines have been identified. These include: (i) VP1 BC/DE loop of all three HEV68 lineages and (ii) VP1+VP2 of HEV71. The RNAV polynucleotide for use in generating vaccines from the HEV71 antigen is construct 72 from Table 28 (ORF SEQ ID NO: 2113, mRNA SEQ ID NO: 2186).

Example 20. MERS-CoV study.

[1047] As MERS-CoV binds cells via DPP4Fc, treatment for MERS-CoV can include an mRNA encoding DPP4-Fc with or without MERS-CoA binding site and mutant binding sites for diabetes signaling and truncated receptor binding domain of MERS-CoV Spike protein. See PLoS One. 2013, 8(12): e81587). This vaccine would act as a decoy for the virus as it will attract the MERS-CoV.

[1048] Another key vaccine for MERS-CoV is identified as an mRNA encoding MERS-CoV Spike Glucoprotein as an antigen. The protein sequence is given here (SEQ ID NO: 2275).

SEQ ID NO. 2275:

mihsvfllmfllltptesyvdvgpdsiksacievdiqqtffdktwprpidvskadgiiypqgrtysnitityqglfpyqgdhgdmyvysaghatgttpqklfvanysqdvkqfangfvvrigaaanst
gtviispstsatirkiypafmlgssvgnfsdgkmgrffnhtlvllpdgcgtllrafycileprsgnhcpagnsytsfatyhtpatdcsdgnynrnaslnsfkeyfnlrnctfmytynitedeilewfgitqt
aqgvhlfssryvdlyggnmfqfatlpvydtikyysiiphsirsiqsdrkawaafyvyklqpltflldfsvdgyirraidcgfndlsqlhcsyesfdvesgvysvssfeakpsgsvveqaegvecdfs
pllsgtppqvynfkrlvftncnynltkllslfsvndftcsqispaaiasncysslildyfsyplsmksdlsvssagpisqfnykqsfsnptclilatvphnlttitkplkysyinkcsrllsddrtevpqlvna
nqyspcvsivpstvwedgdyyrkqlsplegggwlvasgstvamteqlqmgfgitvqygtdtnsvcpklefandtkiasqlgncveyslygvsgrgvfqnctavgvrqqrfvydayqnlvgyy
sddgnyyclracvsvpvsviydketkthatlfgsvacehisstmsqysrstrsmlkrrdstygplqtpvgcvlglvnsslfvedcklplgqslcalpdtpstltprsvrsvpgemrlasiafnhpiqv
dqlnssyfklsiptnfsfgvtqeyiqttiqkvtvdckqyvcngfqkceqllreygqfcskinqalhganlrqddsvrnlfasvkssqsspiipgfggdfnlttllepvsistgsrsarsaiedllfdkvtiad
pgymqgyddcmqqgpasardlicaqyvagykvlpplmdvnmeaaytssllgsiagvgwtaglssfaaipfaqsifyrlngvgitqqvlsenqkliankfnqalgamqtgftttneafhkvqd
avnnnaqalsklaselsntfgaisasigdiiqrldvleqdaqidrlingrlttlnafvaqqlvrsesaalsaqlakdkvnecvkaqskrsgfcgqgthivsfvvnapnglyfmhvgyypsnhievvs
ayglcdaanptnciapvngyfiktnntrivdewsytgssfyapepitslntkyvapqvtyqnistnlpppllgnstgidfqdeldeffknvstsipnfgsltqinttlldltyemlslqqvvkalnesyidl
kelgnytyynkwpwyiwlgfiaglvalalcvffilcctgcgtncmgklkcnrccdryeeydlephkvhvh

Example 21: H10N81n ***Vitro*** **Studies**

[1049] In vitro studies on the translatability of H10 hemagglutinin mRNA were performed in HeLa cells.

[1050] Western blot analysis revealed that HA is expressed in HeLa cells following transfection with the mRNA NAV encoding the HA protein of the H10N8 strain of influenza virus.

Example 22: Influenza Study - Dosing and Formulations

[1051] A mRNA vaccine encoding the HA protein of the H1N1 strain of influenza virus was tested at various doses and in various formulations for the ability ti illicit an immune response in mice.

[1052] The efficacy of different doses and formulations were evaluated in mice. HAI titers of mice determined on Day 35 after vaccination on Day 0 and Day 21 with mRNA encoding HA of the H1N1 virus are shown in Figure 11. Titers were highest at doses of 10 μg mRNA/mouse (400 μg mRNA/kg) with both the KC2 and MC3 formulations administered ID or IM (Table 39).

Table 39: Hemagglutinin inhibition titers in mice following vaccination with different doses and formulations of mRNA encoding the hemagglutinin protein of the H1N1 virus vaccine

Group No. (n=10	Route of	Vaccine	Formulation	Dose	Mean HAI titer ±

per group)	Vaccination on days 0 and 21			(µg/mouse)	SD
1	N/A	None	PBS	0	0
2	ID	H1N1/C1	KC2	10	1792±661
3				2	1088±309
4				0.4	272±152
5				0.08	52±44
6	ID	H1N1/ C0	KC2	10	992±648
7				2	692±743
8				0.4	140±78
9				0.08	35±31
10	ID	H1N1/ C0	MC3	10	1600±867
11				2	1152±588
12				0.4	290±254
13				0.08	37±25
14	IM	H1N1/ C0	MC3	10	1664±1422
15				2	1056±641
16				0.4	800±346
17				0.08	180±105
18	IN	Inactivated PR8 virus	PBS	20 µL	96±47
19	ID	H1N1/G2/C1	PBS	50	11±3

G0 = generation 0, i.e. canonical unmodified nucleotides; G2 = Generation 2 of the modified nucleotides; G5 = Generation 5 of the modified nucleotides (intended for clinical development).

C0 = CAP0; C1 = CAP1 (intended for clinical development)

N/A: Not applicable.

IN: intranasal: ID: intradermal; IM: intramuscular

Example 23: Influenza Study - Onset of Immunity - Survival and HAI

[1053] This study evaluated the efficacy of an mRNA vaccine encoding influenza A/PR/8 (H1N1) in female BALBlc mice following lethal challenge with influenza AIPR18 virus. Groups of animals were vaccinated and challenged after week 1, 2, 3, or 4. Serum was collected from each mouse one day prior to challenge. Individual bleeds were tested for hemagglutination inhibition using Influenza AIPR18 (H1N1). Control groups included unvaccinated mice and mice vaccinated with inactivated PR8 virus as a positive control. These animals were challenged in parallel with groups that were vaccinated with the mRNA vaccine. Challenge was performed with a lethal dose (100 PFU) of AIPR18134 (H1N1) via intranasal instillation. Animals were monitored for morbidity using a health score assigned based on a standard scoring system, weight loss and mortality. All mice receiving the mRNA vaccine showed 100% survival and minimal body weight loss (Figure 12A-12D). At week 1 post vaccination, hemagglutination inhibition titers of mice receiving the mRNA vaccine and mice receiving the positive control vaccine were similar. However, at weeks 2, 3 and 4 post vaccination, mice receiving the mRNA vaccine showed higher mean titers compared with mice receiving the positive control vaccine (Figure 13).

[1054] Groups of 15 female BALB/c mice (6-8 weeks of age) were vaccinated in week 0 with: mRNA encapsulated in LNP at a dose of 0.4 mg/kg by ID injection; inactivated PR8 virus given IN; orvehicle.

[1055] In weeks 1, 2, 3 and 4 after vaccination, animals were inoculated with influenza A/PR/8/34 virus under light anesthesia. The animals received 100 µL of the virus diluted in PBS to a final concentration of 1 x 103 PFU/mL via IN instillation

[1056] Health assessments were performed and weights were recorded daily for 14 days (days 0-13 post infection). Survival and health was evaluated using the scoring system shown in Table 36.

[1057] All animals had chips implanted at least 3 days prior to virus challenge that monitored the body temperature. The temperatures were recorded daily. One day prior to each challenge, serum samples were collected and the serum was analyzed in an HAI assay.

[1058] Survival curves are presented in Figure 15. The animals were monitored for survival for 14 days. Log-rank

analysis was performed in GraphPad Prism v6.

[1059] Mice vaccinated with mRNA vaccine or inactivated PR8 all displayed 100% survival, while mice that were unvaccinated displayed between 0 and 40% survival (Table 40.) No significant decreases in body weight of treated mice were detected.

Table 40: Percent and mean survival of mice after vaccination and challenge with influenza A/PR/8/34 virus

Group (n=15 per group)		% Survival	Mean Survival (Days)	Significance to Relative Vehicle Control (p values)
Vehicle	Week 1	10	8	N/A
	Week 2	40	10	N/A
	Week 3	0	9	N/A
	Week 4	6.67	9	N/A
Inactivated PR8 virus	Week 1	100	Undefined	<0.0001
	Week 2	100	Undefined	0.0004
	Week 3	100	Undefined	<0.0001
	Week 4	100	Undefined	<0.0001
mRNA vaccine encapsulated in LNP	Week 1	100	Undefined	<0.0001
	Week 2	100	Undefined	0.0004
	Week 3	100	Undefined	<0.0001
	Week 4	100	Undefined	<0.0001

[1060] Mean HA titers are shown in Figure 13 and Table 41.

Table 41. Mean hemaglutination inhibition (±SD) titers of mice after vaccination and challenge with influenza A/PR/8/34 virus

Vaccination in week 0	Challenge with influenza A/PR/8/34 virus			
	Week 1	Week 2	Week 3	Week 4
Vehicle	0	0	0	0
Inactivated PR8 virus	25 ± 47	19 ± 26	37 ± 39	43 ± 30
mRNA vaccine	25 ± 17	203 ± 138	379 ± 178	363 ± 154

Values are mean± SD (standard deviation)

[1061] These data demonstrate that mice vaccinated with mRNA vaccine at a dose of 0.4 mg/kg ID or inactivated PR8 virus showed 100% survival after challenge with influenza A/PR/8/34 virus in weeks 1, 2, 3, or 4 indicating that the vaccine conveyed protection against infection with influenza A/PR/8/34 virus under the conditions of this study. HAI activity remained low in weeks 1 through 4 for animals vaccinated with inactivated PR8 virus. Mice vaccinated with mRNA showed low mean HAI titers in week 1 after challenge, but increased titers in weeks 2, 3 and 4, respectively.

Example 24: Evaluation of H1 and H7-specific T cell responses in Influenza vaccine

[1062] Mice were immunized ID at week 0 and 3 with mRNA vaccine encoding the HA protein of influenza H1N1 or H7N9 virus and splenocytes were collected in week 5. T cells were stimulated for 16-18 hours with an HA peptide library (15mers of a full length protein)

that reproduce the H1 or H7 sequence. Non-specific stimulation of T cells was performed using PMA + ionomycin. IFNy ELISpot was used to identify staining for IFNγ. B cells were stimulated with either a peptide library (15mers of a full length protein) that reproduce the H1 or H7 sequence. Polyclonal B cell stimulation was performed using R848 + rIL-2. B cell ELISpot was used to identify staining for antigen-specific IgG.

[1063] IFNy ELISpot results demonstrated increased IFNy secretion by T cells after stimulation with H1 or H7 peptides (Figures 14A and 14B). No increased IFNy secretion was observed in T cells stimulated with PMA + ionomycin (Figure 14C). The results demonstrated that T cell responses are antigenspecific.

[1064] B cell ELISpot results demonstrated IgG secretion upon stimulation of B cells with full antigen, but not with H11H7 peptides (Figures 15A-15D). The lack of stimulation with the peptide library was to be expected because an IgG response is stimulated via MHCII antigens, which tend to be longer than MHCI antigen needed for stimulation of a T cell response.

Example 25: Evaluation of H1, H7, and H10-Specific T and B Cell Response Against Influenza Vaccines in Mouse Splenocytes

[1065] Influenza-specific T and B cell responses were evaluated following administration of mRNA vaccines (NAVs) encoding the H1, H10, and H7 hemagglutinin proteins. Briefly, 36 groups of 5 female 6-8 week old BALBIc mice were vaccinated with various doses via the intradermal route (ID) on day 0. Seven-week post vaccination, nimals were sacrificed and spleens were harvested. The splenocytes were assessed by IFNy ELISpot; Intracellular Cytokine Staining (ICS) for CD3, CD4, CD8, CD45, CCR7, CD44, CD25, IL-2 IFNγ, and TNFα markers, and by B-cell ELISpot.

IFNy ELISpot Analyses

[1066] Splenocytes from groups of mice that had been administered H10N8/N1-methyl pseudouridine /CO were assessed for H1, H7 and H10-specific IFNy production by IFNy ELISpot. Naive mice did not yield measurable IFNy cytokine response, whereas H10 peptide-specific IFNy ELISpot responses were detected in mice vaccinated intradermally with H10N8/N1-methyl pseudouridine/C0. The magnitude of these responses was found to be dependent on the dose of the vaccine administered. Peptide-stimulated splenocytes did not yield detectable H1- or H7-specific IFNy ELISpot responses. All groups produced IFNy ELISpot responses following stimulation with PMA+ionomycin (positive control).

[1067] H1 peptide-specific IFNy ELISpot responses were detected from splenocytes isolated from mice that had been administered H1N1/G2/C1, H1N1/G2/C0 (MC3 formulation), or H1N1/G2/C0 (KC2 formulation) intradermally, as well as from splenocytes from mice that had been administered H1N1/G2/C0 (MC3 formulation) intramuscularly.

[1068] H7 peptide-specific IFNy ELISpot responses were detected from splenocytes isolated from mice that had been administered H7N9/G2/C0 either intradermally or intramuscularly.

[1069] Splenocytes from control mice that had received H1N1/G2/C1/PBS showed no response to peptide stimulation.

B cell ELISpot Analyses

[1070] Briefly, H1-specific IgG and IgM responses were detected in several groups of mice vaccinated with different H1 vaccine formulations. H7-and H10-specific IgG and IgM responses were also elicited in the H7N9/N1-methyl pseudouridine /C0- and H10N8/N1-methyl pseudouridine /C0-vaccinated mice, respectively.

Intracellular Cytokine Staining Results

[1071] Intracellular Cytokine Staining and flow cytometric analysis indicated that H1-specific Th1 (IFNγ, TNFα, IL-2) cytokine responses were induced at both 16-18 hrs and 48 hrs in CD4+ T cells and CD8+ T cells following stimulation of splenocytes from H1-vaccinated mice with corresponding peptide or protein.

[1072] Additionally, H7- and H-10 peptide and protein-specific Th1 (IFNγ, TNFα, IL-2) responses were also detected in CD4+ and CD8+ T cells. H7-specific responses were noted in groups vaccinated with H7 vaccine formulations, and H10-specific responses were seen in groups vaccinated with H10 vaccine formulations

Example 26: Evaluation of H7- and H10-Specific T and B Cell Response Against Influenza Vaccines in Mouse Splenocytes - Time Course

[1073] Influenza-specific T and B cell responses were evaluated following administration of mRNA vaccines (NAVs) encoding the H10 and H7 hemagglutinin proteins. Briefly, 27 groups of 5 female 6-8 week old BALBIc mice (135 mice total) were vaccinated with various doses via the intradermal route (ID) on day 0 shown in Table 42. On days 7, 21, and 84 groups of 5 animals were sacrificed and blood samples and spleens were harvested.

[1074] The mouse spleens were analyzed using IFNγ ELISPOT, IgG ELISPOT (for B cell responses), and Intracellular Cytokine Staining (ICS). Serum samples were analyzed for inhibition of hemagglutin (HAI) by first treating with Receptor Destroying Enzyme (RDE) to inactivate non-specific inhibitors of hemagglutination (false positives) present in the sera. Next, RDE-treated sera were adsorbed using red blood cells (RBCs) to remove non-specific agglutinin (false negatives). Serially-diluted treated sera (2-fold dilutions) were pre-incubated with a standardized quantity of recombinant influenza antigen before RBCs were added to the mixture. Inhibition of hemagglutination indicates the presence of HA-specific antibodies. Pooled anti-H10 mouse serum was used as a positive control and showed HAI titers within range of previously-observed data.

[1075] After receiving a single dose of the mRNA NAVs on day 0, HAI titers were not detected at day 7 (Table 42) but were detected at day 21 (Table 43) and continued to increase through day 84 (Table 44). Day 84 titers were 2-fold to 15-fold higher than those from day 21 (Figures 15 and 16).

[1076] Administration of the H10N8/ N1-methyl pseudouridine IC1 formulation MC3 resulted in higher HAI titers as compared to the HAI titers induced by the H10N8/N1-methyl pseudouridine/C0 formulation MC3 (Figure 17). The MC3 LNP formulation consisted of a cationic lipid (DLin-MC3-DMA, 50 mol%), non-cationic lipid (DSPC, 10 mol%), PEG lipid (PEG-DOMG 1.5 mol%) and a structural lipid (cholesterol, 38.5 mol%).

[1077] Administration of the H10N8/ N1-methyl pseudouridine IC1 formulation MC3 at 10 pg/dose achieved the highest HAI titers, whereas the HAI titers following administration of the 2.0 pg/dose and 0.4 pg/dose are similar to each other (Figure 16).

Table 42: HAI Titers against H10 and H7 at Day 7

mRNA vaccine once at day 0 via ID injection	Dose per mouse (µg)	Group (N=5)	HAI titers of mouse sera Day 0 (11-10-2014) = vaccination day 0 Day 7 (11-17-2014) = terminal bleeds groups 1-9 M1	M2	M3	M4	M5	Mean	Stdev	
H10N8/N1-methyl pseudouridine /C1, formulation MC3	**10**	**1**	<10	<10	<10	<10	<10	< **10**	N/A	HAI titers against H10 protein
	2	**2**	<10	<10	<10	<10	<10	< **10**	N/A	
	0.4	**3**	<10	<10	<10	<10	<10	< **10**	N/A	
H10N8/N1-methyl pseudouridine /C0, formulation MC3	**10**	**4**	<10	<10	<10	<10	<10	< **10**	N/A	
	2	**5**	<10	<10	<10	<10	<10	< **10**	N/A	
	0.4	**6**	<10	<10	<10	<10	<10	< **10**	N/A	
PBS	**0**	**7**	<10	<10	<10	<10	<10	< **10**	N/A	
PBS	**0**	**7**	<10	<10	<10	<10	<10	< **10**	N/A	HAI titers against H7 protein
H7N9/C0	**10**	**8**	<10	<10	<10	<10	<10	< **10**	N/A	
H7N9/C1	**10**	**9**	<10	<10	<10	<10	<10	< **10**	N/A	

Table 43: HAI Titers against H10 and H7 at Day 21

mRNAna vaccine once at day 0 via ID injection	Dose per mouse (µg)	Group (N=5)	HAI titers of mouse sera Day 0 (11-10-2014) = vaccination day 0 Day 21 (12-01-2014) = terminal bleeds group 10-18 M1	M2	M3	M4	M5	Mean	Stdev	
H10N8/N1-methyl pseudouridine/C1, formulation MC3	10	10	160	320	640	80	160	272	223	HAI titers against H10 protein
	2	11	40	80	40	80	160	80	49	
	0.4	12	80	160	40	80	80	88	44	
H10N8/N1-methyl pseudouridine/C0, formulation MC3	10	13	160	40	80	20	40	68	56	
	2	14	20	20	10	10	20	16	5	
	0.4	15	20	10	40	20	80	34	28	
PBS	0	16	<10	<10	<10	<10	<10	< 10	N/A	
PBS	0	16	<10	<10	<10	<10	<10	< 10	0	HAI titers against H7 protein
H7N9/C0	10	17	40	10	10	20	20	26	13	
H7N9C1	10	18	160	160	160	320	80	176	88	

Table 44: HAI Titers against H10 and H7 at Day 84

mRNA vaccine once at day 0 via ID injection	Dose per mouse (µg)	Group (N=5)	HAI titers of mouse sera Day 0 (11-10-2014) = vaccination day 0 Day 84 (02-02-2015) = terminal bleeds groups 19-27								
			M1	M2	M3	M4	M5		Mean	Stdev	
H10N8/N1-methyl pseudouridine /C1, formulation MC3	**10**	**19**	160	320	320	1280	640		**544**	447	HAI titers against H10 protein
	2	**20**	320	320	640	160	320		**352**	175	
	0.4	**21**	640	320	160	320	640		**416**	215	
H10N8/N1-methyl pseudouridine /C0, formulation MC3	**10**	**22**	160	320	320	320	320		**288**	72	
	2	**23**	320	320	160	80	320		**240**	113	
	0.4	**24**	40	320	320	320	80		**216**	143	
PBS	**0**	**25**	<10	<10	<10	<10	<10	**<**	**10**	N/A	
PBS	**0**	**25**	<10	<10	<10	<10	<10	**<**	**10**	N/A	HAI titers against H7 protein
H7N9/C0	**10**	**26**	20	40	80	20	<10	**≤**	**40**	28	
H7N9/C1	**10**	**27**	640	320	320	160	80		**304**	215	

Example 27: Anti-H10 Hemagglutinin Analysis in Non-Human Primates - Dose Range Study

[1078] Groups of cynomoglus monkeys were vaccinated with various doses of NAV encoding H10 HA/LNP formulations (50 pg/dose, 200 pg/dose, 400 pg/dose), NAV encoding H10 HA/LNP delivered with 3M device, or control NAV encoding H10 HA/PBS. Serum samples were taken from the monkeys weekly and evaluated for inhibition of hemagglutinin (Table 45, Figure 19).

Table 45: HAI titers against H10 protein

HAI titers against H10 protein (A/Jiangxi-Donghu/346/2013)								
Cynomolgus monkey ID	Pre-treatment	Week 1	Week 2	Week 3	Week 4	Week 5	Week 6	Week 9

Group 1 (50 µg)	1001	10	40	20	20	320	1280	1280	5120
	1002	20	20	40	40	640	2560	2560	10240
	1003	<10	<10	40	80	640	2560	2560	10240
	Mean	**<13**	**<23**	**33**	**47**	**533**	**2133**	**2133**	**8533**
Group 2 (200 µg)	2001	<10	<10	10	40	640	5120	2560	10240
	2002	20	20	20	40	640	2560	2560	10240
	2003	<10	40	20	40	320	5120	5120	10240
	Mean	**<13**	**<23**	**17**	**40**	**533**	**4267**	**3413**	**10240**
Group 3 (400 µg)	3001	20	<10	40	80	1280	5120	10240	20480
	3002	<10	40	40	160	640	5120	5120	20480
	3003	<10	20	80	160	1280	10240	20480	40960
	Mean	**<13**	**<23**	**53**	**133**	**1067**	**6827**	**11947**	**27307**
Group 4 (vaccine device)	4001	<10	40	<10	20	40	80	80	160
	4002	<10	40	<10	<10	40	160	160	160
	4003	20	40	<10	10	40	80	80	160
	Mean	**<13**	**40**	**<10**	**<13**	**40**	**107**	**107**	**160**
Group 5 (PBS)	5001	<10	20	80	320	1280	5120	5120	10240
	5002	<10	20	40	160	1280	5120	5120	10240
	5003	<10	40	40	80	1280	5120	5120	20480
	Mean	**<10**	**27**	**53**	**187**	**1280**	**5120**	**5120**	**13653**
		Assay Positive Control: Anti-H10 Pool of 9 mouse sera M1, M2, M3, of Group 1, N1-methyl pseudouridine ($m^1\Psi$) and 5-methyl cytidine, Group 3 of study day 14 HAI titer = 160				Assay Positive Control: Anti H-10 Pool of 9 mouse sera M1, M2, M3 of Group 1, N1-methyl pseudouridine ($m^1\Psi$) and 5-methyl cytidine, Group 3 of study day 14 HAI titer = 80			

Example 28: Determination of Time to Onset of Immunity using an H7N9 Vaccine in a Ferret Model

[1079] The time to onset of immunity (as measured by antibody titers and reduction in viral titers following challenge) of an influenza AIAnhui1112013 (H7N9) vaccine was evaluated in a ferrets model of influenza infection. Briefly, twenty groups of 8 ferrets each were vaccinated on day 0 with the H7N9 mRNA NAV vaccine at a dose of 10 pg, 50 pg, or 200 pg, a vH7N9 vaccine lacking the 14 kDa cap at a dose of 200 pg, or PBS control via the intradermal route. Five of the 20 groups received a second vaccination (a booster) to determine if a second dose increased protection. The animals were then exposed to influenza AIAnhui1112013 (H7N9) virus via the intranasal route. The study design is shown in Table 45:

Table 45. Study Design

Group No.	Vaccine Dose	Day of Vaccination	No. of Animals	Day of Challenge	Day of Blood Draws
1	H7N9/N1-methyl pseudouridine /C1/MC3, 200 µg	0	8	7	0, 7
2	H7N9/N1-methyl pseudouridine /C1/MC3, 50 µg	0	8	7	0, 7
3	H7N9/N1-methyl pseudouridine /C1/MC3, 10 µg	0	8	7	0, 7
4	PBS	0	8	7	0, 7
5	H7N9/MC3 (~14 kDa Cap) 200 µg	0	8	7	0 ,7
6	H7N9/N1-methyl pseudouridine /C1/MC3, 200 µg	0	8	21	0, 21
7	H7N9/N1-methyl pseudouridine /C1/MC3, 50 µg	0	8	21	0, 21
8	H7N9/N1-methyl pseudouridine /C1/MC3, 10 µg	0	8	21	0, 21
9	PBS	0	8	21	0, 21
10	H7N9/MC3 (~14 kDa Cap) 200 µg	0	8	21	0, 21
11	H7N9/N1-methyl pseudouridine /C1/MC3, 200 µg	0	8	49	0, 49
12	H7N9/N1-methyl pseudouridine /C1/MC3, 50 µg	0	8	49	0, 49
13	H7N9/N1-methyl pseudouridine /C1/MC3, 10 µg	0	8	49	0, 49
14	PBS	0	8	49	0, 49
15	H7N9/MC3 (~14 kDa Cap) 200 µg	0	8	49	0, 49
16	H7N9/N1-methyl pseudouridine /C1/MC3, 200 µg	4, 25	8	53	4, 25, 53
17	H7N9/N1-methyl pseudouridine /C1/MC3, 50 µg	4, 25	8	53	4, 25, 53
18	H7N9/N1-methyl pseudouridine /C1/MC3, 10 µg	4, 25	8	53	4, 25, 53
19	PBS	4, 25	8	53	4, 25, 53
20	H7N9/MC3 (~14 kDa Cap) 200 µg	4, 25	8	53	4, 25, 53

[1080] Because H7N9 does not induce a lethal disease in ferrets, the primary endpoint was viral burden in tissues (primarily lung) 3 days post-challenge, as determined by TCID50 (50% tissue culture infective dose). In addition to viral titers, blood was collected prior to vaccination and immediately prior to challenge to determine influenza-specific antibody titers, as determined by hemagglutination inhibition (HAI) and microneutralization (MN) assays.

[1081] The lung homogenate data show that a single vaccination at any concentration resulted in a reduction in viral titers, with a time to onset of immunity before 7 days post vaccination (Table 46). There was a 1 log reduction in all vaccine groups challenged 7 days post vaccination. This improved for groups challenged 21 days post vaccination in a dose-dependent fashion to 3, 2, and 1 log reduction for the 200-, 50-, and 10-pg groups, respectively (Figure 20). This further improved for groups challenged 49 days post vaccination to 4, 2, and 3 log reduction for the 200 , 50-, and 10-pg groups, respectively. A statistically significant difference was observed relative to PBS control group at 7 days post vaccination in the 10- and 200-pg groups ($p < 0.05$). The 50-pg dose group also tended towards reduced lung viral titers, but only had statistically significant differences from the PBS group when a boost was administered. This was likely due to the occasional outlying high viral titer sample, which increased the variability in this group.

Table 46: Virus Burden (TCID50 Group Geographic Means)

	Nasal Wash			Lung Homogenate		
	Day 7	Day 21	Day 49	Day 7	Day 21	Day 49
H7N9/N1-methyl pseudouridine /C1/MC3 200 µg	3.7×10^3	3.1×10^3	3.3×10^4	3.0×10^3	4.6×10^1	8.0×10^0
H7N9/N1-methyl pseudouridine /C1/MC3 50 µg	2.4×10^3	1.6×10^4	6.7×10^4	2.0×10^3	5.9×10^2	9.7×10^2
H7N9/N1-methyl pseudouridine /C1/MC3 10 µg	3.7×10^3	8.1×10^2	1.0×10^5	2.3×10^3	2.9×10^3	6.2×10^1
PBS	6.0×10^3	2.9×10^3	1.7×10^5	3.8×10^4	3.2×10^4	2.1×10^4
H7N9/MC3 (~14 kDa Cap) 200 µg	1.4×10^4	1.5×10^3	3.6×10^5	7.9×10^3	1.6×10^3	4.8×10^2

† 158 is the lower limit of detection of the nasal wash $TCID_{50}$ assay. All below LLOQ values were assigned a value of 79 for producing the geomean values for nasal wash. 15.8 is the lower limit of detection of the lung homogenate $TCID_{50}$ assay. All below LLOQ values were assigned a value of 8 for producing the geomean values for nasal wash.

[1082] There was no statistical benefit observed from the boosting vaccination compared to a single vaccination, but this is likely due to the 2-4 log reduction in viral lung titers by 49 days seen in both the single vaccination and 2 (boost) vaccination groups (Table 47).

Table 47: Lung Virus Burden (TCID50 Group Geographic Means)- Boosted vs. Unboosted

	Day 49 Unboosted	Day 49 with the Day 21 Boost
200 µg Vaccine	8.0×10^0	8.0×10^0
50 µg Vaccine	9.7×10^2	8.0×10^0
10 µg Vaccine	6.2×10^1	1.1×10^2
PBS	2.1×10^4	2.2×10^4
Empty Vaccine Vector	4.8×10^2	8.0×10^0

† 15.8 is the lower limit of detection of the lung homogenate $TCID_{50}$ assay. All below LLOQ values were assigned a value of 8 for producing the geomean values for nasal wash.

[1083] Serum was collected from each animal immediately before vaccination and challenge and analyzed by HAI assay. As seen from Table 48, a dose dependent increase was visible 7 days post vaccination, and titers increased over time. The 200-pg to 14-kDa cap vaccine did provide some titer above background, between the 10-pg vaccine and PBS response. The groups receiving a second vaccination showed a dose-dependent, 3 8 fold increase in antibody titers following the second vaccination. The 200-pg to 14-kDa cap vaccine also showed an approximately 2-fold increase in antibody titers following the second vaccination. In all cases, a statistically significant increase relative to the pre-vaccination blood sample was first observed 21 days post-vaccination ($p < 0.05$).

Table 48: HAI Antibody Titer Results

		Geomean			
Cohort	Treatment	Day 0	Day 7	Day 21	Day 49
1	200 µg Vaccine	5.00†	5.45	N/A‡	N/A
1	50 µg Vaccine	5.00	6.48	N/A	N/A
1	10 µg Vaccine	5.00	5.94	N/A	N/A
1	PBS	5.00	5.00	N/A	N/A
1	Empty Vaccine Vector	5.00	5.45	N/A	N/A
2	200 µg Vaccine	5.00	N/A	43.62	N/A
2	50 µg Vaccine	5.00	N/A	25.94	N/A
2	10 µg Vaccine	5.00	N/A	25.94	N/A
2	PBS	5.00	N/A	5.00	N/A
2	Empty Vaccine Vector	5.00	N/A	9.35	N/A
3	200 µg Vaccine	5.00	N/A	N/A	61.69
3	50 µg Vaccine	5.00	N/A	N/A	33.64
3	10 µg Vaccine	5.00	N/A	N/A	25.94
3	PBS	5.00	N/A	N/A	5.00
3	Empty Vaccine Vector	5.00	N/A	N/A	18.34
4	200 µg Vaccine	5.00	N/A	59.93	480.67
4	50 µg Vaccine	5.00	N/A	29.97	195.85
4	10 µg Vaccine	5.00	N/A	40.00	97.92
4	PBS	5.00	N/A	5.00	5.45
4	Empty Vaccine Vector	5.94	N/A	5.96	36.68

† Ten is the lower limit of detection of the HAI assay. All <10 were assigned a value of 5 for producing the geomean values.

‡ NA = Not applicable.

[1084] Serum samples were also analyzed by MN assay. As shown in Table 49, increases in MN titer was observed as early as 7 days post vaccination and continued to increase through 49 days post vaccination. The titers were highest in the 200-µg vaccine group, and were similar in the 50- and 10-pg groups. For the 50- and 200-pg vaccine dose groups, a statistically significant increase was observed by 7 days post-vaccination ($p < 0.05$). For the 10-µg vaccine dose groups, the significant increase was not observed until 21 days post-vaccination ($p < 0.05$). The 200-pg to 14-kDa cap vaccine did provide some titer above background, which was statistically significant by 49 days post vaccination ($p < 0.05$) and were lower than the titers for the 50- and 10-pg groups. The groups receiving a second vaccination experienced a 3.0-4.5 fold increase in titers following the second vaccination. The 200-pg to 14-kDa cap vaccine also showed an approximately 1.5-fold increase in titers following the second vaccination.

Table 49: MN Antibody Titer Results

		Geomean†			
Cohort	Treatment	Day 0	Day 7	Day 21	Day 49
1	H7N9/N1-methyl pseudouridine /C1/MC3 200 µg Vaccine	15.00	33.67	N/A‡	N/A
1	H7N9/N1-methyl pseudouridine /C1/MC3 50 µg Vaccine	17.38	42.29	N/A	N/A
1	H7N9/N1-methyl pseudouridine /C1/MC3 10 µg Vaccine	21.21	25.94	N/A	N/A
1	PBS	19.58	18.47	N/A	N/A
1	H7N9/MC3 (~14 kDa Cap) 200 ug	17.43	20.03	N/A	N/A
2	H7N9/N1-methyl pseudouridine /C1/MC3 200 µg Vaccine	14.69	N/A	92.41	N/A
2	H7N9/N1-methyl pseudouridine /C1/MC3 50 µg Vaccine	17.49	N/A	42.50	N/A
2	H7N9/N1-methyl pseudouridine /C1/MC3 10 µg Vaccine	15.07	N/A	43.48	N/A
2	PBS	17.89	N/A	13.60	N/A
2	H7N9/MC3 (~14 kDa Cap) 200 µg	18.52	N/A	17.43	N/A
3	H7N9/N1-methyl pseudouridine /C1/MC3 200 µg Vaccine	15.51	N/A	N/A	160.08
3	H7N9/N1-methyl pseudouridine /C1/MC3 50 µg Vaccine	13.16	N/A	N/A	89.80
3	H7N9/N1-methyl pseudouridine /C1/MC3 10 µg Vaccine	13.12	N/A	N/A	77.51
3	PBS	12.70	N/A	N/A	10.68
3	H7N9/MC3 (~14 kDa Cap) 200 µg	10.33	N/A	N/A	46.21
4	H7N9/N1-methyl pseudouridine /C1/MC3 200 µg Vaccine	10.33	N/A	63.44	697.88
4	H7N9/N1-methyl pseudouridine /C1/MC3 50 µg Vaccine	10.68	N/A	29.99	273.04
4	H7N9/N1-methyl pseudouridine /C1/MC3 10 µg Vaccine	10.68	N/A	31.97	232.65
4	PBS	12.01	N/A	11.03	10.68
4	H7N9/MC3 (~14 kDa Cap) 200 µg	11.78	N/A	16.47	69.06

† 20 is the lower limit of detection of the MN assay. All < 20 were assigned a value of 10 for producing the geomean values.

‡ N/A = Not applicable.

[1085] Quite surprisingly, the vaccine constructs of the invention reduced viral titers in the lungs when exposed to virus just 7 days following vaccination. Statistically significant increases in antibody titer as measured by HAI and MN were detected as early as 7 days following vaccination. A second vaccination (i.e., booster) did increase antibody titers, but did not statistically reduce the viral titer, as a single vaccination eliminated all virus in most animals. The -14 kDa cap vaccine at 200 pg/animal provided less protection than 10-pg full vaccine, but did reduce viral burden in the lung and increased antibody titers, both relative to PBS control.

Patentkrav

1. Nukleinsyrevaccine, der omfatter:
ét eller flere messenger-RNA- (mRNA) polynukleotider med en åben læseramme, der koder for mindst ét viralt antigent polypeptid,
hvor det ene eller flere mRNA-polynkcleotider omfatter mindst én kemisk modifikation, hvor den mindst ene kemiske modifikation vælges fra gruppen bestående af pseudouridin, N1-methylpseudouridin, 2-thiouridin, 4-thiouridin, 5-methylcytosin, 2-thio-1-methyl-1-deaza-pseudouridin, 2-thio-1-methyl-pseudouridin, 2-thio-5-aza-uridin, 2-thio-dihydropseudouridin, 2-thio-dihydrouridin, 2-thio-pseudouridin, 4-methoxy-2-thio-pseudouridin, 4-methoxy-pseudouridin, 4-thio-1-methyl-pseudouridin, 4-thio-pseudouridin, 5-aza-uridin, dihydropseudouridin, 5-methoxyuridin, og 2'-O-methyluridin; og
hvor det eller flere mRNA-polynukleotider er formuleret i en kationisk lipidnanopartikel med et molforhold på 20-60 % kationisk lipid, 0,5-15 % PEG-modificeret lipid, 25-55 % sterol og 5-25 % ikke-kationisk lipid.

2. Vaccine ifølge krav 1, hvori (a) nukleinsyrevaccinen er multivalent, og/eller (b) den åbne læseramme for det ene eller flere mRNA-polynukleotider koder for mindst 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, 20 eller 50 antigene polypeptider.

3. Vaccine ifølge krav 1 eller krav 2, hvor mindst 80 %, *f.eks.* 100 %, uracil i den åbne læseramme har en kemisk modifikation.

4. Vaccine ifølge krav 3, hvor den kemiske modifikation er i 5-stillingen af uracil.

5. Vaccine ifølge krav 3 eller krav 4, hvor den kemiske modifikation er et N1-methyl-pseudouridin.

6. Vaccine ifølge et hvilket som helst af krav 1-5, hvor det kationiske lipid er et ioniserbart kationisk lipid, det ikke-kationiske lipid er et neutralt lipid, og sterolen er en cholesterol.

7. Vaccine ifølge et hvilket som helst af kravene 1-6, hvor det virale antigene

polypeptid stammer fra et virus valgt blandt adenovirus; herpes simplex, type 1; herpes simplex, type 2; encephalitisvirus, papillomavirus, varicella-zoster-virus; Epstein-barr-virus; human cytomegalovirus; human herpesvirus, type 8; human papillomavirus; BK-virus; JC-virus; kobbevirus; poliovirus, hepatitis B-virus; human bocavirus; parvovirus B19; human astrovirus; Norwalk virus; coxsackievirus; hepatitis A-virus; rhinovirus; svær, akut respiratorisk syndromvirus; hepatitis C-virus; gul febervirus; Dengue-virus; vestnilvirus; rubellavirus; hepatitis E-virus; human immunodeficiensvirus (HIV); influenzavirus, type A eller B; Guanarito-virus; Junin-virus; lassavirus; Machupo-virus; Sabiá-virus; Krim-Congo hæmorrhagisk febervirus; ebolavirus; Marburgvirus; mæslingevirus; fåresygevirus; parainfluenzavirus; respiratorisk syncytialvirus; human metapneumovirus; Hendra-virus; Nipah-virus; rabiesvirus; hepatitis D; rotavirus; orbivirus; coltivirus; human enterovirus; japansk encephalitisvirus; vesikulær exanthernavirus; østlig hesteencephalitis; hantavirus; Middle East Respiratory Coronavirus; chikungunya-virus eller Banna-virus.

8. Vaccine ifølge et hvilket som helst af kravene 1-7 til anvendelse som et terapeutisk eller profylaktisk middel.

9. Vaccine ifølge et hvilket som helst af kravene 1-7 til anvendelse i en fremgangsmåde til forebyggelse eller behandling af en virusinfektion i en menneske- eller dyrekrop.

10. Vaccine til anvendelse ifølge krav 9, hvor virusinfektionen er en luftvejsinfektion, eventuelt hvor luftvejsinfektionen er en adenovirusinfektion, fugleinfluenza (H5N1), influenza, RSV-infektion, svært, akut luftvejssyndrom (SARS), bihulebetændelse eller svineinfluenzainfektion (H1N1).

11. Vaccine til anvendelse ifølge et hvilket som helst af kravene 8-10, hvor vaccinen administreres i en enkelt administrationshændelse eller i to eller flere doser.

12. Vaccine til anvendelse ifølge et hvilket som helstaf kravene 8-11, hvor vaccinen administreres til individet ved intramuskulær (IM) eller intradermal (ID) injektion.

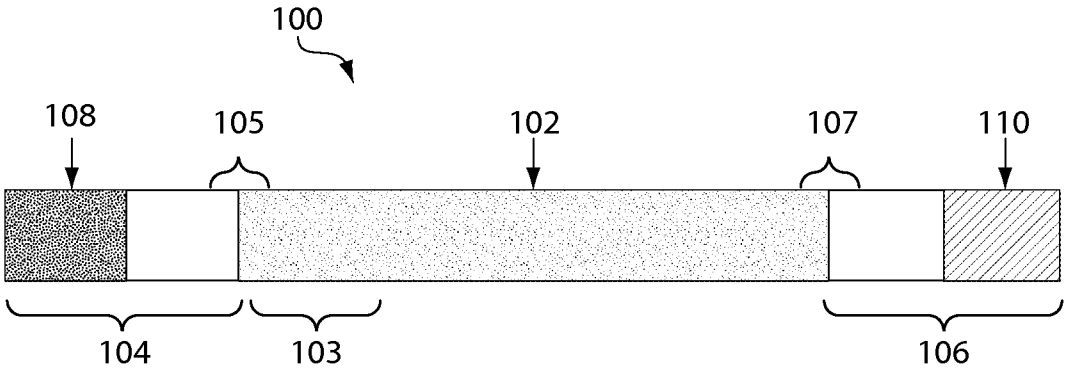


FIGURE 1A

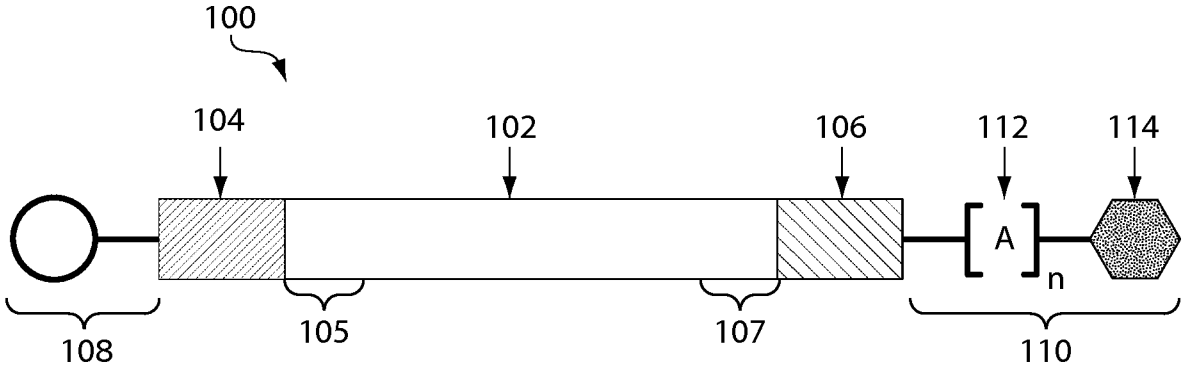


FIGURE 1B

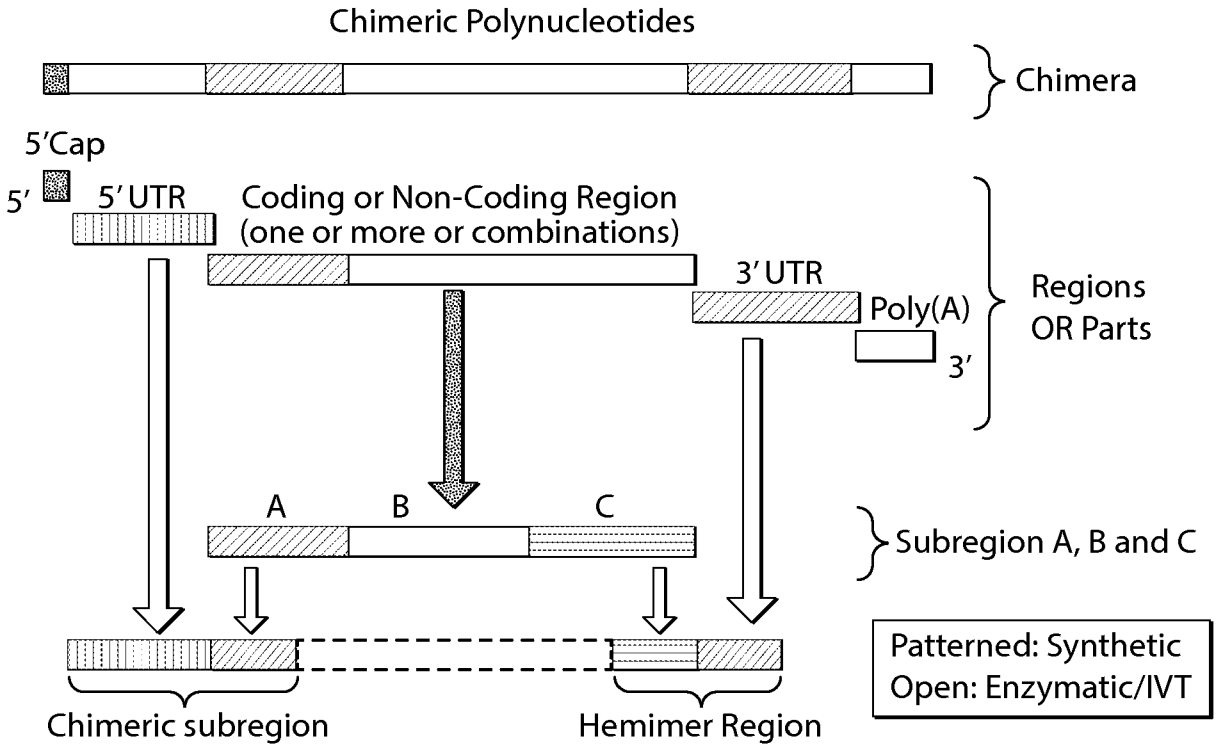

Chimeric Polynucleotides
Chimera
5′Cap
5′
5′ UTR
Coding or Non-Coding Region
(one or more or combinations)
3′ UTR
Poly(A)
3′
Regions
OR Parts
A
B
C
Subregion A, B and C
Patterned: Synthetic
Open: Enzymatic/IVT
Chimeric subregion
Hemimer Region

FIGURE 2

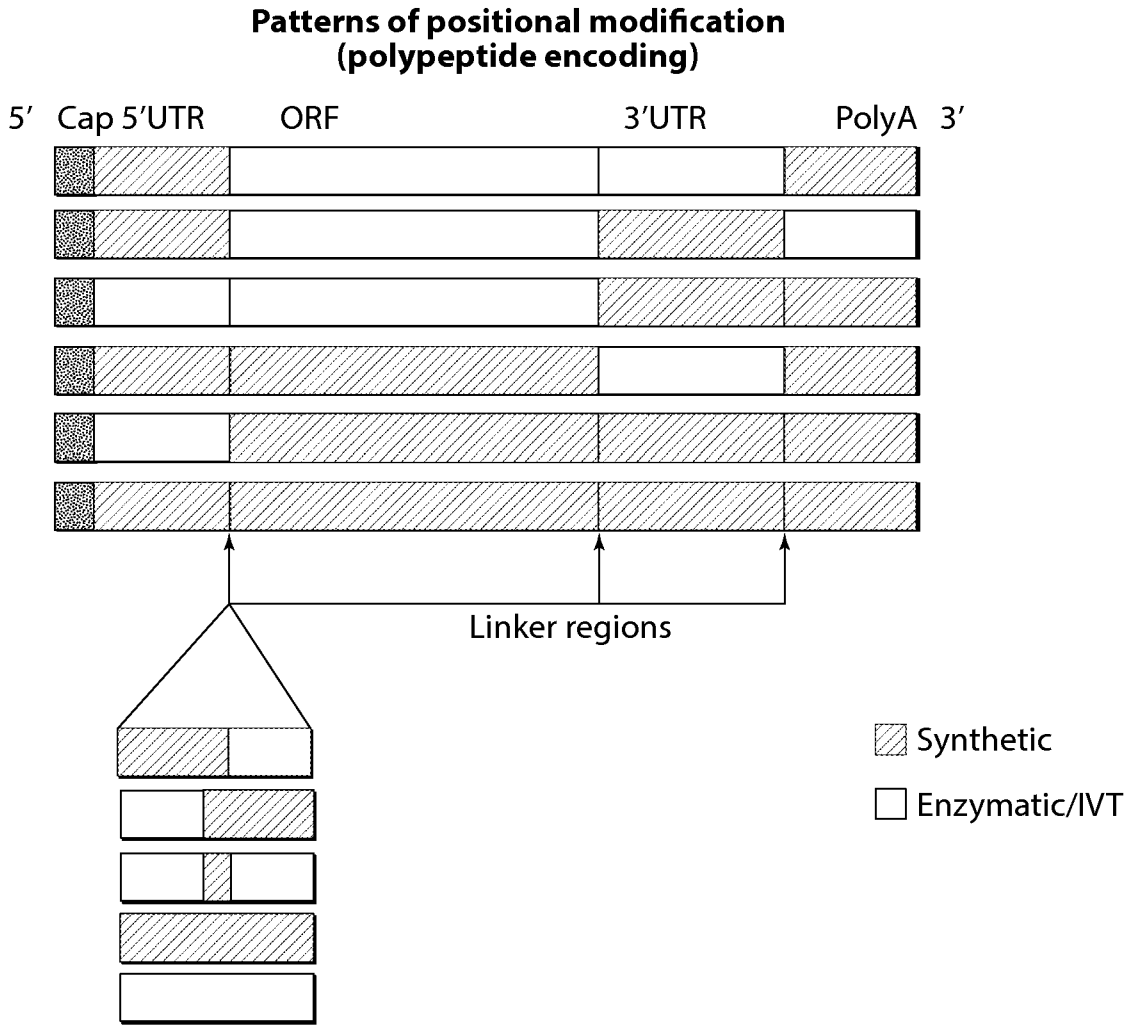

Patterns of positional modification
(polypeptide encoding)
5′
Cap 5′UTR
ORF
3′UTR
PolyA
3′
Linker regions
Synthetic
Enzymatic/IVT

FIGURE 3

Patterns of positional modification

A	(L)	B	(L)	C	(L)

5′[NNNNN$_n$]$_x$ L1 [NNNNNNNNNNNN$_o$]$_y$ L2 [XXXXXXXXXXXX$_p$]$_z$ L3 3′

5′[NNNNN$_n$]$_x$ L1 [XXXXXXXXXXXXXXX$_o$]$_y$ L2 [XXXXXXXXXXXX$_p$]$_z$ L3 3′

5′[XXXXXX$_n$]$_x$ L1 [XXXXXXXXXXXXXXX$_o$]$_y$ L2 [XXXXXXXXXXXX$_p$]$_z$ L3 3′

5′[NNNNN$_n$]$_x$ L1 [NNNNNNNNNNNN$_o$]$_y$ L2 [XXXXXXXXXXXX$_p$]$_z$ L3 3′

A, B, C- Region or Part of chimeric polynucleotide
N-nucleoside
n, o, p-number of nucleosides
x, y, z-number of regions
X-selective placement nucleoside
L-linker (optional)

FIGURE 4

Blocked or structured 3′ termini

A	(L)	B	(L)	C	(L)

5′[NNNNN$_n$]$_x$ L1 [NNNNNNNNNNNN$_o$]$_y$ L2 [XXXXXXXXXXXX$_p$]$_z$ L3 3′

5′[NNNNN$_n$]$_x$ L1 [NNNNNNNNNNNN$_o$]$_y$ L2 [XXXXXXXXXXXX$_p$]$_z$ K 3′

5′[NNNNN$_n$]$_x$ L1 [NNNNNNNNNNNN$_o$]$_y$ L2 [XXXXX**XX**XXXXXXXXX $^{X}_{X}$

3′ [XXXXXXXXXXX ^{X}

5′[NNNNN$_n$]$_x$ L1 [NNNNNNNNNNNN$_o$]$_y$ L2 [XXXX**XXX**XXXXXXXX $^{L}_{L}$ =L3

3′ XXXXXXXXXX $_{L}^{L}$

5′[NNNNN$_n$]$_x$ L1 [NNNNNNNNNNNN$_o$]$_y$ L2 [XXXX**XXX**XXXXXXXX $^{L}_{L}$ =L3

^{X} X XXXXXXXXXX $_{L}^{L}$

^{X} X XXXXXXXXXX 3′

A, B, C- Region or Part of chimeric polynucleotide
N-nucleoside
n, o, p-number of nucleosides
x, y, z-number of regions
X-selective placement nucleoside
K-non-nucleosidic moiety or conjugate
L-linker (optional)

FIGURE 5

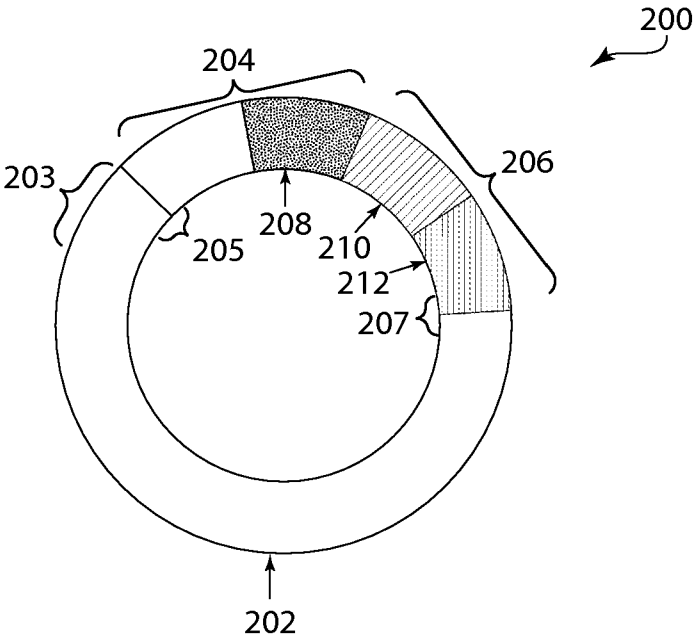


FIGURE 6A

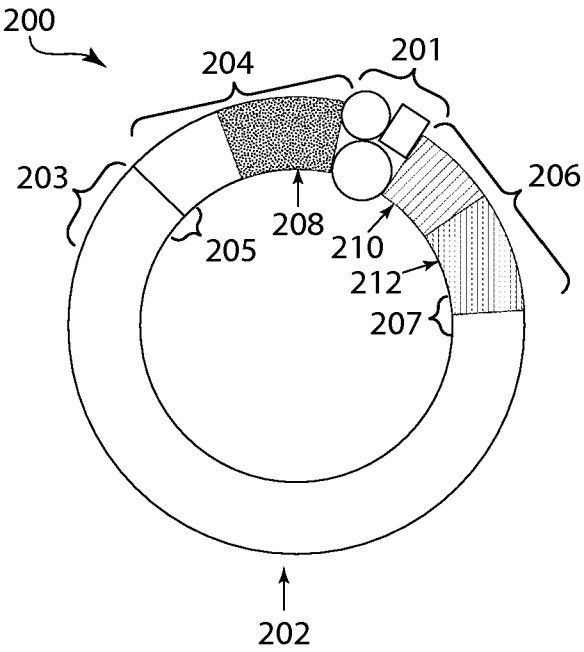


FIGURE 6B

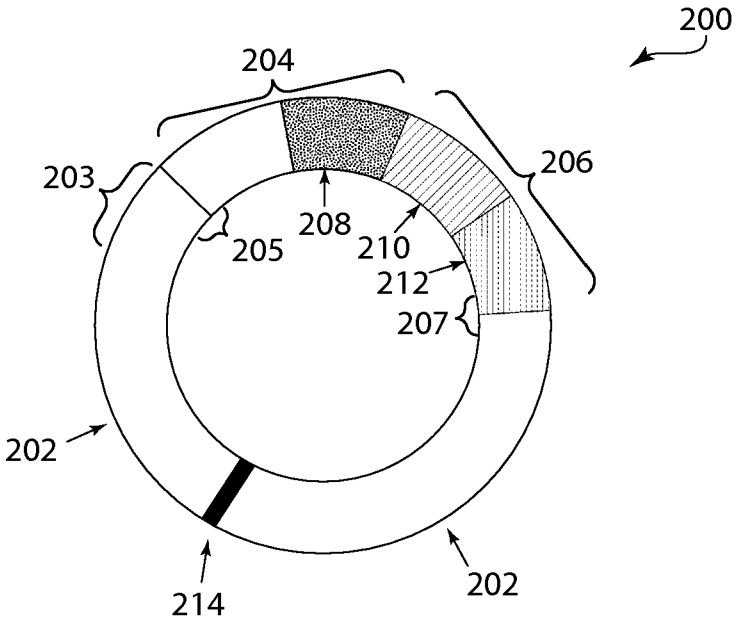


FIGURE 7A

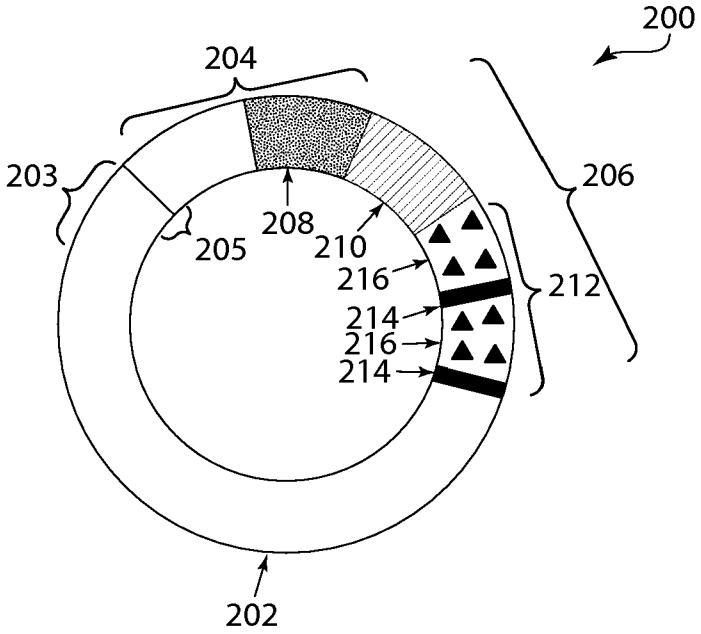


FIGURE 7B

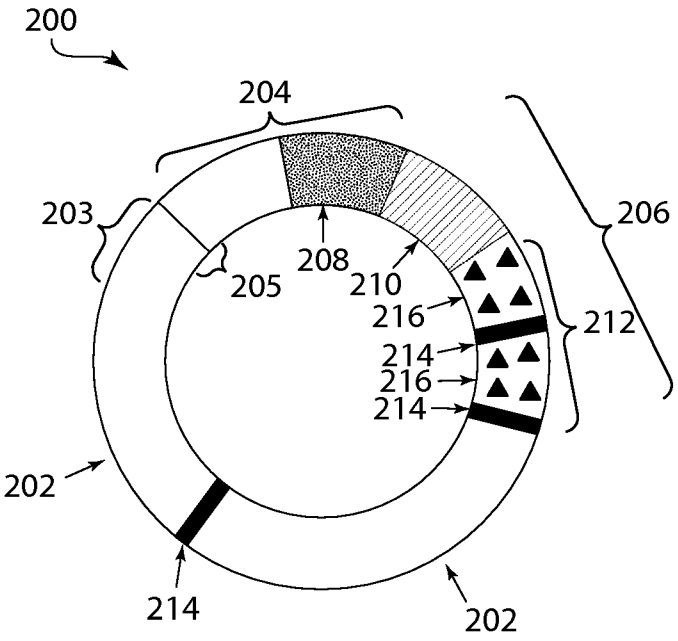


FIGURE 8A

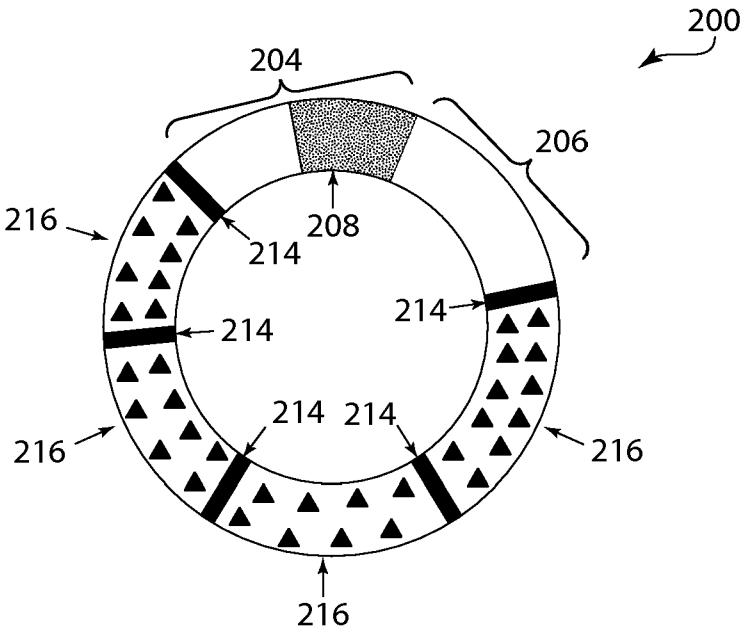


FIGURE 8B

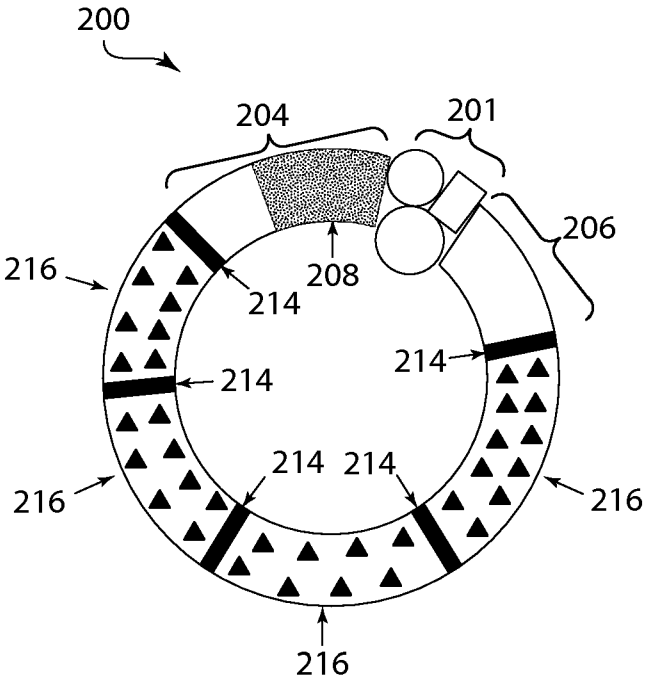


FIGURE 9

Dosing & Clinical assessment

	Control	Protein Vaccine	mRNA H1N1 IM	mRNA H1N1 ID	Positive control
mRNA dose	–	–	8-10µg	8-10µg	80µg
Health Score	7	2	1	2	2
Average weight loss	>25%	10%	<6%	8-10%	8-10%
Lethality	100%	0%	0%	0%	0%

HA neutralization titers

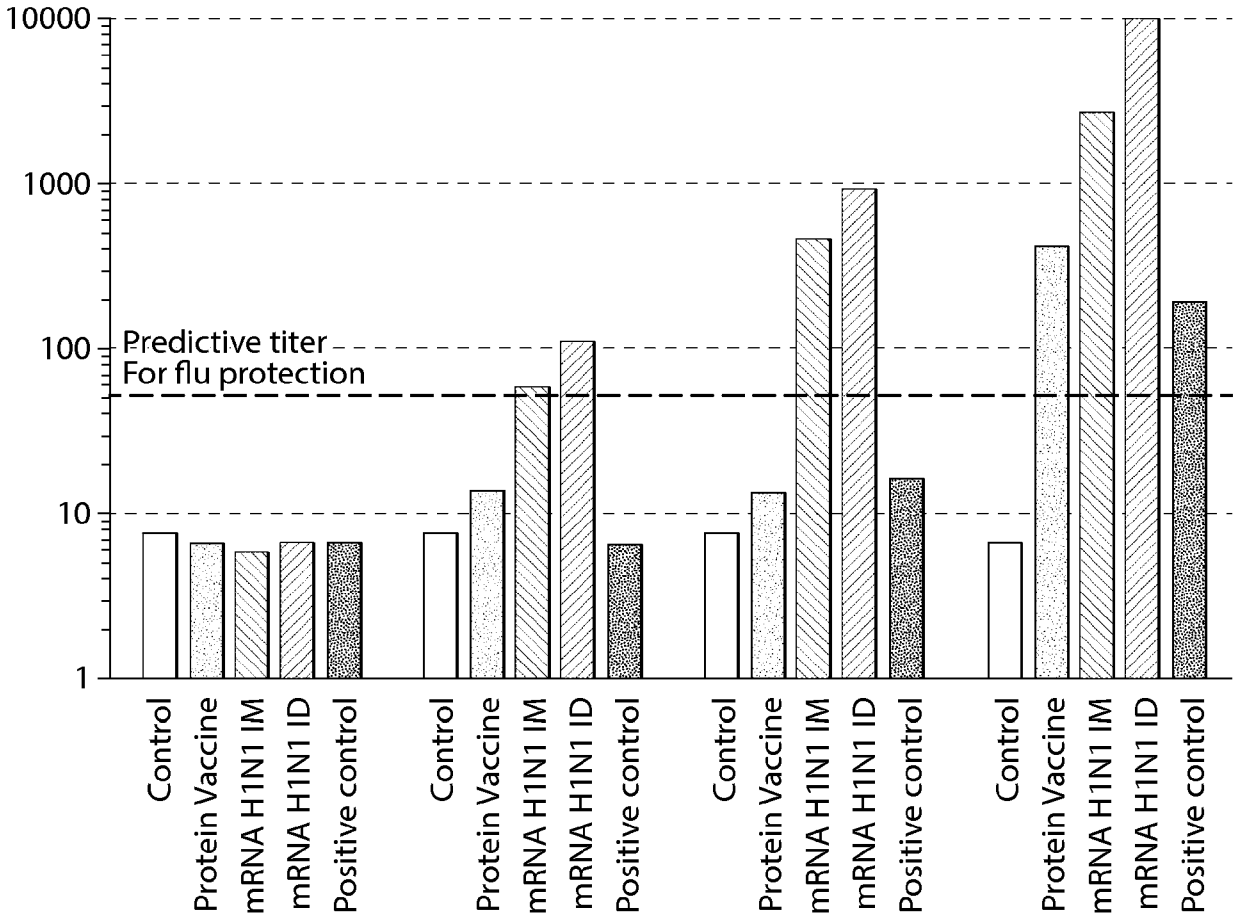


FIGURE 10

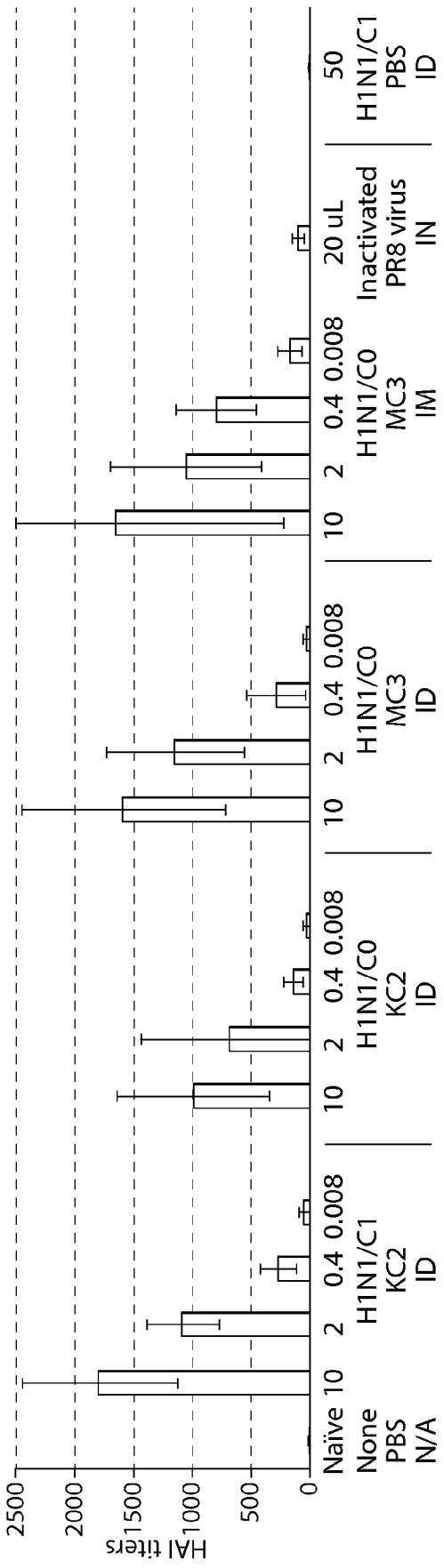

HAI titers
2500
2000
1500
1000
500
0
Naïve
None
PBS
N/A
10
2
0.4
0.008
H1N1/C1
KC2
ID
H1N1/C0
KC2
ID
H1N1/C0
MC3
ID
H1N1/C0
MC3
IM
20 uL
Inactivated
PR8 virus
IN
50
H1N1/C1
PBS
ID

FIGURE 11

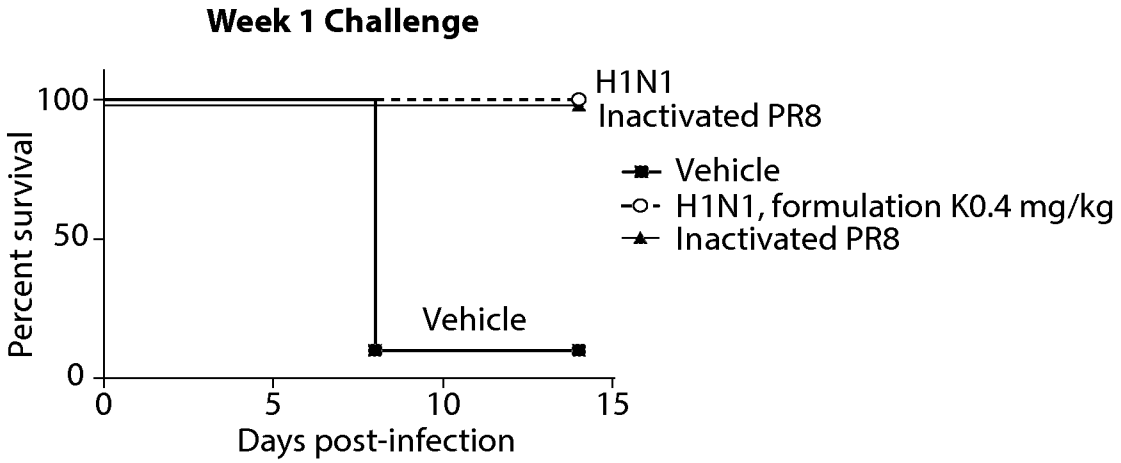


FIGURE 12A

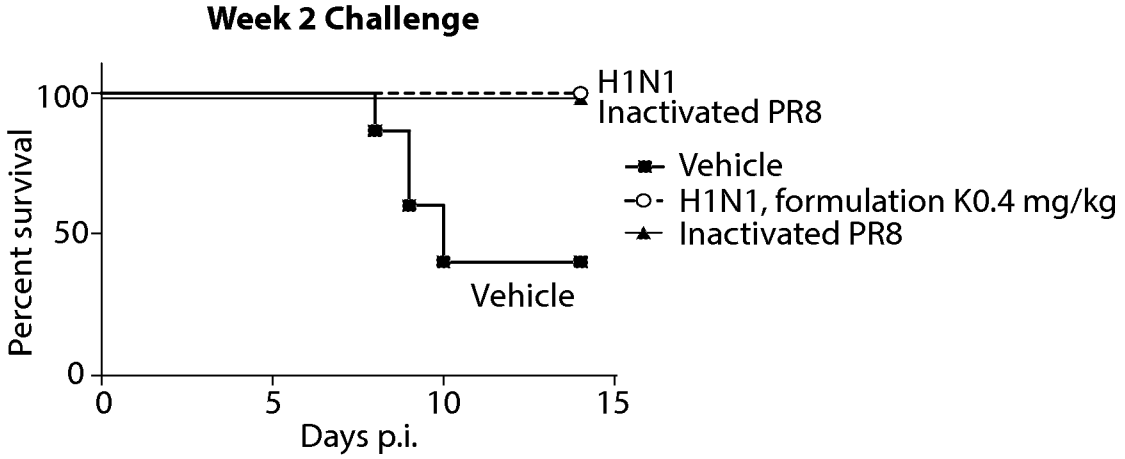


FIGURE 12B

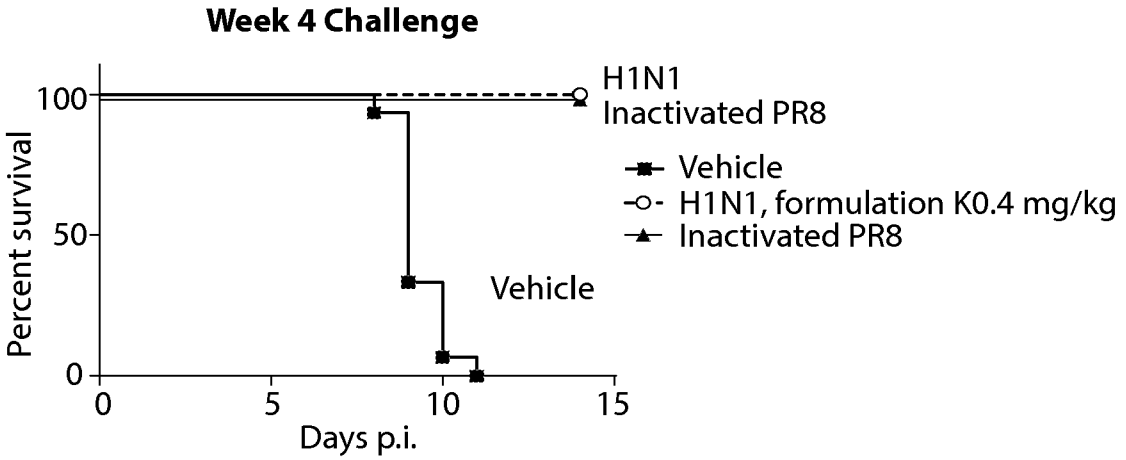


FIGURE 12C

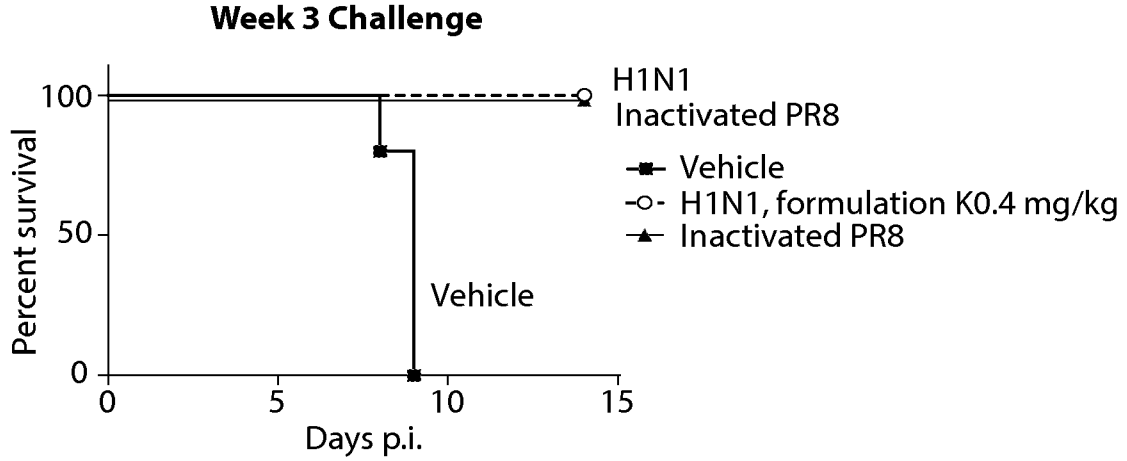


FIGURE 12D

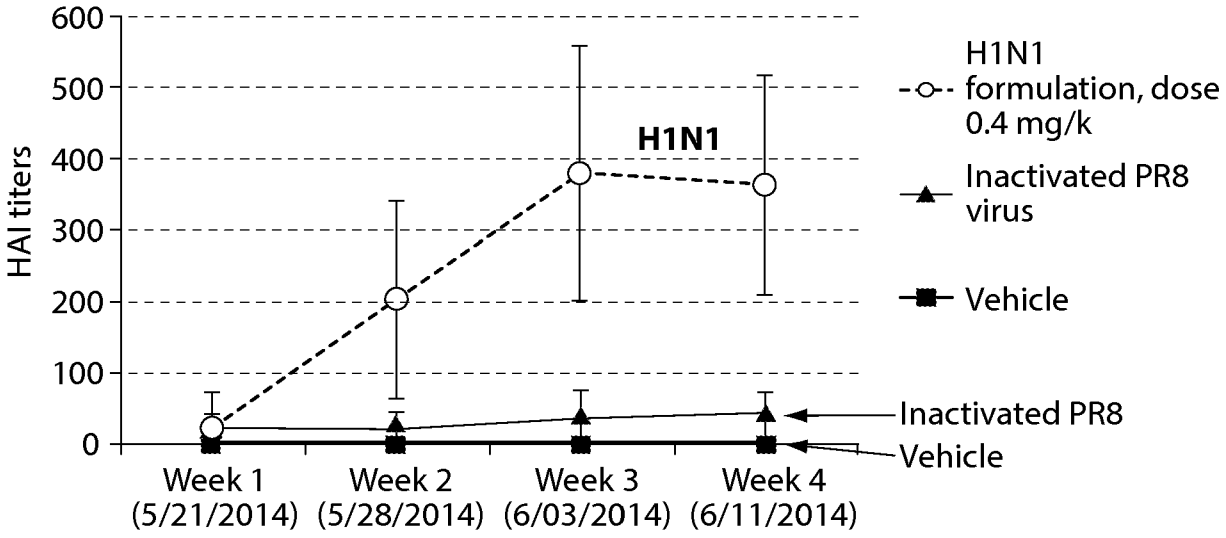

600
500
400
300
200
100
0
HAI titers
H1N1
Week 1
(5/21/2014)
Week 2
(5/28/2014)
Week 3
(6/03/2014)
Week 4
(6/11/2014)
H1N1
formulation, dose
0.4 mg/k
Inactivated PR8
virus
Vehicle
Inactivated PR8
Vehicle

FIGURE 13

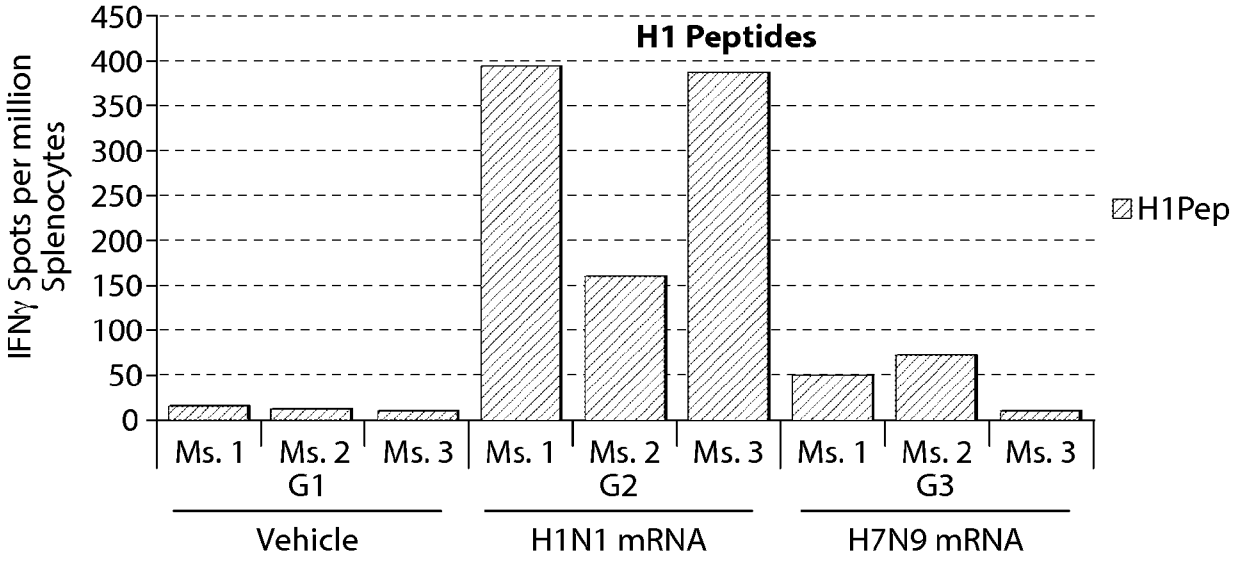


FIGURE 14A

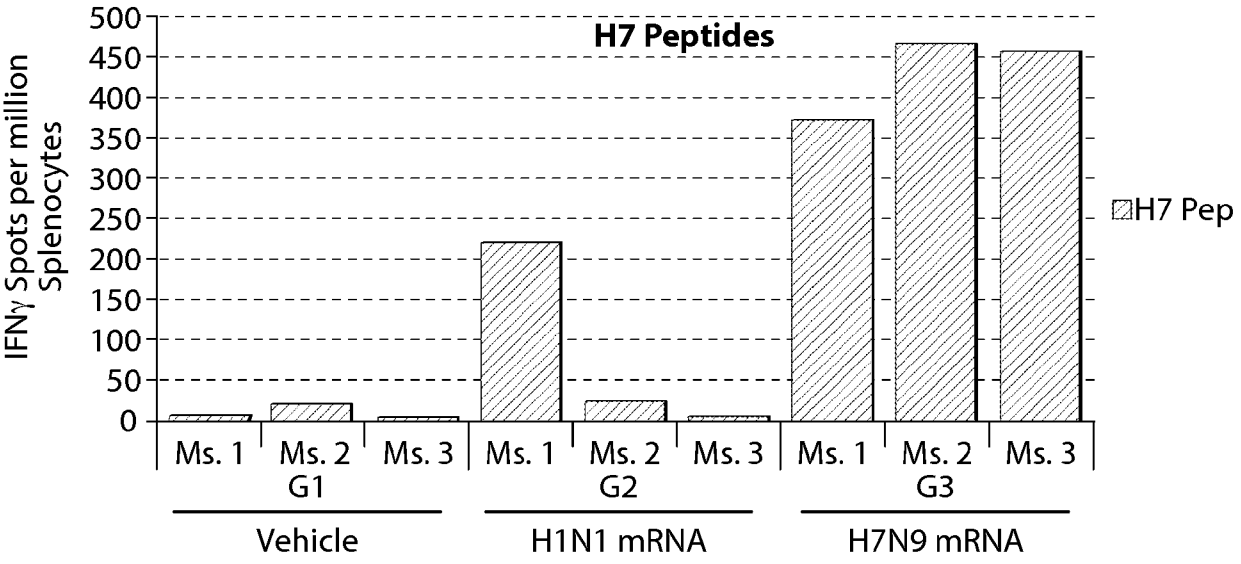


FIGURE 14B

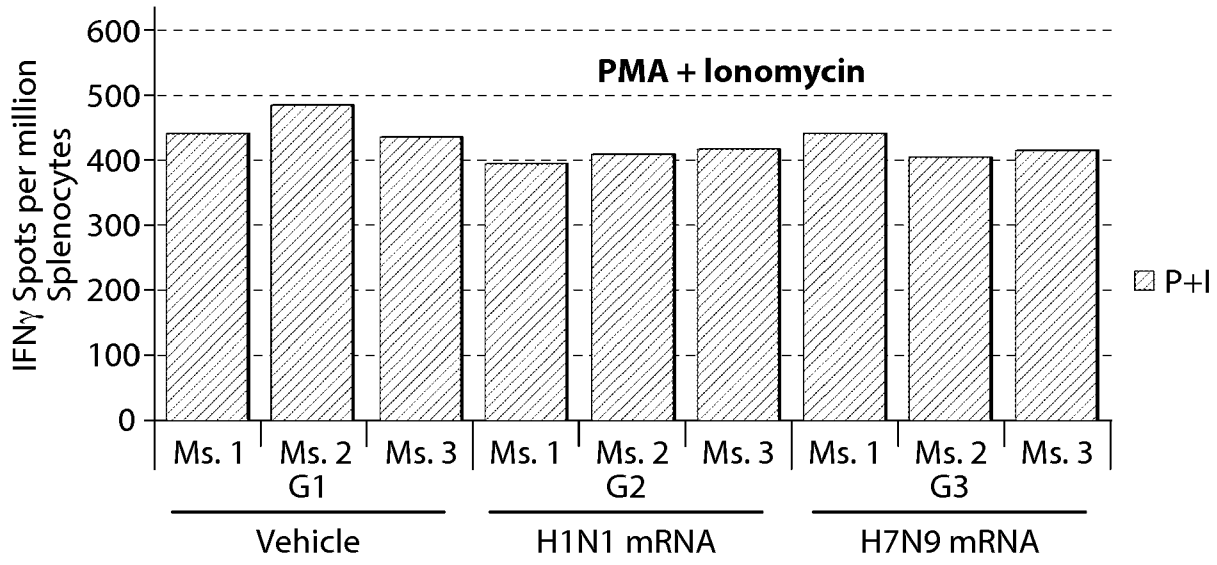

IFNγ Spots per million Splenocytes
600
500
400
300
200
100
0
PMA + Ionomycin
P+I
Ms. 1
Ms. 2
Ms. 3
Ms. 1
Ms. 2
Ms. 3
Ms. 1
Ms. 2
Ms. 3
G1
G2
G3
Vehicle
H1N1 mRNA
H7N9 mRNA

FIGURE 14C

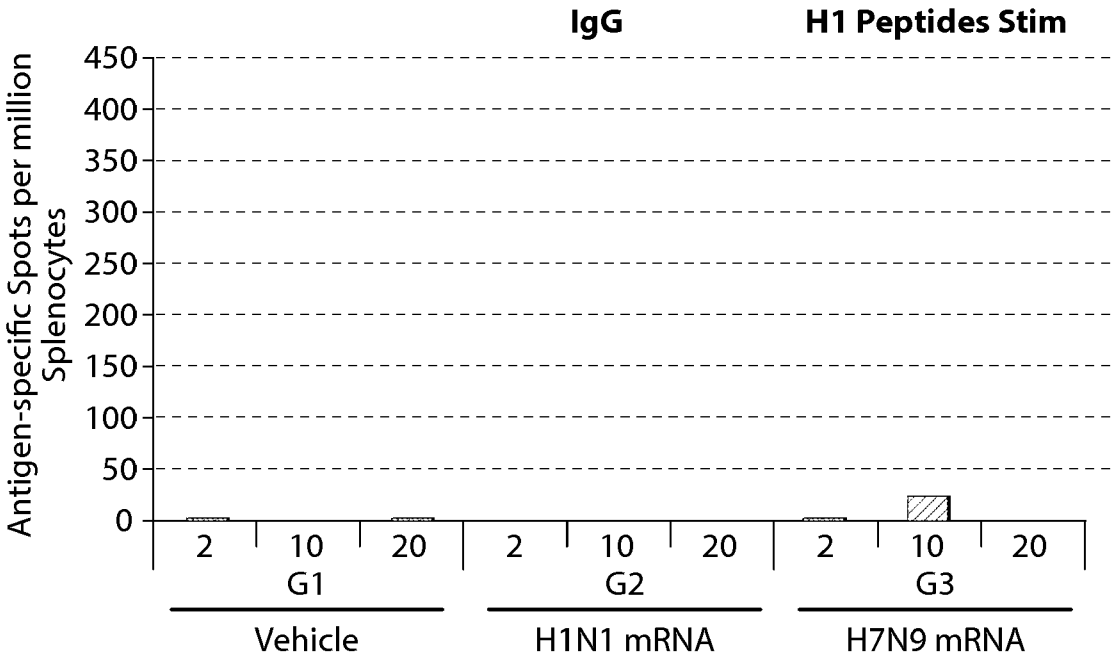


FIGURE 15A

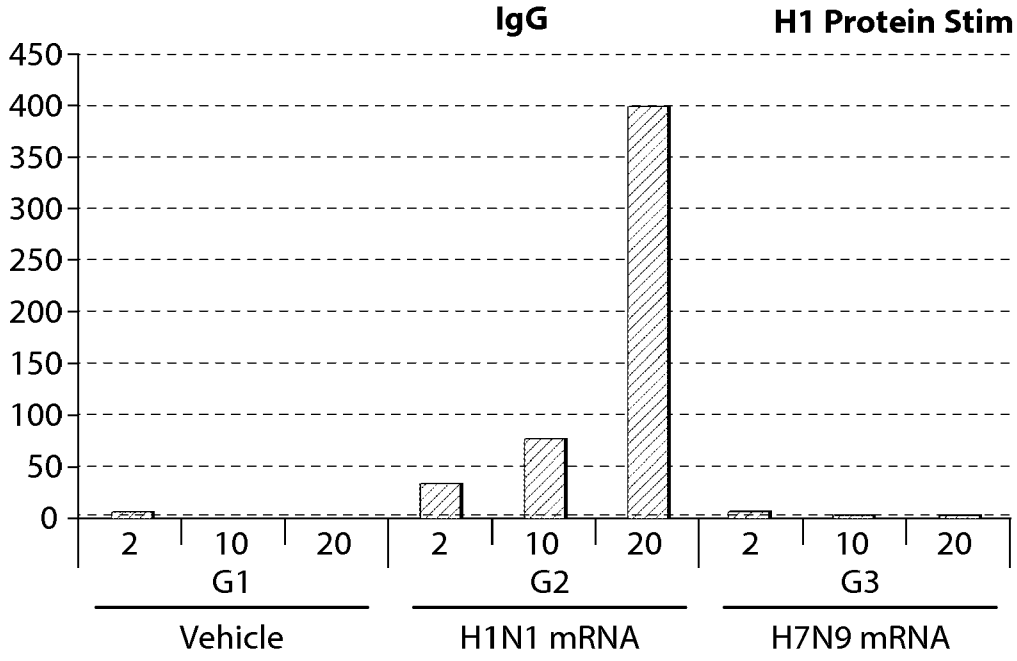


FIGURE 15B

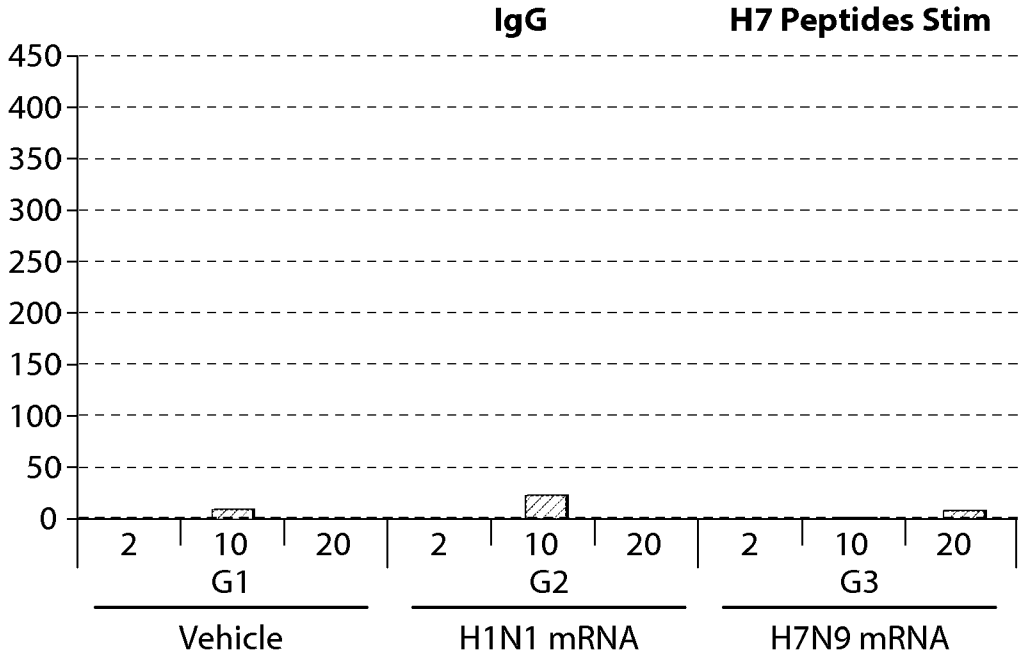


FIGURE 15C

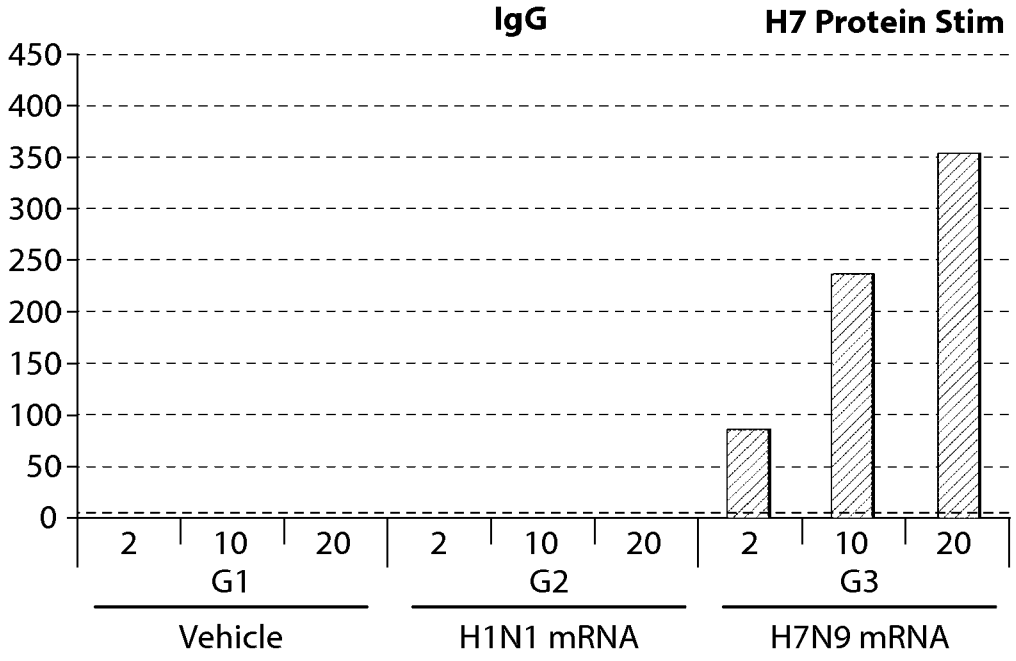


FIGURE 15D

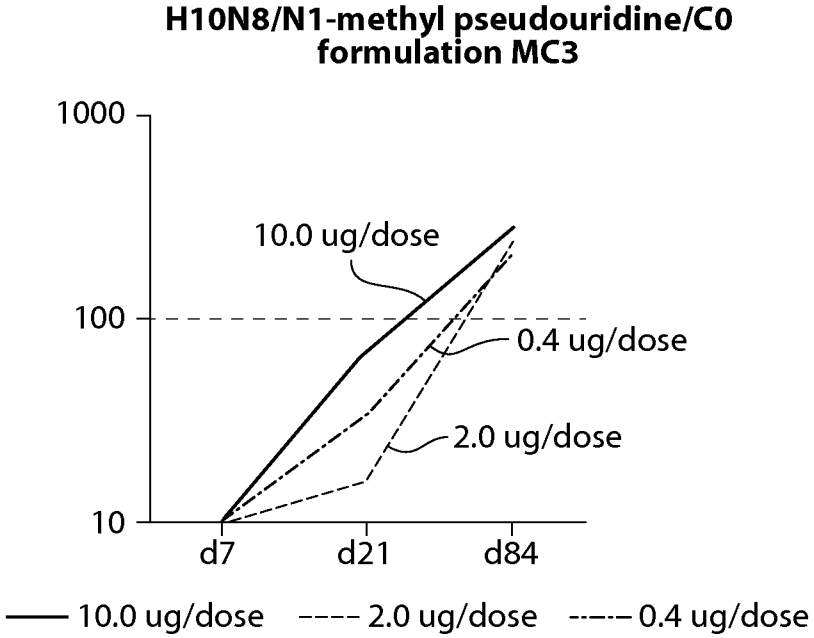


FIGURE 16

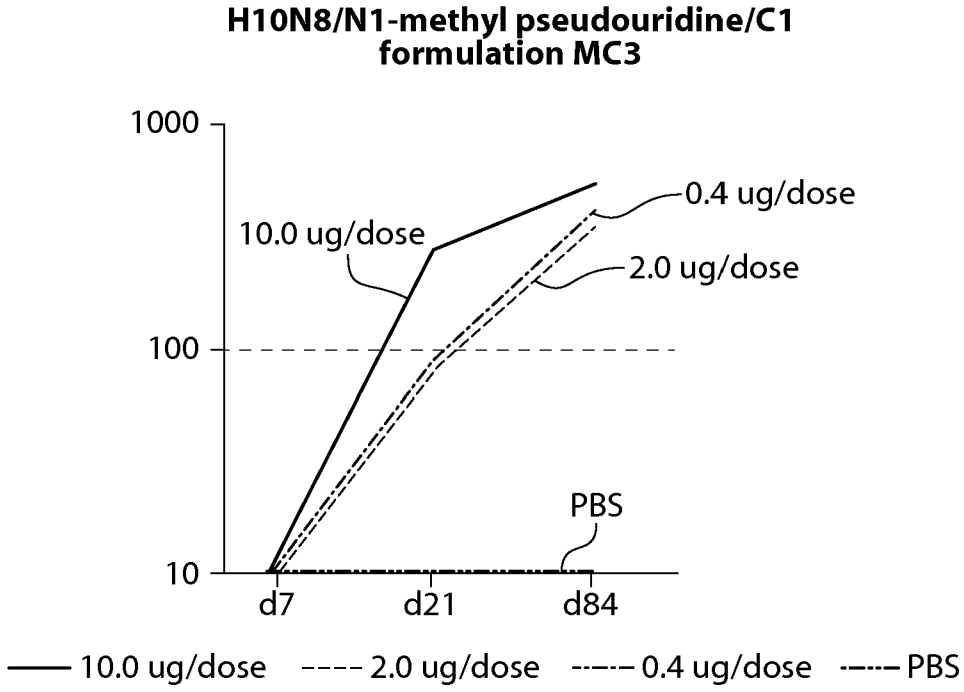


FIGURE 17

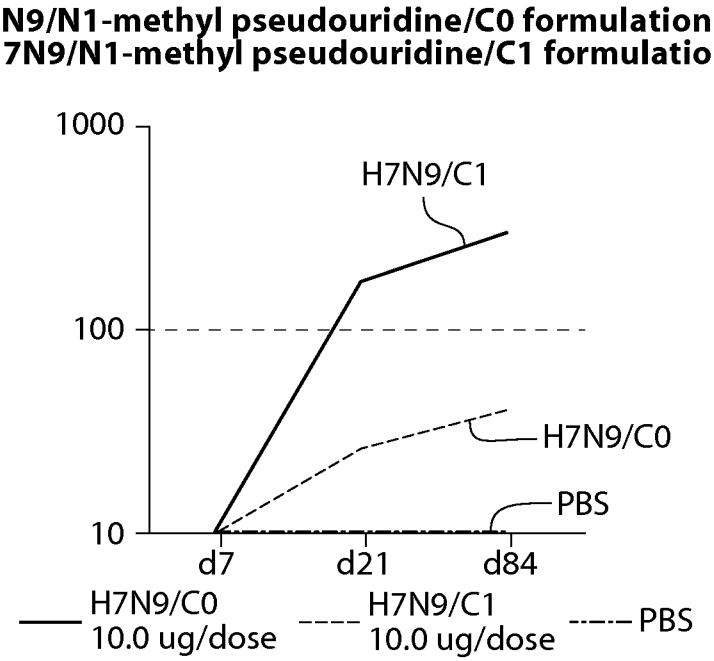


FIGURE 18

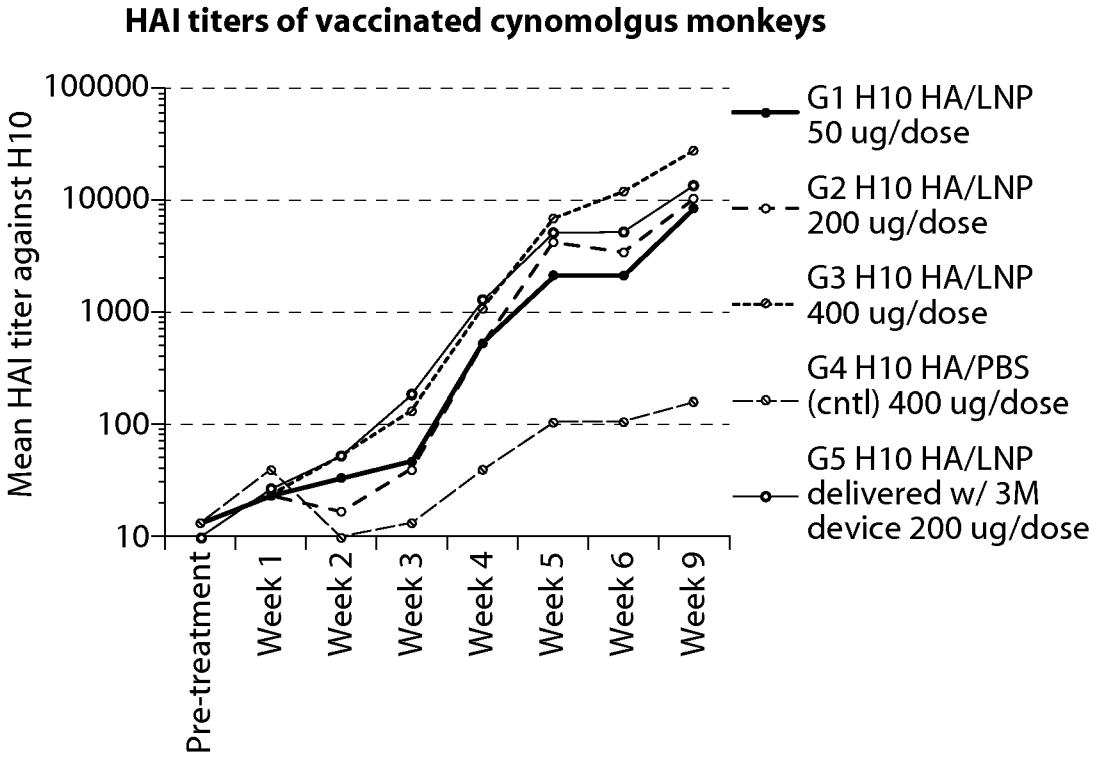


FIGURE 19

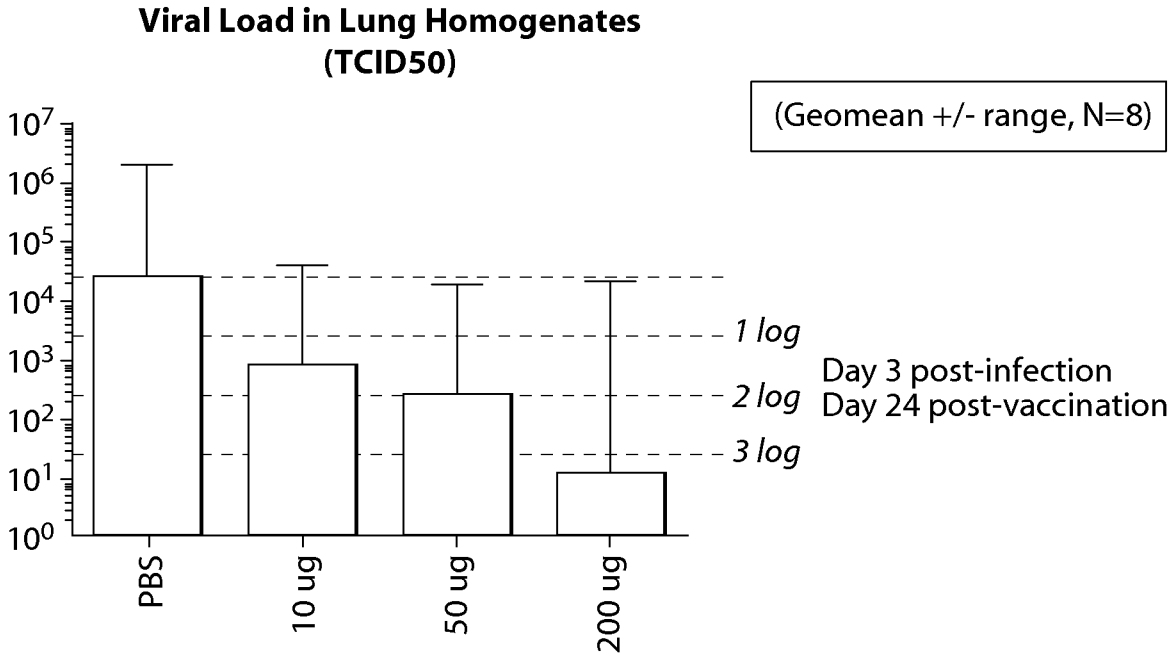

Viral Load in Lung Homogenates
(TCID50)
(Geomean +/- range, N=8)
10^7
10^6
10^5
10^4
10^3
10^2
10^1
10^0
1 log
2 log
3 log
Day 3 post-infection
Day 24 post-vaccination
PBS
10 ug
50 ug
200 ug

FIGURE 20

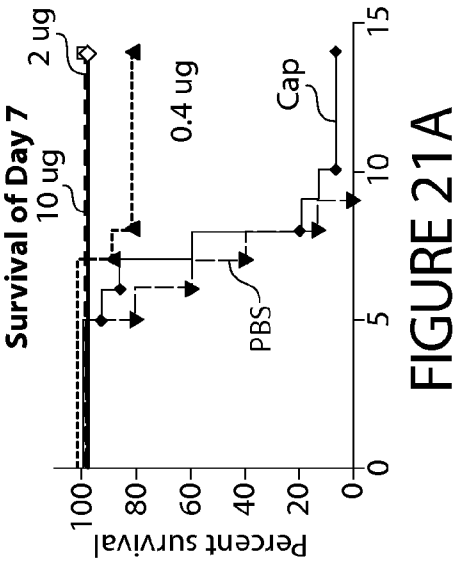


FIGURE 21A

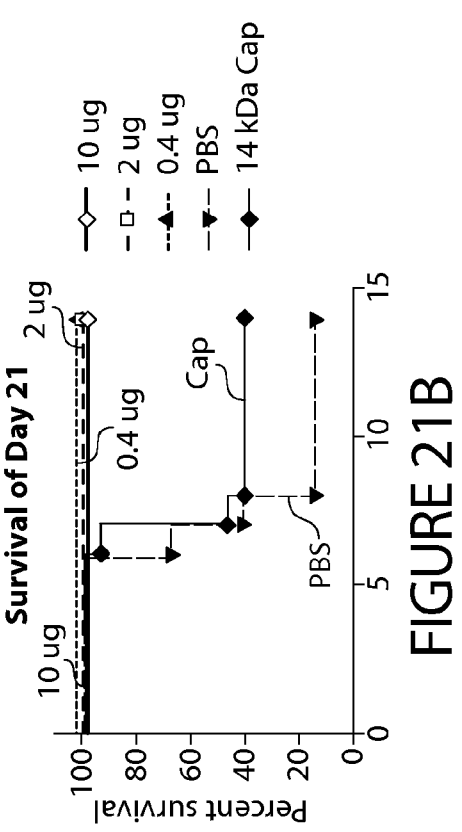


FIGURE 21B

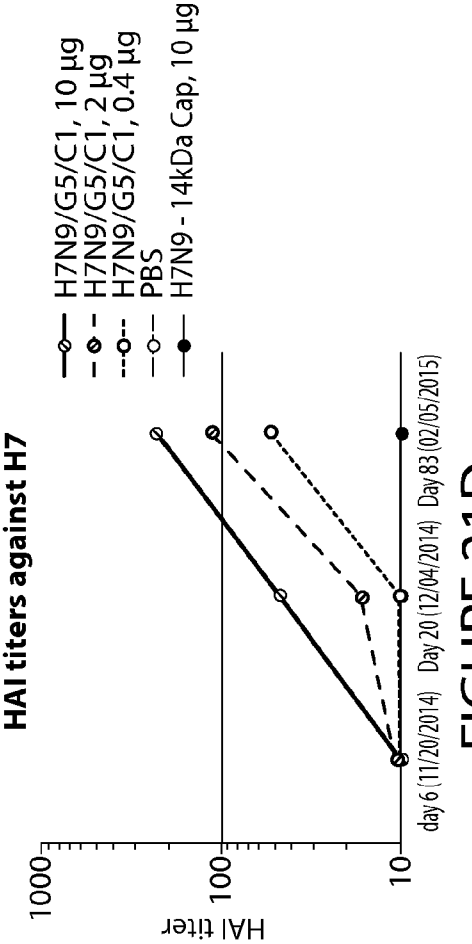


FIGURE 21D

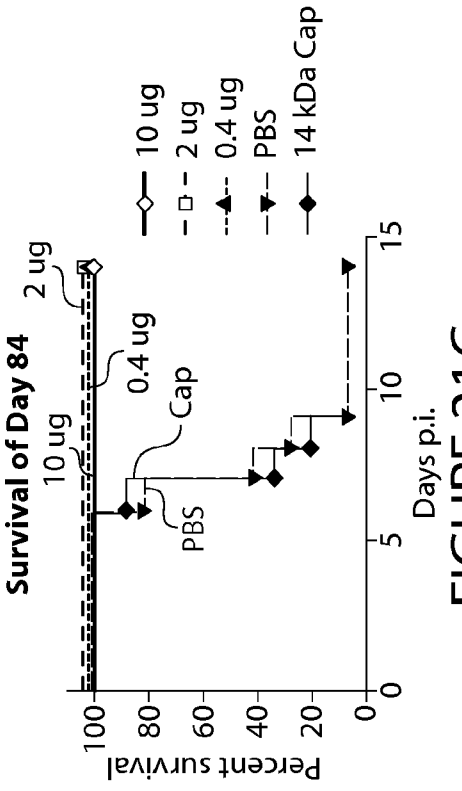


FIGURE 21C

Positions from 1 till 60

	Consensus sequence	MNTQILVFALIAIIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATETVERTNIPRIC
ACX53683	A/goose/Czech Republic/1848-K9/2009(H7N9)	..I.....V.....V....
ADN34727	A/goose/Czech Republic/1848-T14/2009(H7N9)	..I.....V.....V....
CAY39406	A/Anas crecca/Spain/1460/2008(H7N9)	.....V.....V....
AHZ39686	A/Anhui/DEWH72-01/2013(H7N9)	.....
AHZ39710	A/Anhui/DEWH72-03/2013(H7N9)	.....
AHZ39746	A/Anhui/DEWH72-06/2013(H7N9)	.....
AHM24224	A/Beijing/3/2013(H7N9)	.....
AHD25003	A/Guangdong/02/2013(H7N9)	.....
AHH25185	A/Guangdong/04/2013(H7N9)	.....
AGI60301	A/Hangzhou/1/2013(H7N9)	.....
AGK84857	A/Hangzhou/2/2013(H7N9)	.....
AHF20528	A/Hong Kong/470129/2013(H7N9)	.....
AGY41893	A/Huizhou/01/2013(H7N9)	.....
AGO51387	A/Jiangsu/2/2013(H7N9)	.....
AGJ73503	A/Nanjing/1/2013(H7N9)	.....
AGR85026	A/Nanjing/2/2013(H7N9)	.....
AGR84954	A/Nanjing/6/2013(H7N9)	.....
AHK10800	A/Shanghai/01/2014(H7N9)	.....
AGL44438	A/Shanghai/02/2013(H7N9)	.....
AGL33692	A/Shanghai/4655T/2013(H7N9)	----------
AGL33693	A/Shanghai/4659T/2013(H7N9)	----------
AGI60292	A/Shanghai/4664T/2013(H7N9)	.....A.....
AGM53887	A/Shanghai/4701T/2013(H7N9)	----------
AGM53886	A/Shanghai/4842T/2013(H7N9)	----------
AGM53883	A/Shanghai/5083T/2013(H7N9)	----------
AGM53884	A/Shanghai/5180T/2013(H7N9)	----------
AGM53885	A/Shanghai/5240T/2013(H7N9)	----------
AGW82600	A/Shanghai/CN01/2013(H7N9)	--------.....
AHF20568	A/Shanghai/CN02/2013(H7N9)	.....
AGW82612	A/Shanghai/JS01/2013(H7N9)	.....
AID70634	A/Shanghai/Mix1/2014(H7N9)	.....
AHJ57411	A/Shanghai/PD-01/2014(H7N9)	.....
AHJ57418	A/Shanghai/PD-02/2014(H7N9)	.....
AJJ96841	A/Shenzhen/SP139/2014(H7N9)	.....

FIGURE 22-1

AJJ91909	A/Shenzhen/SP26/2014(H7N9)	.....
AJJ91945	A/Shenzhen/SP38/2014(H7N9)	.....A.....
AJJ91885	A/Shenzhen/SP4/2014(H7N9)	.....
AJJ91957	A/Shenzhen/SP44/2014(H7N9)	.....
AJJ91969	A/Shenzhen/SP48/2014(H7N9)	.....
AJJ95596	A/Shenzhen/SP58/2014(H7N9)	.....A.....
AJJ95632	A/Shenzhen/SP62/2014(H7N9)	.....
AJJ95620	A/Shenzhen/SP75/2014(H7N9)	.....
AGR84942	A/Suzhou/5/2013(H7N9)	.....
AGL43637	A/Taiwan/1/2013(H7N9)	.....
AGL95088	A/Taiwan/S02076/2013(H7N9)	------.....
AGL95098	A/Taiwan/T02081/2013(H7N9)	-----.....
AGN69474	A/Wuxi/1/2013(H7N9)	.....
AGN69462	A/Wuxi/2/2013(H7N9)	.....
AGR84990	A/Wuxi/3/2013(H7N9)	.....
AGR84978	A/Wuxi/4/2013(H7N9)	.....
AGO02477	A/Xuzhou/1/2013(H7N9)	.....
AGU02230	A/Zhejiang/DTID-ZJU05/2013(H7N9)	-----.....G.....
AGU02233	A/Zhejiang/DTID-ZJU08/2013(H7N9)	-------.....G.....F....
AHA11500	A/Zhejiang/DTID-ZJU10/2013(H7N9)	--.....
AIN76383	A/Zhejiang/LS01/2014(H7N9)	.....V.....
AGR85002	A/Zhenjiang/1/2013(H7N9)	.....
ADK71137	A/blue-winged teal/Guatemala/CIP049-01/2008(H7N9)	.....ALIACML.GAKG.....A.....I.....TA..KK..
ADK71148	A/blue-winged teal/Guatemala/CIP049-02/2008(H7N9)	.....ALIACML.GAKG.....A.....I....X.....TA..KK..
ABS89409	A/blue-winged teal/Ohio/566/2006(H7N9)	.....ALIACML.GAKG.....A.....I.....TA..KK..
AJJ95048	A/chicken/Dongguan/1009/2014(H7N9)	.....
AJJ94134	A/chicken/Dongguan/1051/2014(H7N9)	.....L.....
AJJ94158	A/chicken/Dongguan/1075/2014(H7N9)	.....
AJJ95572	A/chicken/Dongguan/1100/2014(H7N9)	.....
AJJ94182	A/chicken/Dongguan/1177/2014(H7N9)	.....
AJJ95171	A/chicken/Dongguan/1314/2014(H7N9)	.....
AJJ94254	A/chicken/Dongguan/1374/2014(H7N9)	.....
AJJ95227	A/chicken/Dongguan/1382/2014(H7N9)	.....
AJJ95251	A/chicken/Dongguan/1401/2014(H7N9)	.....
AJJ94356	A/chicken/Dongguan/1456/2014(H7N9)	.....
AJJ94396	A/chicken/Dongguan/1494/2014(H7N9)	.....
AJJ95346	A/chicken/Dongguan/1548/2014(H7N9)	.....
AJJ93857	A/chicken/Dongguan/169/2014(H7N9)	.....
AJJ95382	A/chicken/Dongguan/1690/2014(H7N9)	.....

FIGURE 22-2

AJJ93869	A/chicken/Dongguan/173/2014(H7N9)
AJJ93881	A/chicken/Dongguan/189/2014(H7N9)
AJJ94606	A/chicken/Dongguan/191/2014(H7N9)
AJJ90661	A/chicken/Dongguan/2912/2013(H7N9)
AJJ90951	A/chicken/Dongguan/3544/2013(H7N9)
AJJ91155	A/chicken/Dongguan/4037/2013(H7N9)
AJJ92031	A/chicken/Dongguan/4064/2013(H7N9)
AJJ91993	A/chicken/Dongguan/4119/2013(H7N9)
AJJ93907	A/chicken/Dongguan/449/2014(H7N9)
AJJ93931	A/chicken/Dongguan/536/2014(H7N9)
AJJ93943	A/chicken/Dongguan/568/2014(H7N9)
AJJ94754	A/chicken/Dongguan/748/2014(H7N9)
AJJ94838	A/chicken/Dongguan/835/2014(H7N9)
AJJ94862	A/chicken/Dongguan/843/2014(H7N9)
AJJ94886	A/chicken/Dongguan/851/2014(H7N9)
AJJ94910	A/chicken/Dongguan/874/2014(H7N9)
AHK10585	A/chicken/Guangdong/G1/2013(H7N9)
AJJ91402	A/chicken/Huzhou/4045/2013(H7N9)
AJJ91476	A/chicken/Huzhou/4076/2013(H7N9)
AJJ93051	A/chicken/Jiangxi/10573/2014(H7N9)
AJJ96552	A/chicken/Jiangxi/12206/2014(H7N9)
AJJ96684	A/chicken/Jiangxi/13207/2014(H7N9)
AJJ96720	A/chicken/Jiangxi/13220/2014(H7N9)
AJJ96732	A/chicken/Jiangxi/13223/2014(H7N9)
AJJ97267	A/chicken/Jiangxi/13252/2014(H7N9)
AJJ97291	A/chicken/Jiangxi/13493/2014(H7N9)
AJJ96889	A/chicken/Jiangxi/13496/2014(H7N9)
AJJ96901	A/chicken/Jiangxi/13502/2014(H7N9)
AJJ97331	A/chicken/Jiangxi/13512/2014(H7N9)
AJJ96925	A/chicken/Jiangxi/13513/2014(H7N9)
AJJ97373	A/chicken/Jiangxi/13521/2014(H7N9)
AJJ97443	A/chicken/Jiangxi/13530/2014(H7N9)
AJJ97582	A/chicken/Jiangxi/14023/2014(H7N9)
AJJ97697	A/chicken/Jiangxi/14517/2014(H7N9)
AJJ97709	A/chicken/Jiangxi/14518/2014(H7N9)
AJJ97745	A/chicken/Jiangxi/14554/2014(H7N9)
AJJ97899	A/chicken/Jiangxi/15524/2014(H7N9)
AJJ96817	A/chicken/Jiangxi/9513/2014(H7N9)
AJJ93027	A/chicken/Jiangxi/9558/2014(H7N9)

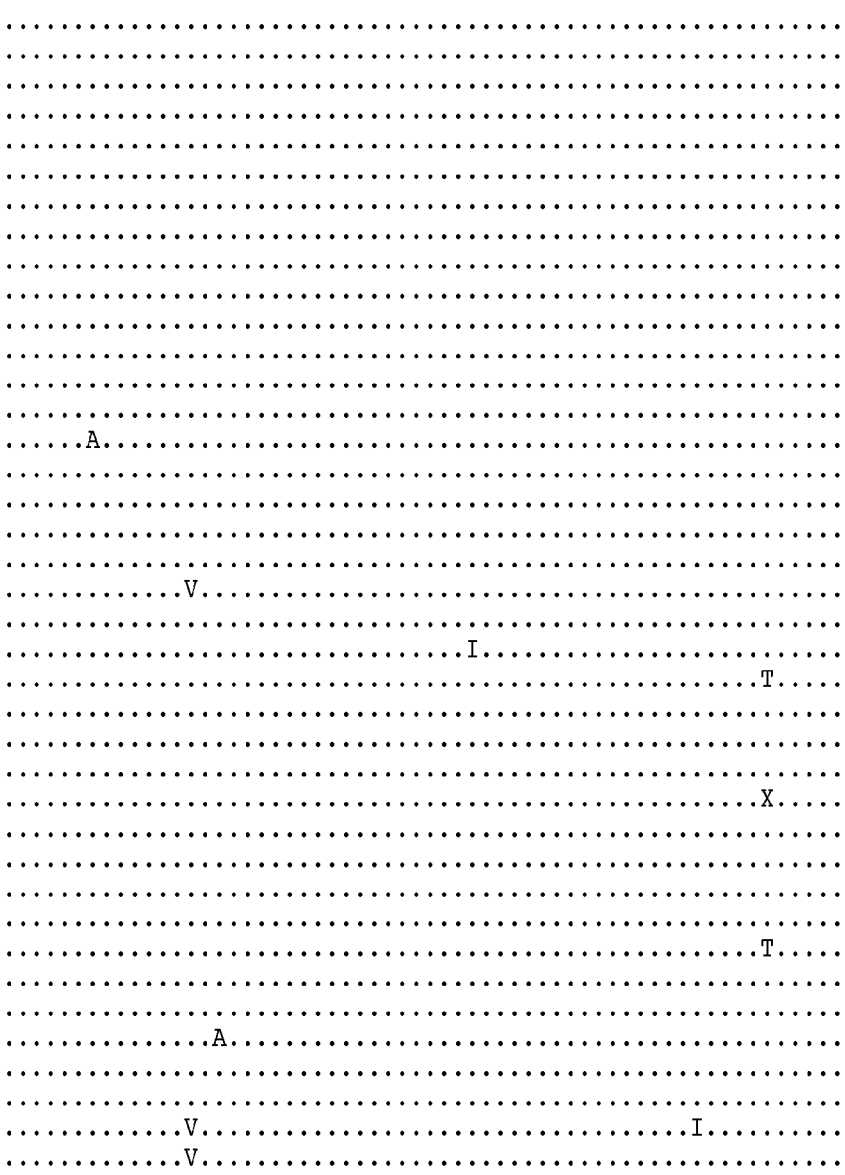


FIGURE 22-3

AGR49399	A/chicken/Jiangxi/SD001/2013(H7N9)	.....
AGQ81043	A/chicken/Rizhao/515/2013(H7N9)	.....
AGR33894	A/chicken/Rizhao/719b/2013(H7N9)	.....
AHN96472	A/chicken/Shanghai/PD-CN-02/2014(H7N9)	.....
AGR49495	A/chicken/Shanghai/S1358/2013(H7N9)	.....
AGR49506	A/chicken/Shanghai/S1410/2013(H7N9)	.....
AJJ97757	A/chicken/Shantou/2537/2014(H7N9)	.....
AJJ97973	A/chicken/Shantou/4325/2014(H7N9)	.....S.....
AJJ97998	A/chicken/Shantou/4816/2014(H7N9)	.....V.....
AJJ91314	A/chicken/Shaoxing/2417/2013(H7N9)	.....
AJJ91725	A/chicken/Shaoxing/5201/2013(H7N9)	.....
AJJ95464	A/chicken/Shenzhen/138/2014(H7N9)	.....
AJJ90576	A/chicken/Shenzhen/1665/2013(H7N9)	.....
AJJ90588	A/chicken/Shenzhen/2110/2013(H7N9)	.....
AJJ91035	A/chicken/Shenzhen/3780/2013(H7N9)	.....
AJJ90490	A/chicken/Shenzhen/742/2013(H7N9)	.....
AJJ92005	A/chicken/Shenzhen/801/2013(H7N9)	.....
AJJ90526	A/chicken/Shenzhen/898/2013(H7N9)	.....
AIU47013	A/chicken/Suzhou/040201H/2013(H7N9)	.....
AGJ72861	A/chicken/Zhejiang/DTID-ZJU01/2013(H7N9)	.....G.....
AIU46619	A/chicken/Zhejiang/DTID-ZJU06/2013(H7N9)	.....
AGR49554	A/chicken/Zhejiang/SD033/2013(H7N9)	.....
AGR49566	A/duck/Anhui/SC702/2013(H7N9)	.....
BAN16711	A/duck/Gunma/466/2011(H7N9)	....V.....M.....V....
AJJ97841	A/duck/Jiangxi/15044/2014(H7N9)	.....
AGQ80952	A/duck/Jiangxi/3096/2009(H7N9)	.....V.....S.....
AGQ80989	A/duck/Jiangxi/3257/2009(H7N9)	.....V.....S.....
BAH22785	A/duck/Mongolia/119/2008(H7N9)	.....V.....
BAN59727	A/duck/Mongolia/129/2010(H7N9)	.....V.....I.V....
BAN59726	A/duck/Mongolia/147/2008(H7N9)	.....V.....
AJK00354	A/duck/Zhejiang/LS02/2014(H7N9)	.....V.....
AFM09439	A/emperor goose/Alaska/44063-061/2006(H7N9)	---...A.IACML.GAKG.....A.....I.....TV..KK..
AFV33947	A/goose/Nebraska/17097-4/2011(H7N9)	......ALIACML.GAKG.....A.....I.....TA..KK..
AFV33945	A/guinea fowl/Nebraska/17096-1/2011(H7N9)	......ALIACML.GAKG.....A.....RI.....TA..KK..
AHA11441	A/guinea fowl/Nebraska/17096/2011(H7N9)	......ALIACML.GAKG.....A.....I.....TA..KK..
AGR49722	A/homing pigeon/Jiangsu/SD184/2013(H7N9)	.....
AGL97639	A/mallard/Minnesota/AI09-3770/2009(H7N9)	--------IACMLVGAKG.....A.....I.....TA..KKL.
AHZ42537	A/mallard/Minnesota/AI09-3770/2009(H7N9)	......A.IACMLVGAKG.....A.....I.....TA..KKL.
ADC29485	A/mallard/Spain/08.00991.3/2005(H7N9)	----------

FIGURE 22-4

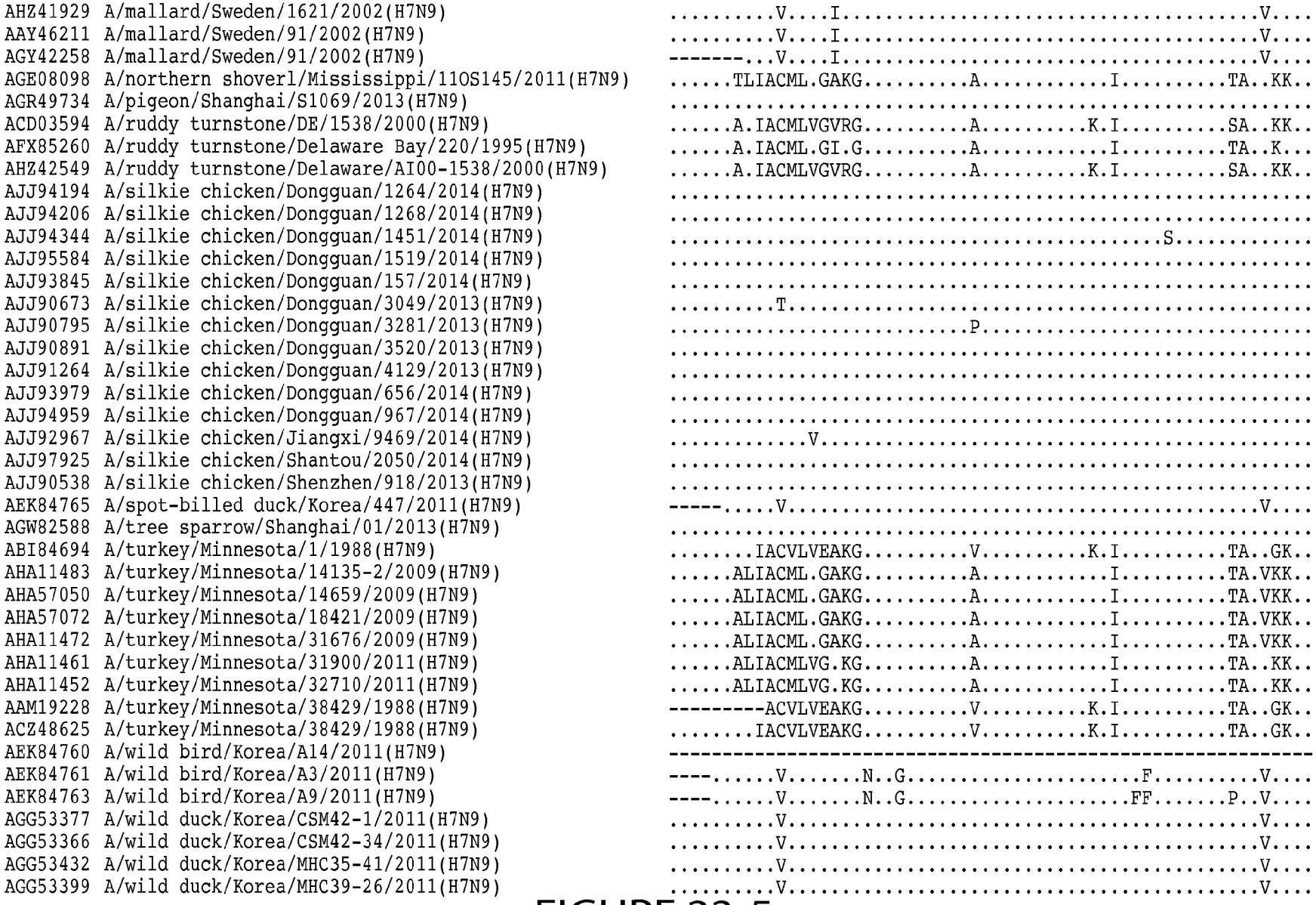

AHZ41929 A/mallard/Sweden/1621/2002(H7N9)
AAY46211 A/mallard/Sweden/91/2002(H7N9)
AGY42258 A/mallard/Sweden/91/2002(H7N9)
AGE08098 A/northern shoverl/Mississippi/11OS145/2011(H7N9)
AGR49734 A/pigeon/Shanghai/S1069/2013(H7N9)
ACD03594 A/ruddy turnstone/DE/1538/2000(H7N9)
AFX85260 A/ruddy turnstone/Delaware Bay/220/1995(H7N9)
AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000(H7N9)
AJJ94194 A/silkie chicken/Dongguan/1264/2014(H7N9)
AJJ94206 A/silkie chicken/Dongguan/1268/2014(H7N9)
AJJ94344 A/silkie chicken/Dongguan/1451/2014(H7N9)
AJJ95584 A/silkie chicken/Dongguan/1519/2014(H7N9)
AJJ93845 A/silkie chicken/Dongguan/157/2014(H7N9)
AJJ90673 A/silkie chicken/Dongguan/3049/2013(H7N9)
AJJ90795 A/silkie chicken/Dongguan/3281/2013(H7N9)
AJJ90891 A/silkie chicken/Dongguan/3520/2013(H7N9)
AJJ91264 A/silkie chicken/Dongguan/4129/2013(H7N9)
AJJ93979 A/silkie chicken/Dongguan/656/2014(H7N9)
AJJ94959 A/silkie chicken/Dongguan/967/2014(H7N9)
AJJ92967 A/silkie chicken/Jiangxi/9469/2014(H7N9)
AJJ97925 A/silkie chicken/Shantou/2050/2014(H7N9)
AJJ90538 A/silkie chicken/Shenzhen/918/2013(H7N9)
AEK84765 A/spot-billed duck/Korea/447/2011(H7N9)
AGW82588 A/tree sparrow/Shanghai/01/2013(H7N9)
ABI84694 A/turkey/Minnesota/1/1988(H7N9)
AHA11483 A/turkey/Minnesota/14135-2/2009(H7N9)
AHA57050 A/turkey/Minnesota/14659/2009(H7N9)
AHA57072 A/turkey/Minnesota/18421/2009(H7N9)
AHA11472 A/turkey/Minnesota/31676/2009(H7N9)
AHA11461 A/turkey/Minnesota/31900/2011(H7N9)
AHA11452 A/turkey/Minnesota/32710/2011(H7N9)
AAM19228 A/turkey/Minnesota/38429/1988(H7N9)
ACZ48625 A/turkey/Minnesota/38429/1988(H7N9)
AEK84760 A/wild bird/Korea/A14/2011(H7N9)
AEK84761 A/wild bird/Korea/A3/2011(H7N9)
AEK84763 A/wild bird/Korea/A9/2011(H7N9)
AGG53377 A/wild duck/Korea/CSM42-1/2011(H7N9)
AGG53366 A/wild duck/Korea/CSM42-34/2011(H7N9)
AGG53432 A/wild duck/Korea/MHC35-41/2011(H7N9)
AGG53399 A/wild duck/Korea/MHC39-26/2011(H7N9)

FIGURE 22-5

AGG53476	A/wild duck/Korea/SH19-27/2010(H7N9)	.....V.....V....
AGG53487	A/wild duck/Korea/SH19-50/2010(H7N9)	.....V.....V....
AGG53520	A/wild duck/Korea/SH20-27/2008(H7N9)	---.....V.....V....
AEM98291	A/wild duck/Mongolia/1-241/2008(H7N9)	---S.....V.....V....
AGR49770	A/wild pigeon/Jiangsu/SD001/2013(H7N9)	.....

Positions from 61 till 120

	Consensus sequence	**SKGKRTVDLGQCGLLGTITGPPQCDQFLEFSADLIIERREGSDVCYPGKFVNEEALRQIL**
ACX53683	A/goose/Czech Republic/1848-K9/2009(H7N9)	.....G.....
ADN34727	A/goose/Czech Republic/1848-T14/2009(H7N9)	.....G.....
CAY39406	A/Anas crecca/Spain/1460/2008(H7N9)	.....
AHZ39686	A/Anhui/DEWH72-01/2013(H7N9)	.....
AHZ39710	A/Anhui/DEWH72-03/2013(H7N9)	.....
AHZ39746	A/Anhui/DEWH72-06/2013(H7N9)	.....
AHM24224	A/Beijing/3/2013(H7N9)	.....K.....
AHD25003	A/Guangdong/02/2013(H7N9)	....K.....
AHH25185	A/Guangdong/04/2013(H7N9)	....K.....
AGI60301	A/Hangzhou/1/2013(H7N9)	.....
AGK84857	A/Hangzhou/2/2013(H7N9)	.....
AHF20528	A/Hong Kong/470129/2013(H7N9)	.....
AGY41893	A/Huizhou/01/2013(H7N9)	....K.....
AGO51387	A/Jiangsu/2/2013(H7N9)	....M.....
AGJ73503	A/Nanjing/1/2013(H7N9)	....M.....
AGR85026	A/Nanjing/2/2013(H7N9)	....M.....
AGR84954	A/Nanjing/6/2013(H7N9)	.....
AHK10800	A/Shanghai/01/2014(H7N9)	.....
AGL44438	A/Shanghai/02/2013(H7N9)	.....
AGL33692	A/Shanghai/4655T/2013(H7N9)	----------
AGL33693	A/Shanghai/4659T/2013(H7N9)	----------
AGI60292	A/Shanghai/4664T/2013(H7N9)	.....
AGM53887	A/Shanghai/4701T/2013(H7N9)	----------
AGM53886	A/Shanghai/4842T/2013(H7N9)	----------
AGM53883	A/Shanghai/5083T/2013(H7N9)	----------
AGM53884	A/Shanghai/5180T/2013(H7N9)	----------
AGM53885	A/Shanghai/5240T/2013(H7N9)	----------
AGW82600	A/Shanghai/CN01/2013(H7N9)	.....
AHF20568	A/Shanghai/CN02/2013(H7N9)	.....
AGW82612	A/Shanghai/JS01/2013(H7N9)	.....

FIGURE 22-6

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AID70634 A/Shanghai/Mix1/2014(H7N9)                                  ..................................................
AHJ57411 A/Shanghai/PD-01/2014(H7N9)                                 ..................................................
AHJ57418 A/Shanghai/PD-02/2014(H7N9)                                 ..........................................I.......
AJJ96841 A/Shenzhen/SP139/2014(H7N9)                                 ....K.............................................
AJJ91909 A/Shenzhen/SP26/2014(H7N9)                                  ....K.....................................I.......
AJJ91945 A/Shenzhen/SP38/2014(H7N9)                                  ....K.............................................
AJJ91885 A/Shenzhen/SP4/2014(H7N9)                                   ....K.............................................
AJJ91957 A/Shenzhen/SP44/2014(H7N9)                                  ..................................................
AJJ91969 A/Shenzhen/SP48/2014(H7N9)                                  ....K.............................................
AJJ95596 A/Shenzhen/SP58/2014(H7N9)                                  ....K.............................................
AJJ95632 A/Shenzhen/SP62/2014(H7N9)                                  ....K.............................................
AJJ95620 A/Shenzhen/SP75/2014(H7N9)                                  ....K.............................................
AGR84942 A/Suzhou/5/2013(H7N9)                                       ..................................................
AGL43637 A/Taiwan/1/2013(H7N9)                                       ..................................................
AGL95088 A/Taiwan/S02076/2013(H7N9)                                  ..................................................
AGL95098 A/Taiwan/T02081/2013(H7N9)                                  ..................................................
AGN69474 A/Wuxi/1/2013(H7N9)                                         ..................................................
AGN69462 A/Wuxi/2/2013(H7N9)                                         ..................................................
AGR84990 A/Wuxi/3/2013(H7N9)                                         ..................................................
AGR84978 A/Wuxi/4/2013(H7N9)                                         ....K.............................................
AGO02477 A/Xuzhou/1/2013(H7N9)                                       ..................................................
AGU02230 A/Zhejiang/DTID-ZJU05/2013(H7N9)                            ..............R...................................
AGU02233 A/Zhejiang/DTID-ZJU08/2013(H7N9)                            ..............R...................................
AHA11500 A/Zhejiang/DTID-ZJU10/2013(H7N9)                            ..................................................
AIN76383 A/Zhejiang/LS01/2014(H7N9)                                  ....K.............................................
AGR85002 A/Zhenjiang/1/2013(H7N9)                                    ....M.............................................
ADK71137 A/blue-winged teal/Guatemala/CIP049-01/2008(H7N9)           TQ...PT..........LI...........D..........T........T...S.....
ADK71148 A/blue-winged teal/Guatemala/CIP049-02/2008(H7N9)           TH...PT..........LI......R....D..........T........T...S.....
ABS89409 A/blue-winged teal/Ohio/566/2006(H7N9)                      TQ...PT..........LI...........DT.........T........T...S.....
AJJ95048 A/chicken/Dongguan/1009/2014(H7N9)                          ..................................................
AJJ94134 A/chicken/Dongguan/1051/2014(H7N9)                          ....K.............................................
AJJ94158 A/chicken/Dongguan/1075/2014(H7N9)                          ....K.............................................
AJJ95572 A/chicken/Dongguan/1100/2014(H7N9)                          ..................................................
AJJ94182 A/chicken/Dongguan/1177/2014(H7N9)                          ....K.............................................
AJJ95171 A/chicken/Dongguan/1314/2014(H7N9)                          ....K.............................................
AJJ94254 A/chicken/Dongguan/1374/2014(H7N9)                          ....K.............................................
AJJ95227 A/chicken/Dongguan/1382/2014(H7N9)                          ....K.....................................I.......
AJJ95251 A/chicken/Dongguan/1401/2014(H7N9)                          ....K.............................................
AJJ94356 A/chicken/Dongguan/1456/2014(H7N9)                          ....K.............................................
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FIGURE 22-7

AJJ94396	A/chicken/Dongguan/1494/2014(H7N9)
AJJ95346	A/chicken/Dongguan/1548/2014(H7N9)
AJJ93857	A/chicken/Dongguan/169/2014(H7N9)
AJJ95382	A/chicken/Dongguan/1690/2014(H7N9)
AJJ93869	A/chicken/Dongguan/173/2014(H7N9)
AJJ93881	A/chicken/Dongguan/189/2014(H7N9)
AJJ94606	A/chicken/Dongguan/191/2014(H7N9)
AJJ90661	A/chicken/Dongguan/2912/2013(H7N9)
AJJ90951	A/chicken/Dongguan/3544/2013(H7N9)
AJJ91155	A/chicken/Dongguan/4037/2013(H7N9)
AJJ92031	A/chicken/Dongguan/4064/2013(H7N9)
AJJ91993	A/chicken/Dongguan/4119/2013(H7N9)
AJJ93907	A/chicken/Dongguan/449/2014(H7N9)
AJJ93931	A/chicken/Dongguan/536/2014(H7N9)
AJJ93943	A/chicken/Dongguan/568/2014(H7N9)
AJJ94754	A/chicken/Dongguan/748/2014(H7N9)
AJJ94838	A/chicken/Dongguan/835/2014(H7N9)
AJJ94862	A/chicken/Dongguan/843/2014(H7N9)
AJJ94886	A/chicken/Dongguan/851/2014(H7N9)
AJJ94910	A/chicken/Dongguan/874/2014(H7N9)
AHK10585	A/chicken/Guangdong/G1/2013(H7N9)
AJJ91402	A/chicken/Huzhou/4045/2013(H7N9)
AJJ91476	A/chicken/Huzhou/4076/2013(H7N9)
AJJ93051	A/chicken/Jiangxi/10573/2014(H7N9)
AJJ96552	A/chicken/Jiangxi/12206/2014(H7N9)
AJJ96684	A/chicken/Jiangxi/13207/2014(H7N9)
AJJ96720	A/chicken/Jiangxi/13220/2014(H7N9)
AJJ96732	A/chicken/Jiangxi/13223/2014(H7N9)
AJJ97267	A/chicken/Jiangxi/13252/2014(H7N9)
AJJ97291	A/chicken/Jiangxi/13493/2014(H7N9)
AJJ96889	A/chicken/Jiangxi/13496/2014(H7N9)
AJJ96901	A/chicken/Jiangxi/13502/2014(H7N9)
AJJ97331	A/chicken/Jiangxi/13512/2014(H7N9)
AJJ96925	A/chicken/Jiangxi/13513/2014(H7N9)
AJJ97373	A/chicken/Jiangxi/13521/2014(H7N9)
AJJ97443	A/chicken/Jiangxi/13530/2014(H7N9)
AJJ97582	A/chicken/Jiangxi/14023/2014(H7N9)
AJJ97697	A/chicken/Jiangxi/14517/2014(H7N9)
AJJ97709	A/chicken/Jiangxi/14518/2014(H7N9)

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FIGURE 22-8

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AJJ97745 A/chicken/Jiangxi/14554/2014(H7N9)                    ...K.....................................................
AJJ97899 A/chicken/Jiangxi/15524/2014(H7N9)                    ....................................................
AJJ96817 A/chicken/Jiangxi/9513/2014(H7N9)                     ...K.....................................................
AJJ93027 A/chicken/Jiangxi/9558/2014(H7N9)                     ...K.....................................................K.....
AGR49399 A/chicken/Jiangxi/SD001/2013(H7N9)                    ....................................................
AGQ81043 A/chicken/Rizhao/515/2013(H7N9)                       ....................................................
AGR33894 A/chicken/Rizhao/719b/2013(H7N9)                      ....................................................
AHN96472 A/chicken/Shanghai/PD-CN-02/2014(H7N9)                ...K.....................................................
AGR49495 A/chicken/Shanghai/S1358/2013(H7N9)                   ...M.....................................................
AGR49506 A/chicken/Shanghai/S1410/2013(H7N9)                   ....................................................
AJJ97757 A/chicken/Shantou/2537/2014(H7N9)                     ...K.....................................................
AJJ97973 A/chicken/Shantou/4325/2014(H7N9)                     ...K.....................................................
AJJ97998 A/chicken/Shantou/4816/2014(H7N9)                     ...K.....................................................
AJJ91314 A/chicken/Shaoxing/2417/2013(H7N9)                    ....................................................
AJJ91725 A/chicken/Shaoxing/5201/2013(H7N9)                    ....................................................
AJJ95464 A/chicken/Shenzhen/138/2014(H7N9)                     ...K.....................................................
AJJ90576 A/chicken/Shenzhen/1665/2013(H7N9)                    ...K.....................................I...............
AJJ90588 A/chicken/Shenzhen/2110/2013(H7N9)                    ...K.....................................................
AJJ91035 A/chicken/Shenzhen/3780/2013(H7N9)                    ...K.....................................................
AJJ90490 A/chicken/Shenzhen/742/2013(H7N9)                     ...K.....................................................
AJJ92005 A/chicken/Shenzhen/801/2013(H7N9)                     ...K.....................................................
AJJ90526 A/chicken/Shenzhen/898/2013(H7N9)                     ...K.....................................I...............
AIU47013 A/chicken/Suzhou/040201H/2013(H7N9)                   ....................................................
AGJ72861 A/chicken/Zhejiang/DTID-ZJU01/2013(H7N9)              ...K.....G.PR.....................M.....................
AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013(H7N9)              ....................................................
AGR49554 A/chicken/Zhejiang/SD033/2013(H7N9)                   ....................................................
AGR49566 A/duck/Anhui/SC702/2013(H7N9)                         ....................................................
BAN16711 A/duck/Gunma/466/2011(H7N9)                           ....................................................
AJJ97841 A/duck/Jiangxi/15044/2014(H7N9)                       ....................................................
AGQ80952 A/duck/Jiangxi/3096/2009(H7N9)                        ....A.....A.....A.....A.....A.....A.....A.....A.....
AGQ80989 A/duck/Jiangxi/3257/2009(H7N9)                        ....A.....A.....A.....A.....A.....A.....A.....A.....
BAH22785 A/duck/Mongolia/119/2008(H7N9)                        ....................................................
BAN59727 A/duck/Mongolia/129/2010(H7N9)                        ....................................................
BAN59726 A/duck/Mongolia/147/2008(H7N9)                        ....................................................
AJK00354 A/duck/Zhejiang/LS02/2014(H7N9)                       ....K.....................................V.....K.T.....
AFM09439 A/emperor goose/Alaska/44063-061/2006(H7N9)           TQ...PT.....................LI.....................D.....K.T.....T...S.....
AFV33947 A/goose/Nebraska/17097-4/2011(H7N9)                   TQ...PT.....................LI.....................D.....T.....T...S.....
AFV33945 A/guinea fowl/Nebraska/17096-1/2011(H7N9)             TQ...PT.....................LI.....................D.....T.....T...S.....
AHA11441 A/guinea fowl/Nebraska/17096/2011(H7N9)               TQ...PT.....................LI.....................D.....T.....T...S.....
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FIGURE 22-9

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AGR49722 A/homing pigeon/Jiangsu/SD184/2013(H7N9)
AGL97639 A/mallard/Minnesota/AI09-3770/2009(H7N9)
AHZ42537 A/mallard/Minnesota/AI09-3770/2009(H7N9)
ADC29485 A/mallard/Spain/08.00991.3/2005(H7N9)
AHZ41929 A/mallard/Sweden/1621/2002(H7N9)
AAY46211 A/mallard/Sweden/91/2002(H7N9)
AGY42258 A/mallard/Sweden/91/2002(H7N9)
AGE08098 A/northern shoveler/Mississippi/11OS145/2011(H7N9)
AGR49734 A/pigeon/Shanghai/S1069/2013(H7N9)
ACD03594 A/ruddy turnstone/DE/1538/2000(H7N9)
AFX85260 A/ruddy turnstone/Delaware Bay/220/1995(H7N9)
AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000(H7N9)
AJU94194 A/silkie chicken/Dongguan/1264/2014(H7N9)
AJU94206 A/silkie chicken/Dongguan/1268/2014(H7N9)
AJU94344 A/silkie chicken/Dongguan/1451/2014(H7N9)
AJU95584 A/silkie chicken/Dongguan/1519/2014(H7N9)
AJU93845 A/silkie chicken/Dongguan/157/2014(H7N9)
AJU90673 A/silkie chicken/Dongguan/3049/2013(H7N9)
AJU90795 A/silkie chicken/Dongguan/3281/2013(H7N9)
AJU90891 A/silkie chicken/Dongguan/3520/2013(H7N9)
AJU91264 A/silkie chicken/Dongguan/4129/2013(H7N9)
AJU93979 A/silkie chicken/Dongguan/656/2014(H7N9)
AJU94959 A/silkie chicken/Dongguan/967/2014(H7N9)
AJU92967 A/silkie chicken/Jiangxi/9469/2014(H7N9)
AJU97925 A/silkie chicken/Shantou/2050/2014(H7N9)
AJU90538 A/silkie chicken/Shenzhen/918/2013(H7N9)
AEK84765 A/spot-billed duck/Korea/447/2011(H7N9)
AGW82588 A/tree sparrow/Shanghai/01/2013(H7N9)
ABI84694 A/turkey/Minnesota/1/1988(H7N9)
AHA11483 A/turkey/Minnesota/14135-2/2009(H7N9)
AHA57050 A/turkey/Minnesota/14659/2009(H7N9)
AHA57072 A/turkey/Minnesota/18421/2009(H7N9)
AHA11472 A/turkey/Minnesota/31676/2009(H7N9)
AHA11461 A/turkey/Minnesota/31900/2011(H7N9)
AHA11452 A/turkey/Minnesota/32710/2011(H7N9)
AAM19228 A/turkey/Minnesota/38429/1988(H7N9)
ACZ48625 A/turkey/Minnesota/38429/1988(H7N9)
AEK84760 A/wild bird/Korea/A14/2011(H7N9)
AEK84761 A/wild bird/Korea/A3/2011(H7N9)
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FIGURE 22-10

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AEK84763  A/wild bird/Korea/A9/2011(H7N9)                    ....K.....................................................K.....
AGG53377  A/wild duck/Korea/CSM42-1/2011(H7N9)               ..........................................................................
AGG53366  A/wild duck/Korea/CSM42-34/2011(H7N9)              ..........................................................................
AGG53432  A/wild duck/Korea/MHC35-41/2011(H7N9)              ..........................................................................
AGG53399  A/wild duck/Korea/MHC39-26/2011(H7N9)              ..........................................................................
AGG53476  A/wild duck/Korea/SH19-27/2010(H7N9)               ..........................................................................
AGG53487  A/wild duck/Korea/SH19-50/2010(H7N9)               ..........................................................................
AGG53520  A/wild duck/Korea/SH20-27/2008(H7N9)               ........................L.............T.....................
AEM98291  A/wild duck/Mongolia/1-241/2008(H7N9)              ..........................................................................
AGR49770  A/wild pigeon/Jiangsu/SD001/2013(H7N9)             ....K.....................................................................

 Positions from 121 till 180
          Consensus sequence                                 RESGGIDKEAMGFTYSGIRTNGATSACRRSGSSSFYAEMKWLLSNTDNAAFPQMTKSYKN
ACX53683  A/goose/Czech Republic/1848-K9/2009(H7N9)          .........T.....................-..............................
ADN34727  A/goose/Czech Republic/1848-T14/2009(H7N9)         .........T............X.......-..............................
CAY39406  A/Anas crecca/Spain/1460/2008(H7N9)                .........T.....................-..............................
AHZ39686  A/Anhui/DEWH72-01/2013(H7N9)                      ...............................-.............D............
AHZ39710  A/Anhui/DEWH72-03/2013(H7N9)                      ....................D..........-.............D............
AHZ39746  A/Anhui/DEWH72-06/2013(H7N9)                      ...............................-..............................
AHM24224  A/Beijing/3/2013(H7N9)                            ...............................-..............................
AHD25003  A/Guangdong/02/2013(H7N9)                         ....................A..........-..............................
AHH25185  A/Guangdong/04/2013(H7N9)                         ......E............A..........-..............................
AGI60301  A/Hangzhou/1/2013(H7N9)                           ...............................-..............................
AGK84857  A/Hangzhou/2/2013(H7N9)                           ...............................-.....................I........
AHF20528  A/Hong Kong/470129/2013(H7N9)                     ....................A..........-..............................
AGY41893  A/Huizhou/01/2013(H7N9)                          ....................A..........-..............................
AGO51387  A/Jiangsu/2/2013(H7N9)                           ...............................-..............................
AGJ73503  A/Nanjing/1/2013(H7N9)                           ...............................-..............................
AGR85026  A/Nanjing/2/2013(H7N9)                           ...............................-..............................
AGR84954  A/Nanjing/6/2013(H7N9)                           ...............................-..............................
AHK10800  A/Shanghai/01/2014(H7N9)                         ...............................-..............................
AGL44438  A/Shanghai/02/2013(H7N9)                         ...............................-..............................
AGL33692  A/Shanghai/4655T/2013(H7N9)                      ------------------------------------------------------------
AGL33693  A/Shanghai/4659T/2013(H7N9)                      ------------------------------------------------------------
AGI60292  A/Shanghai/4664T/2013(H7N9)                      ...............................-..............................
AGM53887  A/Shanghai/4701T/2013(H7N9)                      ------------------------------------------------------------
AGM53886  A/Shanghai/4842T/2013(H7N9)                      ------------------------------------------------------------
AGM53883  A/Shanghai/5083T/2013(H7N9)                      ------------------------------------------------------------
AGM53884  A/Shanghai/5180T/2013(H7N9)                      ------------------------------------------------------------
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FIGURE 22-11

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AGM53885 A/Shanghai/5240T/2013(H7N9)                                ------------------------------------------------------------
AGW82600 A/Shanghai/CN01/2013(H7N9)                                 ............................R-............................
AHF20568 A/Shanghai/CN02/2013(H7N9)                                 ............................-............................
AGW82612 A/Shanghai/JS01/2013(H7N9)                                 ............................-............................
AID70634 A/Shanghai/Mix1/2014(H7N9)                                 ............................-............................
AHJ57411 A/Shanghai/PD-01/2014(H7N9)                                ............................-............................
AHJ57418 A/Shanghai/PD-02/2014(H7N9)                                ............................-............................
AJJ96841 A/Shenzhen/SP139/2014(H7N9)                                .K.....................T.....-............................
AJJ91909 A/Shenzhen/SP26/2014(H7N9)                                 ..................A.......K...-............................
AJJ91945 A/Shenzhen/SP38/2014(H7N9)                                 ..................A...........-............................
AJJ91885 A/Shenzhen/SP4/2014(H7N9)                                  ..................A..V........-............................
AJJ91957 A/Shenzhen/SP44/2014(H7N9)                                 ..................A..T........-............................
AJJ91969 A/Shenzhen/SP48/2014(H7N9)                                 ............................-............................
AJJ95596 A/Shenzhen/SP58/2014(H7N9)                                 ..................A...........-............................
AJJ95632 A/Shenzhen/SP62/2014(H7N9)                                 ............................-.................T..........
AJJ95620 A/Shenzhen/SP75/2014(H7N9)                                 ......................S........-............................
AGR84942 A/Suzhou/5/2013(H7N9)                                      ............................-............................
AGL43637 A/Taiwan/1/2013(H7N9)                                      ............................-............................
AGL95088 A/Taiwan/S02076/2013(H7N9)                                 ............................-............................
AGL95098 A/Taiwan/T02081/2013(H7N9)                                 ............................-............................
AGN69474 A/Wuxi/1/2013(H7N9)                                        ............................-............................
AGN69462 A/Wuxi/2/2013(H7N9)                                        ......................S........-............................
AGR84990 A/Wuxi/3/2013(H7N9)                                        ............................-............................
AGR84978 A/Wuxi/4/2013(H7N9)                                        ............................-............................
AGO02477 A/Xuzhou/1/2013(H7N9)                                      ............................-............................
AGU02230 A/Zhejiang/DTID-ZJU05/2013(H7N9)                           ............................-............................
AGU02233 A/Zhejiang/DTID-ZJU08/2013(H7N9)                           ............................-............................
AHA11500 A/Zhejiang/DTID-ZJU10/2013(H7N9)                           ............................-............................
AIN76383 A/Zhejiang/LS01/2014(H7N9)                                 ......................T........-............................
AGR85002 A/Zhenjiang/1/2013(H7N9)                                   ............................-............................
ADK71137 A/blue-winged teal/Guatemala/CIP049-01/2008(H7N9)          .G.......S.....................-..S.........S............R.
ADK71148 A/blue-winged teal/Guatemala/CIP049-02/2008(H7N9)          .G.......S.....................-..........S............R.
ABS89409 A/blue-winged teal/Ohio/566/2006(H7N9)                     .G.......S.....................-..........S............R.
AJJ95048 A/chicken/Dongguan/1009/2014(H7N9)                         ..................A...........-............................
AJJ94134 A/chicken/Dongguan/1051/2014(H7N9)                         .K..........................-............................
AJJ94158 A/chicken/Dongguan/1075/2014(H7N9)                         ............................-............................
AJJ95572 A/chicken/Dongguan/1100/2014(H7N9)                         ......E.....................A...........-............................
AJJ94182 A/chicken/Dongguan/1177/2014(H7N9)                         ..........................K...-............................
AJJ95171 A/chicken/Dongguan/1314/2014(H7N9)                         ............................-............................
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FIGURE 22-12

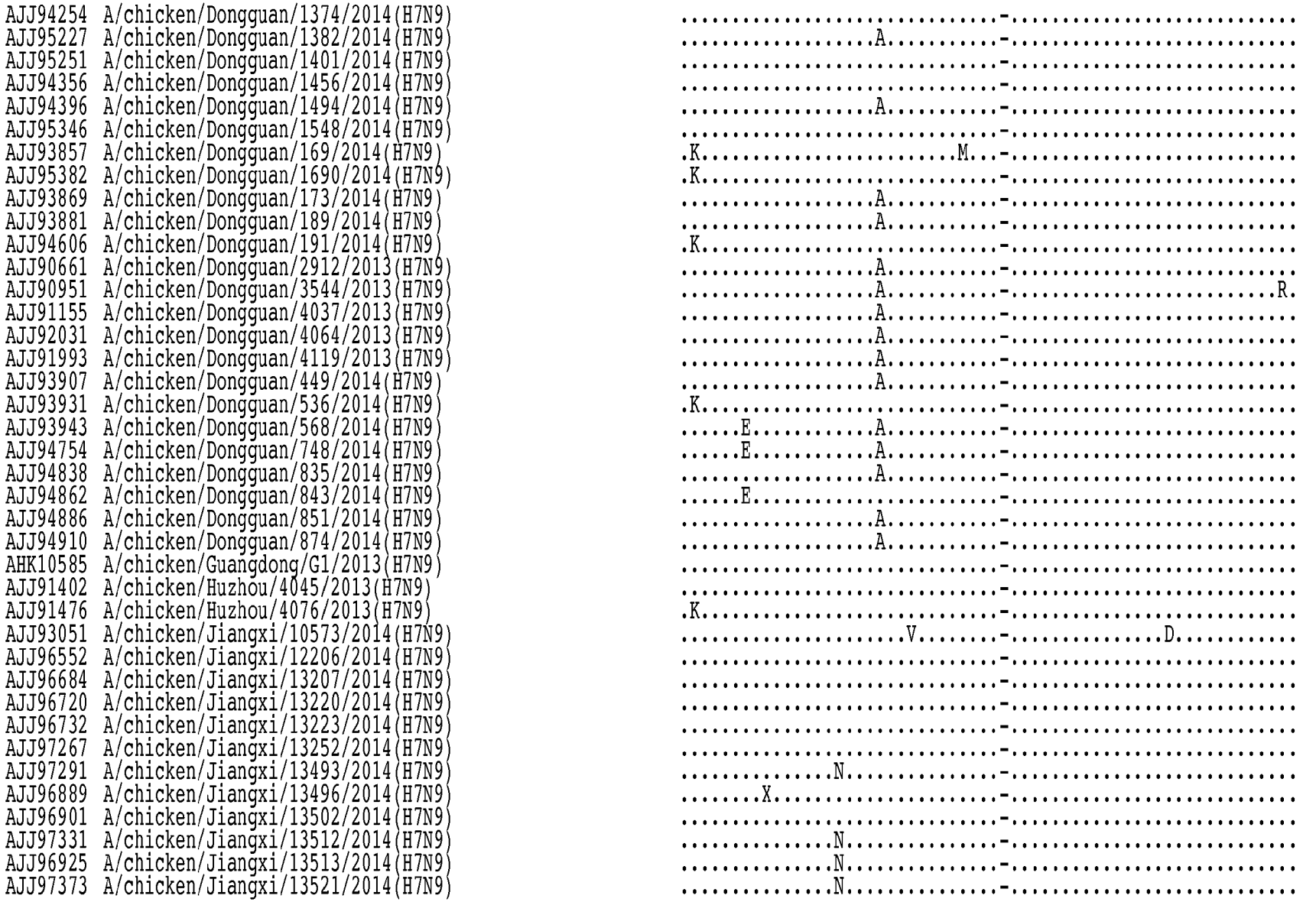

AJJ94254 A/chicken/Dongguan/1374/2014(H7N9)
AJJ95227 A/chicken/Dongguan/1382/2014(H7N9)
AJJ95251 A/chicken/Dongguan/1401/2014(H7N9)
AJJ94356 A/chicken/Dongguan/1456/2014(H7N9)
AJJ94396 A/chicken/Dongguan/1494/2014(H7N9)
AJJ95346 A/chicken/Dongguan/1548/2014(H7N9)
AJJ93857 A/chicken/Dongguan/169/2014(H7N9)
AJJ95382 A/chicken/Dongguan/1690/2014(H7N9)
AJJ93869 A/chicken/Dongguan/173/2014(H7N9)
AJJ93881 A/chicken/Dongguan/189/2014(H7N9)
AJJ94606 A/chicken/Dongguan/191/2014(H7N9)
AJJ90661 A/chicken/Dongguan/2912/2013(H7N9)
AJJ90951 A/chicken/Dongguan/3544/2013(H7N9)
AJJ91155 A/chicken/Dongguan/4037/2013(H7N9)
AJJ92031 A/chicken/Dongguan/4064/2013(H7N9)
AJJ91993 A/chicken/Dongguan/4119/2013(H7N9)
AJJ93907 A/chicken/Dongguan/449/2014(H7N9)
AJJ93931 A/chicken/Dongguan/536/2014(H7N9)
AJJ93943 A/chicken/Dongguan/568/2014(H7N9)
AJJ94754 A/chicken/Dongguan/748/2014(H7N9)
AJJ94838 A/chicken/Dongguan/835/2014(H7N9)
AJJ94862 A/chicken/Dongguan/843/2014(H7N9)
AJJ94886 A/chicken/Dongguan/851/2014(H7N9)
AJJ94910 A/chicken/Dongguan/874/2014(H7N9)
AHK10585 A/chicken/Guangdong/G1/2013(H7N9)
AJJ91402 A/chicken/Huzhou/4045/2013(H7N9)
AJJ91476 A/chicken/Huzhou/4076/2013(H7N9)
AJJ93051 A/chicken/Jiangxi/10573/2014(H7N9)
AJJ96552 A/chicken/Jiangxi/12206/2014(H7N9)
AJJ96684 A/chicken/Jiangxi/13207/2014(H7N9)
AJJ96720 A/chicken/Jiangxi/13220/2014(H7N9)
AJJ96732 A/chicken/Jiangxi/13223/2014(H7N9)
AJJ97267 A/chicken/Jiangxi/13252/2014(H7N9)
AJJ97291 A/chicken/Jiangxi/13493/2014(H7N9)
AJJ96889 A/chicken/Jiangxi/13496/2014(H7N9)
AJJ96901 A/chicken/Jiangxi/13502/2014(H7N9)
AJJ97331 A/chicken/Jiangxi/13512/2014(H7N9)
AJJ96925 A/chicken/Jiangxi/13513/2014(H7N9)
AJJ97373 A/chicken/Jiangxi/13521/2014(H7N9)
.....-.....
.....A.....-.....
.....-.....
.....-.....
.....A.....-.....
.....-.....
.K.....M...-.....
.K.....-.....
.....A.....-.....
.....A.....-.....
.K.....-.....
.....A.....-.....
.....A.....-.....R.
.....A.....-.....
.....A.....-.....
.....A.....-.....
.....A.....-.....
.K.....-.....
.....E.....A.....-.....
.....E.....A.....-.....
.....A.....-.....
.....E.....-.....
.....A.....-.....
.....A.....-.....
.....-.....
.....-.....
.K.....-.....
.....V.....-.....D.....
.....-.....
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.....-.....
.....-.....
.....N.....-.....
.....X.....-.....
.....-.....
.....N.....-.....
.....N.....-.....
.....N.....-.....

FIGURE 22-13

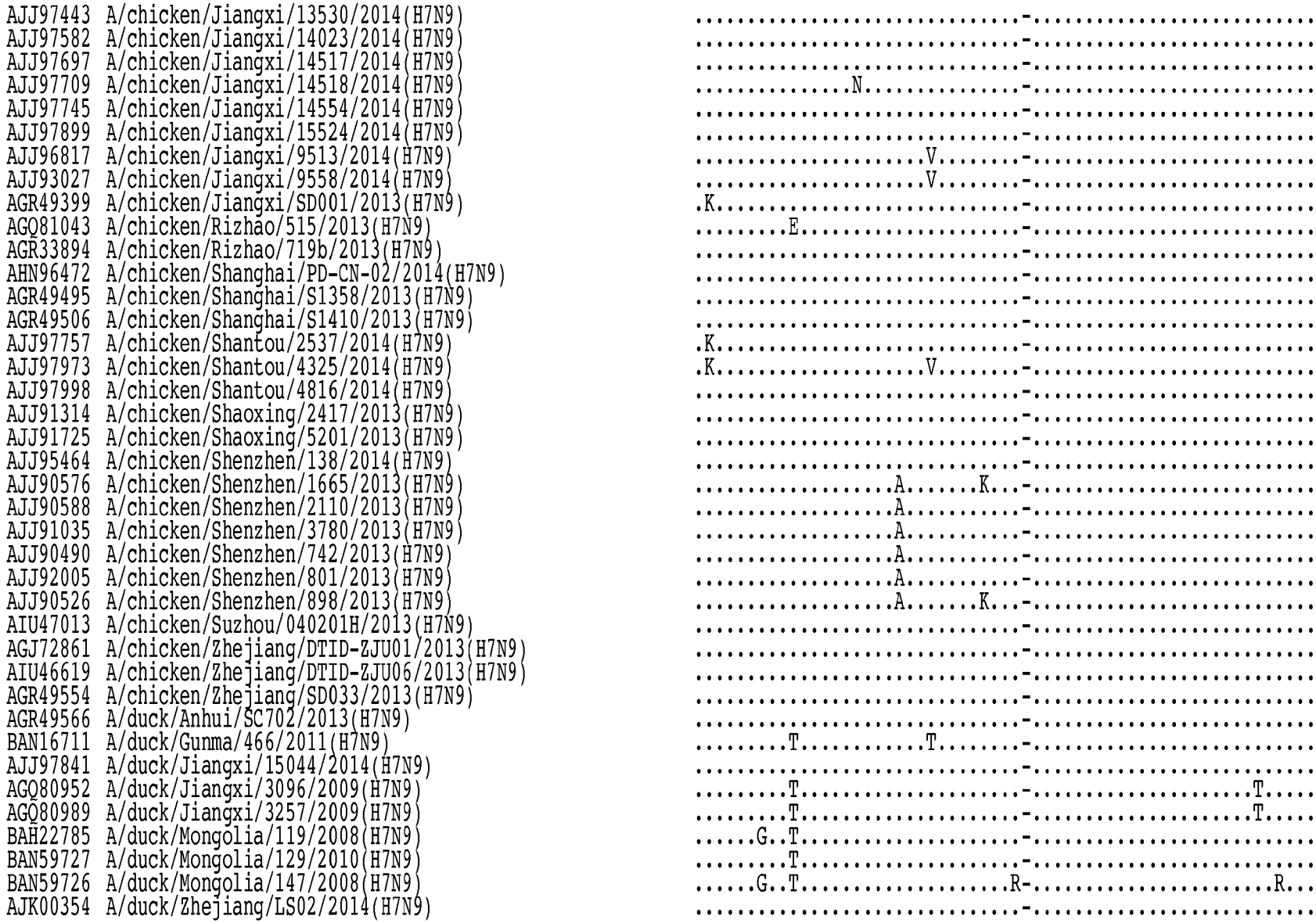

AJJ97443 A/chicken/Jiangxi/13530/2014(H7N9)
AJJ97582 A/chicken/Jiangxi/14023/2014(H7N9)
AJJ97697 A/chicken/Jiangxi/14517/2014(H7N9)
AJJ97709 A/chicken/Jiangxi/14518/2014(H7N9)
AJJ97745 A/chicken/Jiangxi/14554/2014(H7N9)
AJJ97899 A/chicken/Jiangxi/15524/2014(H7N9)
AJJ96817 A/chicken/Jiangxi/9513/2014(H7N9)
AJJ93027 A/chicken/Jiangxi/9558/2014(H7N9)
AGR49399 A/chicken/Jiangxi/SD001/2013(H7N9)
AGQ81043 A/chicken/Rizhao/515/2013(H7N9)
AGR33894 A/chicken/Rizhao/719b/2013(H7N9)
AHN96472 A/chicken/Shanghai/PD-CN-02/2014(H7N9)
AGR49495 A/chicken/Shanghai/S1358/2013(H7N9)
AGR49506 A/chicken/Shanghai/S1410/2013(H7N9)
AJJ97757 A/chicken/Shantou/2537/2014(H7N9)
AJJ97973 A/chicken/Shantou/4325/2014(H7N9)
AJJ97998 A/chicken/Shantou/4816/2014(H7N9)
AJJ91314 A/chicken/Shaoxing/2417/2013(H7N9)
AJJ91725 A/chicken/Shaoxing/5201/2013(H7N9)
AJJ95464 A/chicken/Shenzhen/138/2014(H7N9)
AJJ90576 A/chicken/Shenzhen/1665/2013(H7N9)
AJJ90588 A/chicken/Shenzhen/2110/2013(H7N9)
AJJ91035 A/chicken/Shenzhen/3780/2013(H7N9)
AJJ90490 A/chicken/Shenzhen/742/2013(H7N9)
AJJ92005 A/chicken/Shenzhen/801/2013(H7N9)
AJJ90526 A/chicken/Shenzhen/898/2013(H7N9)
AIU47013 A/chicken/Suzhou/040201H/2013(H7N9)
AGJ72861 A/chicken/Zhejiang/DTID-ZJU01/2013(H7N9)
AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013(H7N9)
AGR49554 A/chicken/Zhejiang/SD033/2013(H7N9)
AGR49566 A/duck/Anhui/SC702/2013(H7N9)
BAN16711 A/duck/Gunma/466/2011(H7N9)
AJJ97841 A/duck/Jiangxi/15044/2014(H7N9)
AGQ80952 A/duck/Jiangxi/3096/2009(H7N9)
AGQ80989 A/duck/Jiangxi/3257/2009(H7N9)
BAH22785 A/duck/Mongolia/119/2008(H7N9)
BAN59727 A/duck/Mongolia/129/2010(H7N9)
BAN59726 A/duck/Mongolia/147/2008(H7N9)
AJK00354 A/duck/Zhejiang/LS02/2014(H7N9)

FIGURE 22-14

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AFM09439  A/emperor goose/Alaska/44063-061/2006(H7N9)                .G.......S.....................-.............S............R.
AFV33947  A/goose/Nebraska/17097-4/2011(H7N9)                        .G.......S.....................-.............S............R.
AFV33945  A/guinea fowl/Nebraska/17096-1/2011(H7N9)                  .G.......S.....................-.............SN...........R.
AHA11441  A/guinea fowl/Nebraska/17096/2011(H7N9)                    .G.......S.....................-.............SN...........R.
AGR49722  A/homing pigeon/Jiangsu/SD184/2013(H7N9)                   ................E..............-..............................
AGL97639  A/mallard/Minnesota/AI09-3770/2009(H7N9)                  .G.......S.....................-.............S............R.
AHZ42537  A/mallard/Minnesota/AI09-3770/2009(H7N9)                  .G.......S.....................-.............S............R.
ADC29485  A/mallard/Spain/08.00991.3/2005(H7N9)                     ------------------------------------------------------------
AHZ41929  A/mallard/Sweden/1621/2002(H7N9)                          .........T.....................-..............................
AAY46211  A/mallard/Sweden/91/2002(H7N9)                            .........T.............P.......-..............................
AGY42258  A/mallard/Sweden/91/2002(H7N9)                            .........T.............X.......-..............................
AGE08098  A/northern shoverl/Mississippi/11OS145/2011(H7N9)         .G.......S.....................-.............S............R.
AGR49734  A/pigeon/Shanghai/S1069/2013(H7N9)                        ...............................-..............................
ACD03594  A/ruddy turnstone/DE/1538/2000(H7N9)                      .G.......S...................L.-.............S............R.
AFX85260  A/ruddy turnstone/Delaware Bay/220/1995(H7N9)             .G.......S.................I.L.-.............S............R.
AHZ42549  A/ruddy turnstone/Delaware/AI00-1538/2000(H7N9)           .G.......S...................L.-.............S............R.
AJJ94194  A/silkie chicken/Dongguan/1264/2014(H7N9)                 ...............................-..............................
AJJ94206  A/silkie chicken/Dongguan/1268/2014(H7N9)                 ..................A............-..............................
AJJ94344  A/silkie chicken/Dongguan/1451/2014(H7N9)                 .K.............................-..............................
AJJ95584  A/silkie chicken/Dongguan/1519/2014(H7N9)                 ...............................-..............................
AJJ93845  A/silkie chicken/Dongguan/157/2014(H7N9)                  ..................A............-..............................
AJJ90673  A/silkie chicken/Dongguan/3049/2013(H7N9)                 ..................A............-..............................
AJJ90795  A/silkie chicken/Dongguan/3281/2013(H7N9)                 ..................A............-..............................
AJJ90891  A/silkie chicken/Dongguan/3520/2013(H7N9)                 ..................A............-..............................
AJJ91264  A/silkie chicken/Dongguan/4129/2013(H7N9)                 ..................A............-..............................
AJJ93979  A/silkie chicken/Dongguan/656/2014(H7N9)                  ..................A............-..............................
AJJ94959  A/silkie chicken/Dongguan/967/2014(H7N9)                  ..................A.......X...-..............................
AJJ92967  A/silkie chicken/Jiangxi/9469/2014(H7N9)                  ....................V.........-..............................
AJJ97925  A/silkie chicken/Shantou/2050/2014(H7N9)                  .K.............................-..............................
AJJ90538  A/silkie chicken/Shenzhen/918/2013(H7N9)                  ..................A............-..............................
AEK84765  A/spot-billed duck/Korea/447/2011(H7N9)                   .........T.....................-..............................
AGW82588  A/tree sparrow/Shanghai/01/2013(H7N9)                     ...............................-..............................
ABI84694  A/turkey/Minnesota/1/1988(H7N9)                          .G.......S.....................-.............S............R.
AHA11483  A/turkey/Minnesota/14135-2/2009(H7N9)                    .G.......S.....................-.............SN...........R.
AHA57050  A/turkey/Minnesota/14659/2009(H7N9)                      .G.......S.....................-.............SN...........R.
AHA57072  A/turkey/Minnesota/18421/2009(H7N9)                      .G.......S.....................-.............SND..........R.
AHA11472  A/turkey/Minnesota/31676/2009(H7N9)                      .G.......S............E........-.............SN...........R.
AHA11461  A/turkey/Minnesota/31900/2011(H7N9)                      .G.......S..............T.....-.............SN...........R.
AHA11452  A/turkey/Minnesota/32710/2011(H7N9)                      .G.......S..............T.....-.............SN...........R.
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FIGURE 22-15

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AAM19228 A/turkey/Minnesota/38429/1988(H7N9)        .G.......S........I............-.............S............R.
ACZ48625 A/turkey/Minnesota/38429/1988(H7N9)        .G.......S.....................-.............S............R.
AEK84760 A/wild bird/Korea/A14/2011(H7N9)           ------------------------------------------------------------
AEK84761 A/wild bird/Korea/A3/2011(H7N9)            .........T...A.N..K...PPI...K..-.....K.........K............
AEK84763 A/wild bird/Korea/A9/2011(H7N9)            .........T...A....K...PPI...K..-.....K.........K............
AGG53377 A/wild duck/Korea/CSM42-1/2011(H7N9)       .........T..L.................-.............................
AGG53366 A/wild duck/Korea/CSM42-34/2011(H7N9)      .........T..L.................-.............................
AGG53432 A/wild duck/Korea/MHC35-41/2011(H7N9)      .........T....................-.............................
AGG53399 A/wild duck/Korea/MHC39-26/2011(H7N9)      .........T....................-.............................
AGG53476 A/wild duck/Korea/SH19-27/2010(H7N9)       .........T....................-.............................
AGG53487 A/wild duck/Korea/SH19-50/2010(H7N9)       .........T....................-.............................
AGG53520 A/wild duck/Korea/SH20-27/2008(H7N9)       ......E..T....................-.............................
AEM98291 A/wild duck/Mongolia/1-241/2008(H7N9)      .........T....................-.............................
AGR49770 A/wild pigeon/Jiangsu/SD001/2013(H7N9)     ..............................-.............................

Positions from 181 till 240
         Consensus sequence                         TRKSPALIVWGIHHSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARPQVNGLSGRI
ACX53683 A/goose/Czech Republic/1848-K9/2009(H7N9)  ...D....I......G..T.........S..I.......................Q....
ADN34727 A/goose/Czech Republic/1848-T14/2009(H7N9) ...D....I......G..T.........S..I.......................Q....
CAY39406 A/Anas crecca/Spain/1460/2008(H7N9)        ...D....I......G..T.........S..I.......................Q....
AHZ39686 A/Anhui/DEWH72-01/2013(H7N9)               ............................................................
AHZ39710 A/Anhui/DEWH72-03/2013(H7N9)               ............................................................
AHZ39746 A/Anhui/DEWH72-06/2013(H7N9)               ...........................................................E........
AHM24224 A/Beijing/3/2013(H7N9)                     ............................................................
AHD25003 A/Guangdong/02/2013(H7N9)                  ............................................................
AHH25185 A/Guangdong/04/2013(H7N9)                  ............................................................
AGI60301 A/Hangzhou/1/2013(H7N9)                    ...........................................................I....
AGK84857 A/Hangzhou/2/2013(H7N9)                    ............................................................
AHF20528 A/Hong Kong/470129/2013(H7N9)              ............................................................
AGY41893 A/Huizhou/01/2013(H7N9)                    ............................................................
AGO51387 A/Jiangsu/2/2013(H7N9)                     ............................................................
AGJ73503 A/Nanjing/1/2013(H7N9)                     ............................................................
AGR85026 A/Nanjing/2/2013(H7N9)                     ............................................................
AGR84954 A/Nanjing/6/2013(H7N9)                     ............................................................
AHK10800 A/Shanghai/01/2014(H7N9)                   ............................................................
AGL44438 A/Shanghai/02/2013(H7N9)                   ............................................................
AGL33692 A/Shanghai/4655T/2013(H7N9)                ------------------------------------------------------------
AGL33693 A/Shanghai/4659T/2013(H7N9)                ------------------------------------------------------------
AGI60292 A/Shanghai/4664T/2013(H7N9)                ............................................................
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FIGURE 22-16

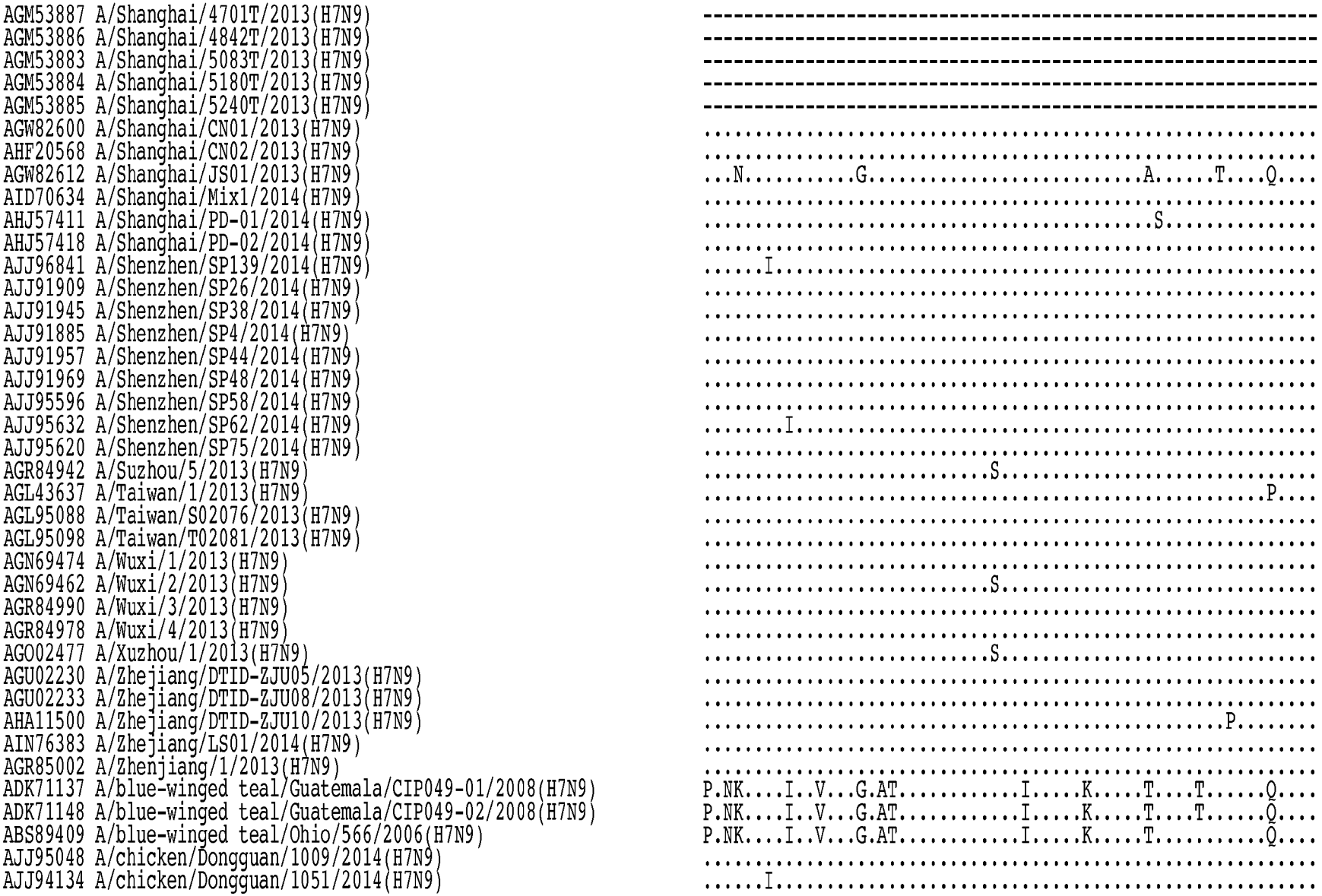

AGM53887 A/Shanghai/4701T/2013(H7N9)
AGM53886 A/Shanghai/4842T/2013(H7N9)
AGM53883 A/Shanghai/5083T/2013(H7N9)
AGM53884 A/Shanghai/5180T/2013(H7N9)
AGM53885 A/Shanghai/5240T/2013(H7N9)
AGW82600 A/Shanghai/CN01/2013(H7N9)
AHF20568 A/Shanghai/CN02/2013(H7N9)
AGW82612 A/Shanghai/JS01/2013(H7N9)
AID70634 A/Shanghai/Mix1/2014(H7N9)
AHJ57411 A/Shanghai/PD-01/2014(H7N9)
AHJ57418 A/Shanghai/PD-02/2014(H7N9)
AJJ96841 A/Shenzhen/SP139/2014(H7N9)
AJJ91909 A/Shenzhen/SP26/2014(H7N9)
AJJ91945 A/Shenzhen/SP38/2014(H7N9)
AJJ91885 A/Shenzhen/SP4/2014(H7N9)
AJJ91957 A/Shenzhen/SP44/2014(H7N9)
AJJ91969 A/Shenzhen/SP48/2014(H7N9)
AJJ95596 A/Shenzhen/SP58/2014(H7N9)
AJJ95632 A/Shenzhen/SP62/2014(H7N9)
AJJ95620 A/Shenzhen/SP75/2014(H7N9)
AGR84942 A/Suzhou/5/2013(H7N9)
AGL43637 A/Taiwan/1/2013(H7N9)
AGL95088 A/Taiwan/S02076/2013(H7N9)
AGL95098 A/Taiwan/T02081/2013(H7N9)
AGN69474 A/Wuxi/1/2013(H7N9)
AGN69462 A/Wuxi/2/2013(H7N9)
AGR84990 A/Wuxi/3/2013(H7N9)
AGR84978 A/Wuxi/4/2013(H7N9)
AGO02477 A/Xuzhou/1/2013(H7N9)
AGU02230 A/Zhejiang/DTID-ZJU05/2013(H7N9)
AGU02233 A/Zhejiang/DTID-ZJU08/2013(H7N9)
AHA11500 A/Zhejiang/DTID-ZJU10/2013(H7N9)
AIN76383 A/Zhejiang/LS01/2014(H7N9)
AGR85002 A/Zhenjiang/1/2013(H7N9)
ADK71137 A/blue-winged teal/Guatemala/CIP049-01/2008(H7N9)
ADK71148 A/blue-winged teal/Guatemala/CIP049-02/2008(H7N9)
ABS89409 A/blue-winged teal/Ohio/566/2006(H7N9)
AJJ95048 A/chicken/Dongguan/1009/2014(H7N9)
AJJ94134 A/chicken/Dongguan/1051/2014(H7N9)

FIGURE 22-17

```
AJJ94158 A/chicken/Dongguan/1075/2014(H7N9)  ..................................................
AJJ95572 A/chicken/Dongguan/1100/2014(H7N9)  ..................................................
AJJ94182 A/chicken/Dongguan/1177/2014(H7N9)  ..........I.......................................
AJJ95171 A/chicken/Dongguan/1314/2014(H7N9)  ..................................................
AJJ94254 A/chicken/Dongguan/1374/2014(H7N9)  ..................................................
AJJ95227 A/chicken/Dongguan/1382/2014(H7N9)  ..................................................
AJJ95251 A/chicken/Dongguan/1401/2014(H7N9)  ..................................................
AJJ94356 A/chicken/Dongguan/1456/2014(H7N9)  ..................................................
AJJ94396 A/chicken/Dongguan/1494/2014(H7N9)  ..................................................
AJJ95346 A/chicken/Dongguan/1548/2014(H7N9)  ..................................................
AJJ93857 A/chicken/Dongguan/169/2014(H7N9)   .....I............................................
AJJ95382 A/chicken/Dongguan/1690/2014(H7N9)  .....I............................................
AJJ93869 A/chicken/Dongguan/173/2014(H7N9)   ..................................................
AJJ93881 A/chicken/Dongguan/189/2014(H7N9)   ..................................................
AJJ94606 A/chicken/Dongguan/191/2014(H7N9)   .....I............................................
AJJ90661 A/chicken/Dongguan/2912/2013(H7N9)  ..................................................
AJJ90951 A/chicken/Dongguan/3544/2013(H7N9)  ..................................................
AJJ91155 A/chicken/Dongguan/4037/2013(H7N9)  ..................................................
AJJ92031 A/chicken/Dongguan/4064/2013(H7N9)  ..........E.......................................
AJJ91993 A/chicken/Dongguan/4119/2013(H7N9)  ..................................................
AJJ93907 A/chicken/Dongguan/449/2014(H7N9)   ..................................................
AJJ93931 A/chicken/Dongguan/536/2014(H7N9)   .....I............................................
AJJ93943 A/chicken/Dongguan/568/2014(H7N9)   ..................................................
AJJ94754 A/chicken/Dongguan/748/2014(H7N9)   ..................N...............................
AJJ94838 A/chicken/Dongguan/835/2014(H7N9)   ..................A...............................
AJJ94862 A/chicken/Dongguan/843/2014(H7N9)   ..................................................
AJJ94886 A/chicken/Dongguan/851/2014(H7N9)   ..................................................
AJJ94910 A/chicken/Dongguan/874/2014(H7N9)   ..................A...............................
AHK10585 A/chicken/Guangdong/G1/2013(H7N9)   ..................................................
AJJ91402 A/chicken/Huzhou/4045/2013(H7N9)    ..................................................
AJJ91476 A/chicken/Huzhou/4076/2013(H7N9)    ..................................................
AJJ93051 A/chicken/Jiangxi/10573/2014(H7N9)  ..................................................
AJJ96552 A/chicken/Jiangxi/12206/2014(H7N9)  ..................................................
AJJ96684 A/chicken/Jiangxi/13207/2014(H7N9)  ..................................................
AJJ96720 A/chicken/Jiangxi/13220/2014(H7N9)  ..................................................
AJJ96732 A/chicken/Jiangxi/13223/2014(H7N9)  ..................................................
AJJ97267 A/chicken/Jiangxi/13252/2014(H7N9)  ..................................................
AJJ97291 A/chicken/Jiangxi/13493/2014(H7N9)  .....I............................................
AJJ96889 A/chicken/Jiangxi/13496/2014(H7N9)  ..................................................
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FIGURE 22-18

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AJJ96901 A/chicken/Jiangxi/13502/2014(H7N9)  .....I.....T.....
AJJ97331 A/chicken/Jiangxi/13512/2014(H7N9)  .....I.....T.....
AJJ96925 A/chicken/Jiangxi/13513/2014(H7N9)  .....I.....T.....
AJJ97373 A/chicken/Jiangxi/13521/2014(H7N9)  .....I.....T.....
AJJ97443 A/chicken/Jiangxi/13530/2014(H7N9)  .....I.....T.....
AJJ97582 A/chicken/Jiangxi/14023/2014(H7N9)  .....I.....T.....
AJJ97697 A/chicken/Jiangxi/14517/2014(H7N9)  .....I.....T.....
AJJ97709 A/chicken/Jiangxi/14518/2014(H7N9)  .....I.....T.....
AJJ97745 A/chicken/Jiangxi/14554/2014(H7N9)  .....I.....T.....
AJJ97899 A/chicken/Jiangxi/15524/2014(H7N9)  .....I.....T.....
AJJ96817 A/chicken/Jiangxi/9513/2014(H7N9)   .....I.....T.....
AJJ93027 A/chicken/Jiangxi/9558/2014(H7N9)   .....I.....T.....
AGR49399 A/chicken/Jiangxi/SD001/2013(H7N9)  .....I.....T.....
AGQ81043 A/chicken/Rizhao/515/2013(H7N9)     .....I.....T.....
AGR33894 A/chicken/Rizhao/719b/2013(H7N9)    .....I.....T.....
AHN96472 A/chicken/Shanghai/PD-CN-02/2014(H7N9)  .....I.....T.....
AGR49495 A/chicken/Shanghai/S1358/2013(H7N9)  .....I.....T.....
AGR49506 A/chicken/Shanghai/S1410/2013(H7N9)  .....I.....T.....Q....
AJJ97757 A/chicken/Shantou/2537/2014(H7N9)   .....I.....T.....
AJJ97973 A/chicken/Shantou/4325/2014(H7N9)   .....I.....T.....
AJJ97998 A/chicken/Shantou/4816/2014(H7N9)   .....I.....T.....
AJJ91314 A/chicken/Shaoxing/2417/2013(H7N9)  .....I.....T.....P.....
AJJ91725 A/chicken/Shaoxing/5201/2013(H7N9)  .....I.....T.....
AJJ95464 A/chicken/Shenzhen/138/2014(H7N9)   .....I.....T.....
AJJ90576 A/chicken/Shenzhen/1665/2013(H7N9)  .....I.....T.....
AJJ90588 A/chicken/Shenzhen/2110/2013(H7N9)  .....I.....T.....
AJJ91035 A/chicken/Shenzhen/3780/2013(H7N9)  .....I.....T.....
AJJ90490 A/chicken/Shenzhen/742/2013(H7N9)   .....I.....T.....
AJJ92005 A/chicken/Shenzhen/801/2013(H7N9)   .....I.....T.....
AJJ90526 A/chicken/Shenzhen/898/2013(H7N9)   .....I.....T.....
AIU47013 A/chicken/Suzhou/040201H/2013(H7N9)  .....I.....T.....
AGJ72861 A/chicken/Zhejiang/DTID-ZJU01/2013(H7N9)  .....I.....T.....Q....
AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013(H7N9)  .....I.....T.....
AGR49554 A/chicken/Zhejiang/SD033/2013(H7N9)  .....I.....T.....
AGR49566 A/duck/Anhui/SC702/2013(H7N9)       .....I.....T.....
BAN16711 A/duck/Gunma/466/2011(H7N9)         .....I.....T.....
AJJ97841 A/duck/Jiangxi/15044/2014(H7N9)     .....I.....T.....Q....
AGQ80952 A/duck/Jiangxi/3096/2009(H7N9)      .....I.....T.....
AGQ80989 A/duck/Jiangxi/3257/2009(H7N9)      .....I.....T.....Q....
```

FIGURE 22-19

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BAH22785 A/duck/Mongolia/119/2008(H7N9)                          ...D....I......G..T...........I.....................Q....
BAN59727 A/duck/Mongolia/129/2010(H7N9)                          ...D....I......G..T.........S..I.....................Q....
BAN59726 A/duck/Mongolia/147/2008(H7N9)                          ...D....I......G..T...........I.....................Q....
AJK00354 A/duck/Zhejiang/LS02/2014(H7N9)                         ...D....I......G..T...........I..................L...Q....
AFM09439 A/emperor goose/Alaska/44063-061/2006(H7N9)             P.NK....I..V...G.AT...........I.....K...............Q....
AFV33947 A/goose/Nebraska/17097-4/2011(H7N9)                     P.NK.......V...A.AT.........S..I.....K.....T.........Q....
AFV33945 A/guinea fowl/Nebraska/17096-1/2011(H7N9)               P.NK.......V...G.AT.........S..I.....K.....T.........Q....
AHA11441 A/guinea fowl/Nebraska/17096/2011(H7N9)                 P.NK.......V...G.AT.........S..I.....K.....T.........Q....
AGR49722 A/homing pigeon/Jiangsu/SD184/2013(H7N9)                ..................................................Q....
AGL97639 A/mallard/Minnesota/AI09-3770/2009(H7N9)                P.NK....I..V...G.AT...........I.....K.....T.........Q....
AHZ42537 A/mallard/Minnesota/AI09-3770/2009(H7N9)                P.NK....I..V...G.AT...........I.....K.....T.........Q....
ADC29485 A/mallard/Spain/08.00991.3/2005(H7N9)                   --------------------------------------------------
AHZ41929 A/mallard/Sweden/1621/2002(H7N9)                        ..ND....I......G..T...........I.....................Q....
AAY46211 A/mallard/Sweden/91/2002(H7N9)                          ..ND....I......G..T...........I.....................Q....
AGY42258 A/mallard/Sweden/91/2002(H7N9)                          ..ND....I......G..T...........I.....................Q....
AGE08098 A/northern shoverl/Mississippi/11OS145/2011(H7N9)       P.NK....I..V...G.AT...........I.....K.....T.........Q....
AGR49734 A/pigeon/Shanghai/S1069/2013(H7N9)                      ..................................................
ACD03594 A/ruddy turnstone/DE/1538/2000(H7N9)                    P.NK....I..V...G.AN...........I.....K.....T.........Q....
AFX85260 A/ruddy turnstone/Delaware Bay/220/1995(H7N9)           P.NK....I..V...G.AN...........I.....K.....T.........Q....
AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000(H7N9)         P.NK....I..V...G.AN...........I.....K.....T.........Q....
AJJ94194 A/silkie chicken/Dongguan/1264/2014(H7N9)               ..................................................
AJJ94206 A/silkie chicken/Dongguan/1268/2014(H7N9)               ..................................................
AJJ94344 A/silkie chicken/Dongguan/1451/2014(H7N9)               ......I...........................................
AJJ95584 A/silkie chicken/Dongguan/1519/2014(H7N9)               ..................................................
AJJ93845 A/silkie chicken/Dongguan/157/2014(H7N9)                ..................................................
AJJ90673 A/silkie chicken/Dongguan/3049/2013(H7N9)               ..................................................
AJJ90795 A/silkie chicken/Dongguan/3281/2013(H7N9)               ..................................................
AJJ90891 A/silkie chicken/Dongguan/3520/2013(H7N9)               ...X..............................................
AJJ91264 A/silkie chicken/Dongguan/4129/2013(H7N9)               ..................................................
AJJ93979 A/silkie chicken/Dongguan/656/2014(H7N9)                ..................................................
AJJ94959 A/silkie chicken/Dongguan/967/2014(H7N9)                ..................................................
AJJ92967 A/silkie chicken/Jiangxi/9469/2014(H7N9)                ..................................................
AJJ97925 A/silkie chicken/Shantou/2050/2014(H7N9)                ......I...........................................
AJJ90538 A/silkie chicken/Shenzhen/918/2013(H7N9)                ..................................................
AEK84765 A/spot-billed duck/Korea/447/2011(H7N9)                 ..RD..........G..T.........S..I.....................Q....
AGW82588 A/tree sparrow/Shanghai/01/2013(H7N9)                   ..................................................
ABI84694 A/turkey/Minnesota/1/1988(H7N9)                         P.NK..........G..T...........I.....K.....T.........Q....
AHA11483 A/turkey/Minnesota/14135-2/2009(H7N9)                   P.DK....I..V...G.AT...........I.....K.....T.........Q....
AHA57050 A/turkey/Minnesota/14659/2009(H7N9)                     P.DK....I..V...G.AT...........I.....K.....T.........Q....
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FIGURE 22-20

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AHA57072 A/turkey/Minnesota/18421/2009(H7N9)
AHA11472 A/turkey/Minnesota/31676/2009(H7N9)
AHA11461 A/turkey/Minnesota/31900/2011(H7N9)
AHA11452 A/turkey/Minnesota/32710/2011(H7N9)
AAM19228 A/turkey/Minnesota/38429/1988(H7N9)
ACZ48625 A/turkey/Minnesota/38429/1988(H7N9)
AEK84760 A/wild bird/Korea/A14/2011(H7N9)
AEK84761 A/wild bird/Korea/A3/2011(H7N9)
AEK84763 A/wild bird/Korea/A9/2011(H7N9)
AGG53377 A/wild duck/Korea/CSM42-1/2011(H7N9)
AGG53366 A/wild duck/Korea/CSM42-34/2011(H7N9)
AGG53432 A/wild duck/Korea/MHC35-41/2011(H7N9)
AGG53399 A/wild duck/Korea/MHC39-26/2011(H7N9)
AGG53476 A/wild duck/Korea/SH19-27/2010(H7N9)
AGG53487 A/wild duck/Korea/SH19-50/2010(H7N9)
AGG53520 A/wild duck/Korea/SH20-27/2008(H7N9)
AEM98291 A/wild duck/Mongolia/1-241/2008(H7N9)
AGR49770 A/wild pigeon/Jiangsu/SD001/2013(H7N9)

Positions from 241 till 300
        Consensus sequence
ACX53683 A/goose/Czech Republic/1848-K9/2009(H7N9)
ADN34727 A/goose/Czech Republic/1848-T14/2009(H7N9)
CAY39406 A/Anas crecca/Spain/1460/2008(H7N9)
AHZ39686 A/Anhui/DEWH72-01/2013(H7N9)
AHZ39710 A/Anhui/DEWH72-03/2013(H7N9)
AHZ39746 A/Anhui/DEWH72-06/2013(H7N9)
AHM24224 A/Beijing/3/2013(H7N9)
AHD25003 A/Guangdong/02/2013(H7N9)
AHH25185 A/Guangdong/04/2013(H7N9)
AGI60301 A/Hangzhou/1/2013(H7N9)
AGK84857 A/Hangzhou/2/2013(H7N9)
AHF20528 A/Hong Kong/470129/2013(H7N9)
AGY41893 A/Huizhou/01/2013(H7N9)
AGO51387 A/Jiangsu/2/2013(H7N9)
AGJ73503 A/Nanjing/1/2013(H7N9)
AGR85026 A/Nanjing/2/2013(H7N9)
AGR84954 A/Nanjing/6/2013(H7N9)
AHK10800 A/Shanghai/01/2014(H7N9)
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P.DK....I..V...G.AT...........I.....K.....T..........Q....
P.DK....I..V...G.AT...........I.....K.....T..........Q....
P.NK.......V...G.AT.........S..I.....K.....T..........Q....
P.NK.......V...G.AT.........S..I.....K.....T..........Q....
P.NK..........G..T...........I..E..K.....T..........E....
P.NK..........G..T...........I.....K.....T..........Q....
------------------------------------------------------------
..RN..........G..TK......I.SN.I.....................Q....
I.RD..........G..TK..N...I.SN.I.....................Q....
..RD..........G.ST.........S..I.....................Q....
..RD..........G.ST.........S..I.....................Q....
..RD..........G..T.........S..I.....................Q....
..RD..........G..T.........S..I.....................Q....
..RD..........G..T.........S..I.....................Q....
..RD..........G..T.........S..I.....................Q....
...D....I.....G..T.........S..I.....................Q....
...D....I.....G..T.........S..I.....................Q....
...................................................Q....
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DFHWLMLNPNDTVTFSFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIIS
............................K............................
............................K............................
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.....................................................D......
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FIGURE 22-21

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AGL44438  A/Shanghai/02/2013(H7N9)                              ..................................................
AGL33692  A/Shanghai/4655T/2013(H7N9)                           --------------------------------------------------
AGL33693  A/Shanghai/4659T/2013(H7N9)                           --------------------------------------------------
AGI60292  A/Shanghai/4664T/2013(H7N9)                           ..................................................H.......
AGM53887  A/Shanghai/4701T/2013(H7N9)                           --------------------------------------------------
AGM53886  A/Shanghai/4842T/2013(H7N9)                           --------------------------------------------------
AGM53883  A/Shanghai/5083T/2013(H7N9)                           --------------------------------------------------
AGM53884  A/Shanghai/5180T/2013(H7N9)                           --------------------------------------------------
AGM53885  A/Shanghai/5240T/2013(H7N9)                           --------------------------------------------------
AGW82600  A/Shanghai/CN01/2013(H7N9)                            ..................................................M.
AHF20568  A/Shanghai/CN02/2013(H7N9)                            ..................................................M.
AGW82612  A/Shanghai/JS01/2013(H7N9)                            ..........................................D.......
AID70634  A/Shanghai/Mix1/2014(H7N9)                           ..................................................
AHJ57411  A/Shanghai/PD-01/2014(H7N9)                          ..........................................K.......
AHJ57418  A/Shanghai/PD-02/2014(H7N9)                          ........................K.........................
AJJ96841  A/Shenzhen/SP139/2014(H7N9)                         ......................C.................D.......
AJJ91909  A/Shenzhen/SP26/2014(H7N9)                           ..................................................
AJJ91945  A/Shenzhen/SP38/2014(H7N9)                           ..................................................
AJJ91885  A/Shenzhen/SP4/2014(H7N9)                            ..................................................
AJJ91957  A/Shenzhen/SP44/2014(H7N9)                           ..................................................
AJJ91969  A/Shenzhen/SP48/2014(H7N9)                           ..................................................
AJJ95596  A/Shenzhen/SP58/2014(H7N9)                           ..................................................
AJJ95632  A/Shenzhen/SP62/2014(H7N9)                           ..................................................
AJJ95620  A/Shenzhen/SP75/2014(H7N9)                           ..................................................
AGR84942  A/Suzhou/5/2013(H7N9)                                ..................................................
AGL43637  A/Taiwan/1/2013(H7N9)                                ..................................................N
AGL95088  A/Taiwan/S02076/2013(H7N9)                           ..................................................
AGL95098  A/Taiwan/T02081/2013(H7N9)                           ..................................................
AGN69474  A/Wuxi/1/2013(H7N9)                                  ..................................................
AGN69462  A/Wuxi/2/2013(H7N9)                                  ..................................................
AGR84990  A/Wuxi/3/2013(H7N9)                                  ..................................................
AGR84978  A/Wuxi/4/2013(H7N9)                                  ..................................................
AGO02477  A/Xuzhou/1/2013(H7N9)                                ..................................................
AGU02230  A/Zhejiang/DTID-ZJU05/2013(H7N9)                      ..................................................
AGU02233  A/Zhejiang/DTID-ZJU08/2013(H7N9)                      ..................................................
AHA11500  A/Zhejiang/DTID-ZJU10/2013(H7N9)                      ..................................................
AIN76383  A/Zhejiang/LS01/2014(H7N9)                           ..................................................
AGR85002  A/Zhenjiang/1/2013(H7N9)                             ..................................................
ADK71137  A/blue-winged teal/Guatemala/CIP049-01/2008(H7N9)     .....L.D.......T.................L....D.PL.SG.....F......V.
```

FIGURE 22-22

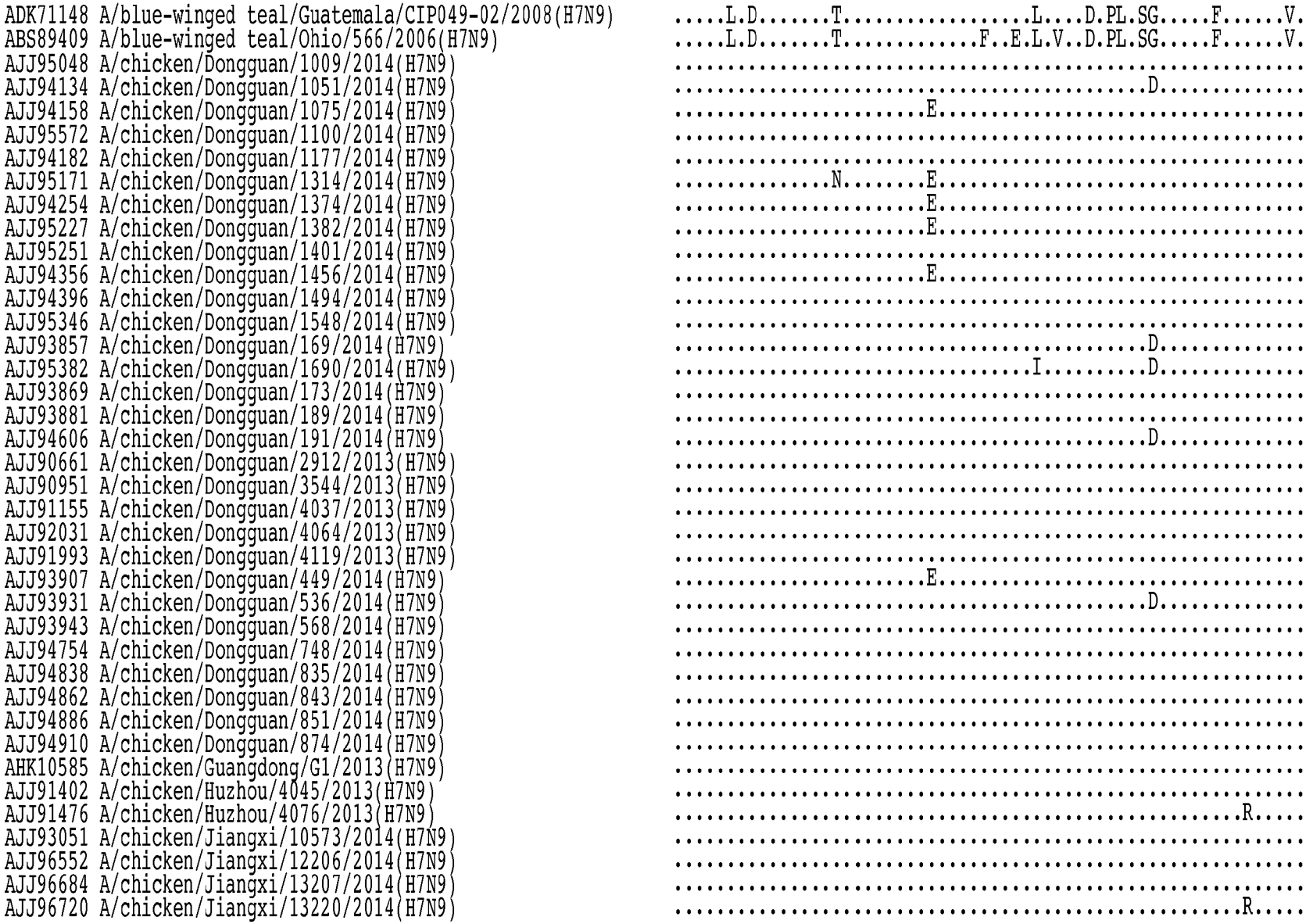

ADK71148 A/blue-winged teal/Guatemala/CIP049-02/2008(H7N9) .....L.D.....T.....L....D.PL.SG.....F.....V.
ABS89409 A/blue-winged teal/Ohio/566/2006(H7N9) .....L.D.....T.....F..E.L.V..D.PL.SG.....F.....V.
AJJ95048 A/chicken/Dongguan/1009/2014(H7N9) .....
AJJ94134 A/chicken/Dongguan/1051/2014(H7N9) .....D.....
AJJ94158 A/chicken/Dongguan/1075/2014(H7N9) .....E.....
AJJ95572 A/chicken/Dongguan/1100/2014(H7N9) .....
AJJ94182 A/chicken/Dongguan/1177/2014(H7N9) .....
AJJ95171 A/chicken/Dongguan/1314/2014(H7N9) .....N.....E.....
AJJ94254 A/chicken/Dongguan/1374/2014(H7N9) .....E.....
AJJ95227 A/chicken/Dongguan/1382/2014(H7N9) .....E.....
AJJ95251 A/chicken/Dongguan/1401/2014(H7N9) .....
AJJ94356 A/chicken/Dongguan/1456/2014(H7N9) .....E.....
AJJ94396 A/chicken/Dongguan/1494/2014(H7N9) .....
AJJ95346 A/chicken/Dongguan/1548/2014(H7N9) .....
AJJ93857 A/chicken/Dongguan/169/2014(H7N9) .....D.....
AJJ95382 A/chicken/Dongguan/1690/2014(H7N9) .....I.....D.....
AJJ93869 A/chicken/Dongguan/173/2014(H7N9) .....
AJJ93881 A/chicken/Dongguan/189/2014(H7N9) .....
AJJ94606 A/chicken/Dongguan/191/2014(H7N9) .....D.....
AJJ90661 A/chicken/Dongguan/2912/2013(H7N9) .....
AJJ90951 A/chicken/Dongguan/3544/2013(H7N9) .....
AJJ91155 A/chicken/Dongguan/4037/2013(H7N9) .....
AJJ92031 A/chicken/Dongguan/4064/2013(H7N9) .....
AJJ91993 A/chicken/Dongguan/4119/2013(H7N9) .....
AJJ93907 A/chicken/Dongguan/449/2014(H7N9) .....E.....
AJJ93931 A/chicken/Dongguan/536/2014(H7N9) .....D.....
AJJ93943 A/chicken/Dongguan/568/2014(H7N9) .....
AJJ94754 A/chicken/Dongguan/748/2014(H7N9) .....
AJJ94838 A/chicken/Dongguan/835/2014(H7N9) .....
AJJ94862 A/chicken/Dongguan/843/2014(H7N9) .....
AJJ94886 A/chicken/Dongguan/851/2014(H7N9) .....
AJJ94910 A/chicken/Dongguan/874/2014(H7N9) .....
AHK10585 A/chicken/Guangdong/G1/2013(H7N9) .....
AJJ91402 A/chicken/Huzhou/4045/2013(H7N9) .....
AJJ91476 A/chicken/Huzhou/4076/2013(H7N9) .....R.....
AJJ93051 A/chicken/Jiangxi/10573/2014(H7N9) .....
AJJ96552 A/chicken/Jiangxi/12206/2014(H7N9) .....
AJJ96684 A/chicken/Jiangxi/13207/2014(H7N9) .....
AJJ96720 A/chicken/Jiangxi/13220/2014(H7N9) .....R.....

FIGURE 22-23

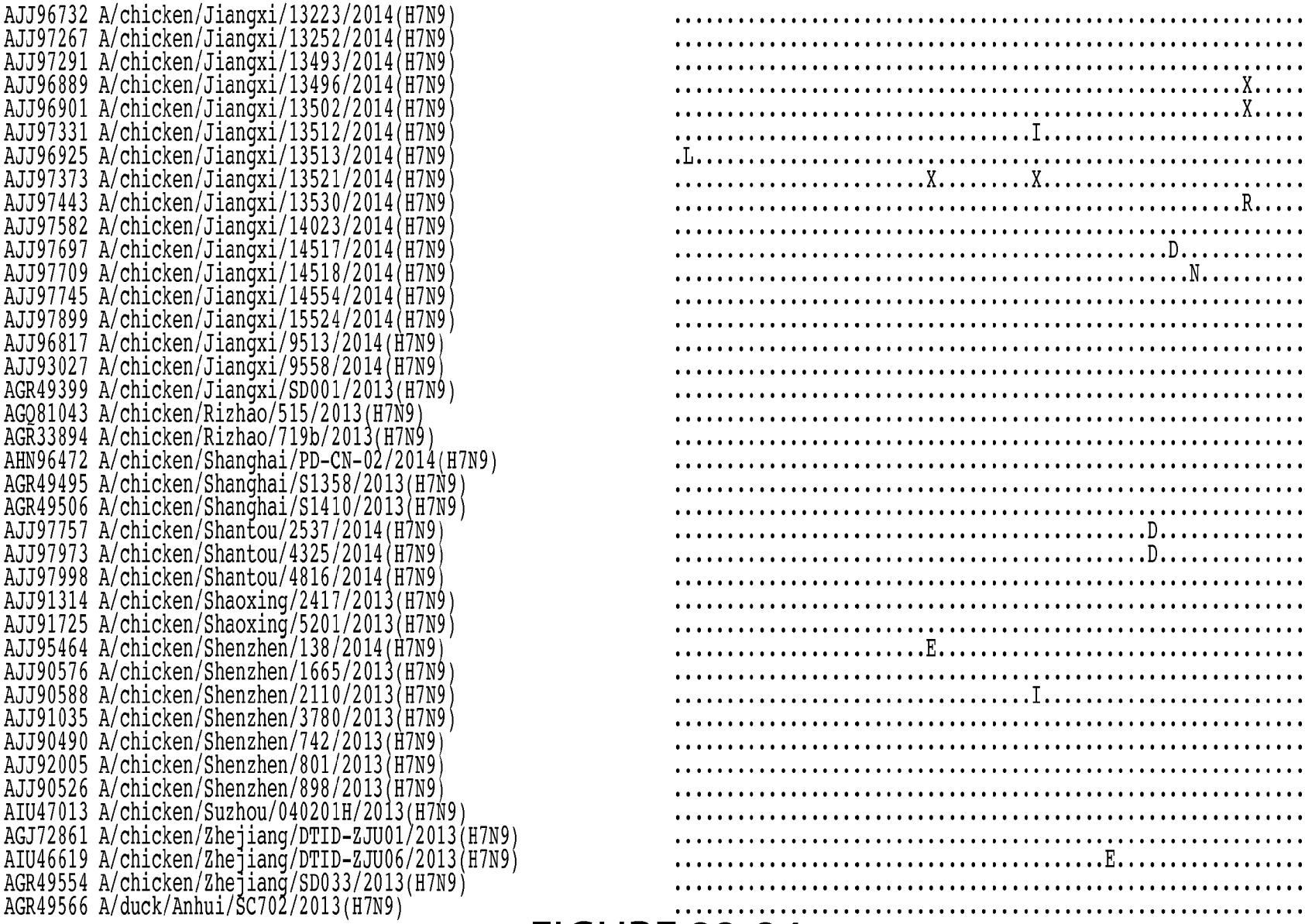

AJJ96732 A/chicken/Jiangxi/13223/2014(H7N9)
AJJ97267 A/chicken/Jiangxi/13252/2014(H7N9)
AJJ97291 A/chicken/Jiangxi/13493/2014(H7N9)
AJJ96889 A/chicken/Jiangxi/13496/2014(H7N9)
AJJ96901 A/chicken/Jiangxi/13502/2014(H7N9)
AJJ97331 A/chicken/Jiangxi/13512/2014(H7N9)
AJJ96925 A/chicken/Jiangxi/13513/2014(H7N9)
AJJ97373 A/chicken/Jiangxi/13521/2014(H7N9)
AJJ97443 A/chicken/Jiangxi/13530/2014(H7N9)
AJJ97582 A/chicken/Jiangxi/14023/2014(H7N9)
AJJ97697 A/chicken/Jiangxi/14517/2014(H7N9)
AJJ97709 A/chicken/Jiangxi/14518/2014(H7N9)
AJJ97745 A/chicken/Jiangxi/14554/2014(H7N9)
AJJ97899 A/chicken/Jiangxi/15524/2014(H7N9)
AJJ96817 A/chicken/Jiangxi/9513/2014(H7N9)
AJJ93027 A/chicken/Jiangxi/9558/2014(H7N9)
AGR49399 A/chicken/Jiangxi/SD001/2013(H7N9)
AGQ81043 A/chicken/Rizhao/515/2013(H7N9)
AGR33894 A/chicken/Rizhao/719b/2013(H7N9)
AHN96472 A/chicken/Shanghai/PD-CN-02/2014(H7N9)
AGR49495 A/chicken/Shanghai/S1358/2013(H7N9)
AGR49506 A/chicken/Shanghai/S1410/2013(H7N9)
AJJ97757 A/chicken/Shantou/2537/2014(H7N9)
AJJ97973 A/chicken/Shantou/4325/2014(H7N9)
AJJ97998 A/chicken/Shantou/4816/2014(H7N9)
AJJ91314 A/chicken/Shaoxing/2417/2013(H7N9)
AJJ91725 A/chicken/Shaoxing/5201/2013(H7N9)
AJJ95464 A/chicken/Shenzhen/138/2014(H7N9)
AJJ90576 A/chicken/Shenzhen/1665/2013(H7N9)
AJJ90588 A/chicken/Shenzhen/2110/2013(H7N9)
AJJ91035 A/chicken/Shenzhen/3780/2013(H7N9)
AJJ90490 A/chicken/Shenzhen/742/2013(H7N9)
AJJ92005 A/chicken/Shenzhen/801/2013(H7N9)
AJJ90526 A/chicken/Shenzhen/898/2013(H7N9)
AIU47013 A/chicken/Suzhou/040201H/2013(H7N9)
AGJ72861 A/chicken/Zhejiang/DTID-ZJU01/2013(H7N9)
AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013(H7N9)
AGR49554 A/chicken/Zhejiang/SD033/2013(H7N9)
AGR49566 A/duck/Anhui/SC702/2013(H7N9)

FIGURE 22-24

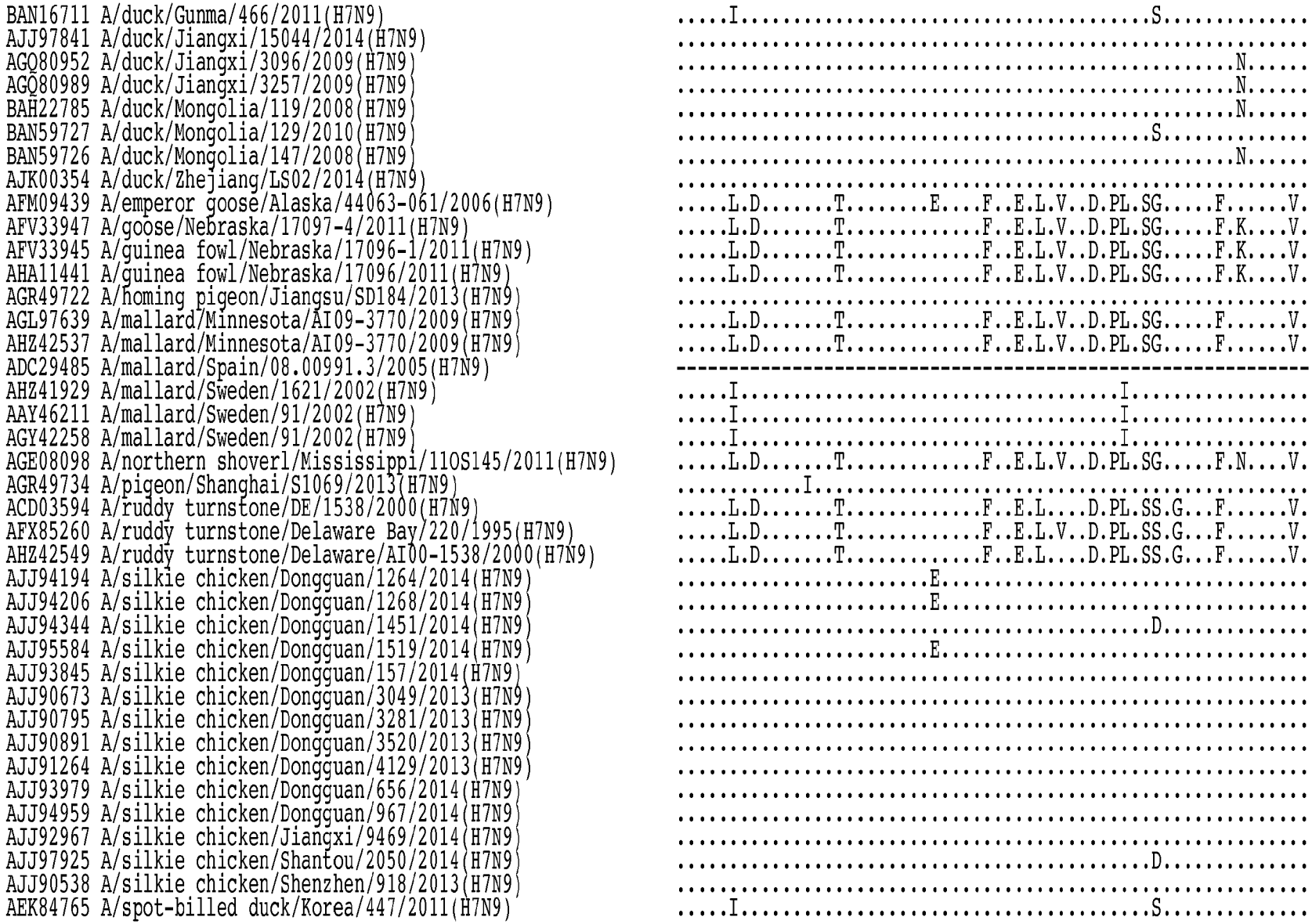

BAN16711 A/duck/Gunma/466/2011(H7N9)
AJJ97841 A/duck/Jiangxi/15044/2014(H7N9)
AGQ80952 A/duck/Jiangxi/3096/2009(H7N9)
AGQ80989 A/duck/Jiangxi/3257/2009(H7N9)
BAH22785 A/duck/Mongolia/119/2008(H7N9)
BAN59727 A/duck/Mongolia/129/2010(H7N9)
BAN59726 A/duck/Mongolia/147/2008(H7N9)
AJK00354 A/duck/Zhejiang/LS02/2014(H7N9)
AFM09439 A/emperor goose/Alaska/44063-061/2006(H7N9)
AFV33947 A/goose/Nebraska/17097-4/2011(H7N9)
AFV33945 A/guinea fowl/Nebraska/17096-1/2011(H7N9)
AHA11441 A/guinea fowl/Nebraska/17096/2011(H7N9)
AGR49722 A/homing pigeon/Jiangsu/SD184/2013(H7N9)
AGL97639 A/mallard/Minnesota/AI09-3770/2009(H7N9)
AHZ42537 A/mallard/Minnesota/AI09-3770/2009(H7N9)
ADC29485 A/mallard/Spain/08.00991.3/2005(H7N9)
AHZ41929 A/mallard/Sweden/1621/2002(H7N9)
AAY46211 A/mallard/Sweden/91/2002(H7N9)
AGY42258 A/mallard/Sweden/91/2002(H7N9)
AGE08098 A/northern shoverl/Mississippi/11OS145/2011(H7N9)
AGR49734 A/pigeon/Shanghai/S1069/2013(H7N9)
ACD03594 A/ruddy turnstone/DE/1538/2000(H7N9)
AFX85260 A/ruddy turnstone/Delaware Bay/220/1995(H7N9)
AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000(H7N9)
AJJ94194 A/silkie chicken/Dongguan/1264/2014(H7N9)
AJJ94206 A/silkie chicken/Dongguan/1268/2014(H7N9)
AJJ94344 A/silkie chicken/Dongguan/1451/2014(H7N9)
AJJ95584 A/silkie chicken/Dongguan/1519/2014(H7N9)
AJJ93845 A/silkie chicken/Dongguan/157/2014(H7N9)
AJJ90673 A/silkie chicken/Dongguan/3049/2013(H7N9)
AJJ90795 A/silkie chicken/Dongguan/3281/2013(H7N9)
AJJ90891 A/silkie chicken/Dongguan/3520/2013(H7N9)
AJJ91264 A/silkie chicken/Dongguan/4129/2013(H7N9)
AJJ93979 A/silkie chicken/Dongguan/656/2014(H7N9)
AJJ94959 A/silkie chicken/Dongguan/967/2014(H7N9)
AJJ92967 A/silkie chicken/Jiangxi/9469/2014(H7N9)
AJJ97925 A/silkie chicken/Shantou/2050/2014(H7N9)
AJJ90538 A/silkie chicken/Shenzhen/918/2013(H7N9)
AEK84765 A/spot-billed duck/Korea/447/2011(H7N9)

FIGURE 22-25

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AGW82588 A/tree sparrow/Shanghai/01/2013(H7N9)
ABI84694 A/turkey/Minnesota/1/1988(H7N9)
AHA11483 A/turkey/Minnesota/14135-2/2009(H7N9)
AHA57050 A/turkey/Minnesota/14659/2009(H7N9)
AHA57072 A/turkey/Minnesota/18421/2009(H7N9)
AHA11472 A/turkey/Minnesota/31676/2009(H7N9)
AHA11461 A/turkey/Minnesota/31900/2011(H7N9)
AHA11452 A/turkey/Minnesota/32710/2011(H7N9)
AAM19228 A/turkey/Minnesota/38429/1988(H7N9)
ACZ48625 A/turkey/Minnesota/38429/1988(H7N9)
AEK84760 A/wild bird/Korea/A14/2011(H7N9)
AEK84761 A/wild bird/Korea/A3/2011(H7N9)
AEK84763 A/wild bird/Korea/A9/2011(H7N9)
AGG53377 A/wild duck/Korea/CSM42-1/2011(H7N9)
AGG53366 A/wild duck/Korea/CSM42-34/2011(H7N9)
AGG53432 A/wild duck/Korea/MHC35-41/2011(H7N9)
AGG53399 A/wild duck/Korea/MHC39-26/2011(H7N9)
AGG53476 A/wild duck/Korea/SH19-27/2010(H7N9)
AGG53487 A/wild duck/Korea/SH19-50/2010(H7N9)
AGG53520 A/wild duck/Korea/SH20-27/2008(H7N9)
AEM98291 A/wild duck/Mongolia/1-241/2008(H7N9)
AGR49770 A/wild pigeon/Jiangsu/SD001/2013(H7N9)

Positions from 301 till 360
        Consensus sequence
ACX53683 A/goose/Czech Republic/1848-K9/2009(H7N9)
ADN34727 A/goose/Czech Republic/1848-T14/2009(H7N9)
CAY39406 A/Anas crecca/Spain/1460/2008(H7N9)
AHZ39686 A/Anhui/DEWH72-01/2013(H7N9)
AHZ39710 A/Anhui/DEWH72-03/2013(H7N9)
AHZ39746 A/Anhui/DEWH72-06/2013(H7N9)
AHM24224 A/Beijing/3/2013(H7N9)
AHD25003 A/Guangdong/02/2013(H7N9)
AHH25185 A/Guangdong/04/2013(H7N9)
AGI60301 A/Hangzhou/1/2013(H7N9)
AGK84857 A/Hangzhou/2/2013(H7N9)
AHF20528 A/Hong Kong/470129/2013(H7N9)
AGY41893 A/Huizhou/01/2013(H7N9)
AGO51387 A/Jiangsu/2/2013(H7N9)
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....................................................
....ML.D.......T.............FK.E.L.V..D.PL.SS.G...F......V.
.....L.D.......T.............F..E.L.V..D.PL.SG.....F......V.
.....L.D.......T.............F..E.L.V..D.PL.SG.....F......V.
.....L.D.......T.............F..E.L.V..D.PL.SG.....F......V.
.....L.D.......T.............F..E.L.V..D.PL.SG.....F......V.
.....L.D.......T.............F..E.L.V..D.PL.SG.....F.K....V.
.....L.D.......T.............F..E.L.V..D.PL.SG.....F.K....V.
....ML.D.......T.............FK.E.L.V..D.PL.SS.G...F......V.
....ML.D.......T.............FK.E.L.V..D.PL.SS.G...F......V.
----------------P.........................S.............
.....I................P.....................S.............
.....I.........I..............I.............S.............
.....I.....................................S.............
.....I.....................................S.............
.....I.....................................S.............
.....I.....................................S.............
.....I.....................................S.............
.....I.....................................S.............
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....................................................S...
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NLPFQNIDSRAVGKCPRYVKQRSLLLATGMKNVPEIPKGRGLFGAIAGFIENGWEGLIDG
.......N.............E..M..........L.....................
.......N.............E..M..........L.....................
.......N.............E..M.....................................
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S...................................................
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FIGURE 22-26

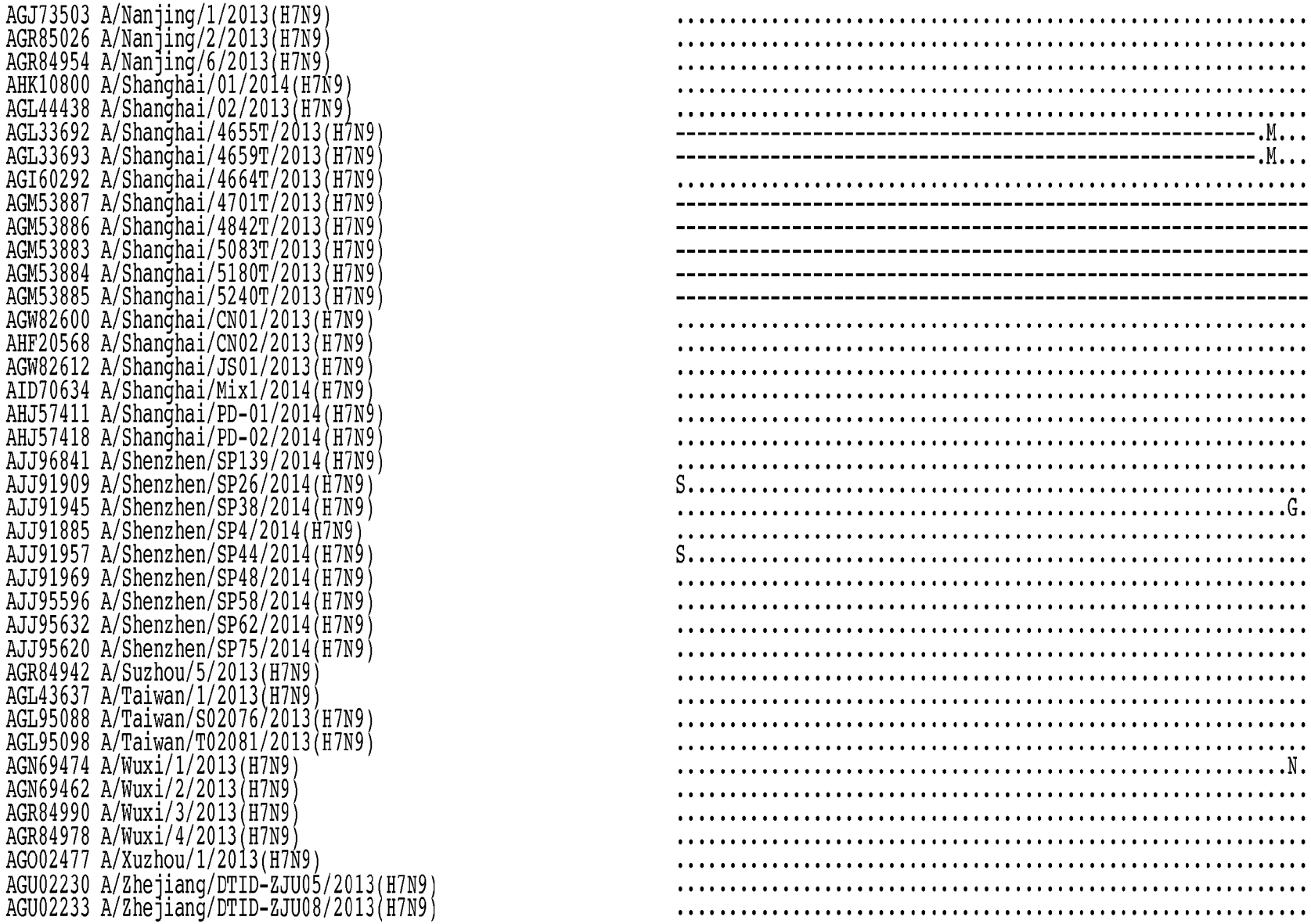

AGJ73503 A/Nanjing/1/2013(H7N9)
AGR85026 A/Nanjing/2/2013(H7N9)
AGR84954 A/Nanjing/6/2013(H7N9)
AHK10800 A/Shanghai/01/2014(H7N9)
AGL44438 A/Shanghai/02/2013(H7N9)
AGL33692 A/Shanghai/4655T/2013(H7N9)
AGL33693 A/Shanghai/4659T/2013(H7N9)
AGI60292 A/Shanghai/4664T/2013(H7N9)
AGM53887 A/Shanghai/4701T/2013(H7N9)
AGM53886 A/Shanghai/4842T/2013(H7N9)
AGM53883 A/Shanghai/5083T/2013(H7N9)
AGM53884 A/Shanghai/5180T/2013(H7N9)
AGM53885 A/Shanghai/5240T/2013(H7N9)
AGW82600 A/Shanghai/CN01/2013(H7N9)
AHF20568 A/Shanghai/CN02/2013(H7N9)
AGW82612 A/Shanghai/JS01/2013(H7N9)
AID70634 A/Shanghai/Mix1/2014(H7N9)
AHJ57411 A/Shanghai/PD-01/2014(H7N9)
AHJ57418 A/Shanghai/PD-02/2014(H7N9)
AJJ96841 A/Shenzhen/SP139/2014(H7N9)
AJJ91909 A/Shenzhen/SP26/2014(H7N9)
AJJ91945 A/Shenzhen/SP38/2014(H7N9)
AJJ91885 A/Shenzhen/SP4/2014(H7N9)
AJJ91957 A/Shenzhen/SP44/2014(H7N9)
AJJ91969 A/Shenzhen/SP48/2014(H7N9)
AJJ95596 A/Shenzhen/SP58/2014(H7N9)
AJJ95632 A/Shenzhen/SP62/2014(H7N9)
AJJ95620 A/Shenzhen/SP75/2014(H7N9)
AGR84942 A/Suzhou/5/2013(H7N9)
AGL43637 A/Taiwan/1/2013(H7N9)
AGL95088 A/Taiwan/S02076/2013(H7N9)
AGL95098 A/Taiwan/T02081/2013(H7N9)
AGN69474 A/Wuxi/1/2013(H7N9)
AGN69462 A/Wuxi/2/2013(H7N9)
AGR84990 A/Wuxi/3/2013(H7N9)
AGR84978 A/Wuxi/4/2013(H7N9)
AGO02477 A/Xuzhou/1/2013(H7N9)
AGU02230 A/Zhejiang/DTID-ZJU05/2013(H7N9)
AGU02233 A/Zhejiang/DTID-ZJU08/2013(H7N9)

FIGURE 22-27

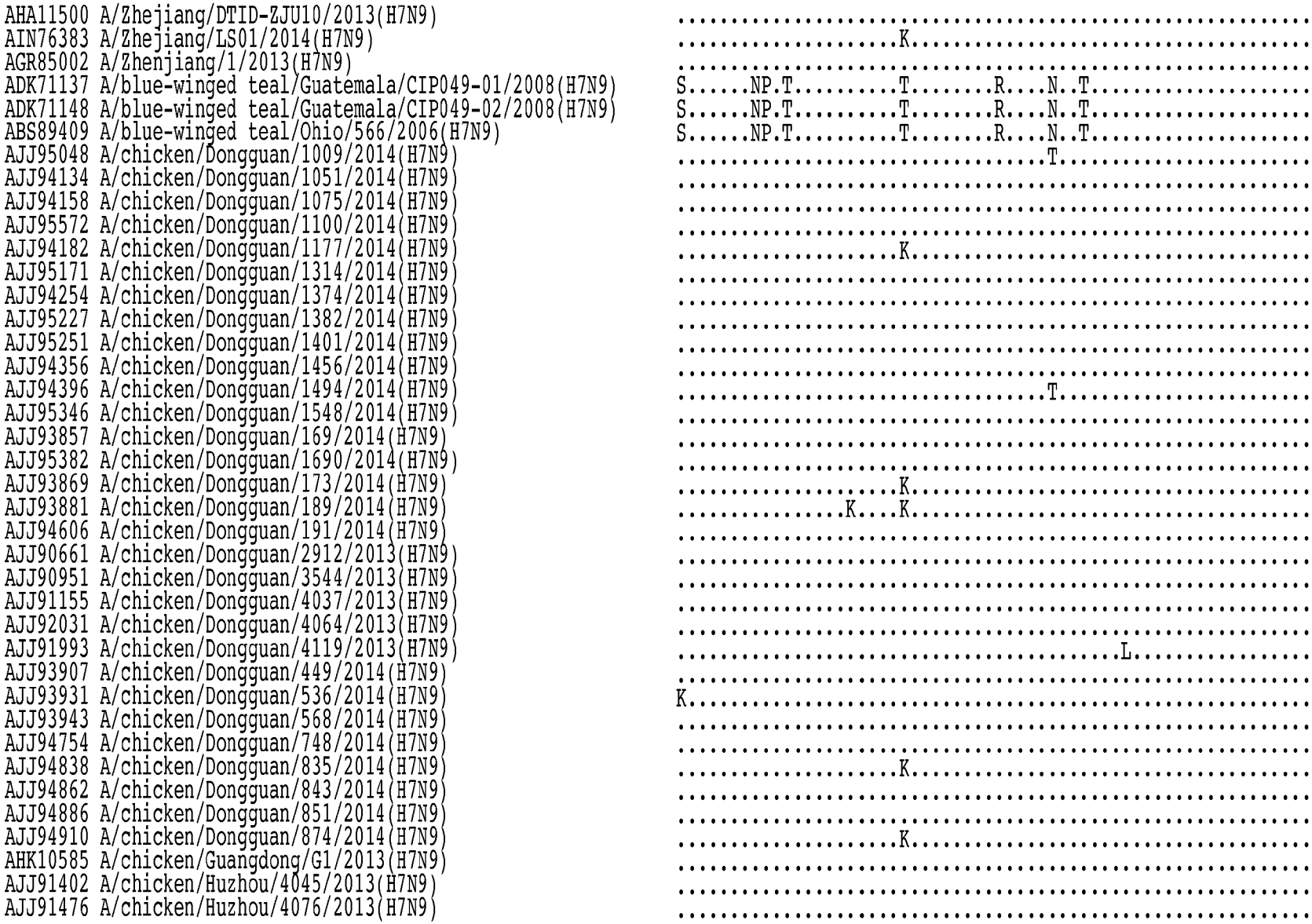

AHA11500 A/Zhejiang/DTID-ZJU10/2013(H7N9)
AIN76383 A/Zhejiang/LS01/2014(H7N9)
AGR85002 A/Zhenjiang/1/2013(H7N9)
ADK71137 A/blue-winged teal/Guatemala/CIP049-01/2008(H7N9)
ADK71148 A/blue-winged teal/Guatemala/CIP049-02/2008(H7N9)
ABS89409 A/blue-winged teal/Ohio/566/2006(H7N9)
AJJ95048 A/chicken/Dongguan/1009/2014(H7N9)
AJJ94134 A/chicken/Dongguan/1051/2014(H7N9)
AJJ94158 A/chicken/Dongguan/1075/2014(H7N9)
AJJ95572 A/chicken/Dongguan/1100/2014(H7N9)
AJJ94182 A/chicken/Dongguan/1177/2014(H7N9)
AJJ95171 A/chicken/Dongguan/1314/2014(H7N9)
AJJ94254 A/chicken/Dongguan/1374/2014(H7N9)
AJJ95227 A/chicken/Dongguan/1382/2014(H7N9)
AJJ95251 A/chicken/Dongguan/1401/2014(H7N9)
AJJ94356 A/chicken/Dongguan/1456/2014(H7N9)
AJJ94396 A/chicken/Dongguan/1494/2014(H7N9)
AJJ95346 A/chicken/Dongguan/1548/2014(H7N9)
AJJ93857 A/chicken/Dongguan/169/2014(H7N9)
AJJ95382 A/chicken/Dongguan/1690/2014(H7N9)
AJJ93869 A/chicken/Dongguan/173/2014(H7N9)
AJJ93881 A/chicken/Dongguan/189/2014(H7N9)
AJJ94606 A/chicken/Dongguan/191/2014(H7N9)
AJJ90661 A/chicken/Dongguan/2912/2013(H7N9)
AJJ90951 A/chicken/Dongguan/3544/2013(H7N9)
AJJ91155 A/chicken/Dongguan/4037/2013(H7N9)
AJJ92031 A/chicken/Dongguan/4064/2013(H7N9)
AJJ91993 A/chicken/Dongguan/4119/2013(H7N9)
AJJ93907 A/chicken/Dongguan/449/2014(H7N9)
AJJ93931 A/chicken/Dongguan/536/2014(H7N9)
AJJ93943 A/chicken/Dongguan/568/2014(H7N9)
AJJ94754 A/chicken/Dongguan/748/2014(H7N9)
AJJ94838 A/chicken/Dongguan/835/2014(H7N9)
AJJ94862 A/chicken/Dongguan/843/2014(H7N9)
AJJ94886 A/chicken/Dongguan/851/2014(H7N9)
AJJ94910 A/chicken/Dongguan/874/2014(H7N9)
AHK10585 A/chicken/Guangdong/G1/2013(H7N9)
AJJ91402 A/chicken/Huzhou/4045/2013(H7N9)
AJJ91476 A/chicken/Huzhou/4076/2013(H7N9)

FIGURE 22-28

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AJJ93051 A/chicken/Jiangxi/10573/2014(H7N9)                                                
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AGJ72861 A/chicken/Zhejiang/DTID-ZJU01/2013(H7N9)
AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013(H7N9)
AGR49554 A/chicken/Zhejiang/SD033/2013(H7N9)
AGR49566 A/duck/Anhui/SC702/2013(H7N9)
BAN16711 A/duck/Gunma/466/2011(H7N9)
AJJ97841 A/duck/Jiangxi/15044/2014(H7N9)
AGQ80952 A/duck/Jiangxi/3096/2009(H7N9)
AGQ80989 A/duck/Jiangxi/3257/2009(H7N9)
BAH22785 A/duck/Mongolia/119/2008(H7N9)
BAN59727 A/duck/Mongolia/129/2010(H7N9)
BAN59726 A/duck/Mongolia/147/2008(H7N9)
AJK00354 A/duck/Zhejiang/LS02/2014(H7N9)
AFM09439 A/emperor goose/Alaska/44063-061/2006(H7N9)
AFV33947 A/goose/Nebraska/17097-4/2011(H7N9)
AFV33945 A/guinea fowl/Nebraska/17096-1/2011(H7N9)
AHA11441 A/guinea fowl/Nebraska/17096/2011(H7N9)
AGR49722 A/homing pigeon/Jiangsu/SD184/2013(H7N9)
AGL97639 A/mallard/Minnesota/AI09-3770/2009(H7N9)
AHZ42537 A/mallard/Minnesota/AI09-3770/2009(H7N9)
ADC29485 A/mallard/Spain/08.00991.3/2005(H7N9)
AHZ41929 A/mallard/Sweden/1621/2002(H7N9)
AAY46211 A/mallard/Sweden/91/2002(H7N9)
AGY42258 A/mallard/Sweden/91/2002(H7N9)
AGE08098 A/northern shoverl/Mississippi/11OS145/2011(H7N9)
AGR49734 A/pigeon/Shanghai/S1069/2013(H7N9)
ACD03594 A/ruddy turnstone/DE/1538/2000(H7N9)
AFX85260 A/ruddy turnstone/Delaware Bay/220/1995(H7N9)
AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000(H7N9)
AJJ94194 A/silkie chicken/Dongguan/1264/2014(H7N9)
AJJ94206 A/silkie chicken/Dongguan/1268/2014(H7N9)
AJJ94344 A/silkie chicken/Dongguan/1451/2014(H7N9)
AJJ95584 A/silkie chicken/Dongguan/1519/2014(H7N9)
AJJ93845 A/silkie chicken/Dongguan/157/2014(H7N9)
AJJ90673 A/silkie chicken/Dongguan/3049/2013(H7N9)
AJJ90795 A/silkie chicken/Dongguan/3281/2013(H7N9)
AJJ90891 A/silkie chicken/Dongguan/3520/2013(H7N9)
AJJ91264 A/silkie chicken/Dongguan/4129/2013(H7N9)
AJJ93979 A/silkie chicken/Dongguan/656/2014(H7N9)
AJJ94959 A/silkie chicken/Dongguan/967/2014(H7N9)
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........N............................................
.......N.............E..M.........L..................
.....................................................
.......N.............E................................
.......N.............E................................
.......N..T..........E................................
.......N.............E..M.........L..................
.......N..T..........E................................
.......N.............E.............V..................
S......NP.T..........T........R....N..T...............
S......NP.T..........T........R....N..T...............
S......NP.T..........T........R....N..T...............
S......NP.T..........T........R....N..T...............
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S......NP.T..........T........R....N..T...............
S......NP.T..........T........R....N..T...............
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.......N.............E................................
.......N.............E................................
.......N.............E................................
S......NP.T..........T........R....N..T...............
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S......NP.T..........T........R....N..T...............
S......NP.T..R.......T.............N..T...............
S......NP.T..........T........R....N..T...............
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D....................................................
..........T..........................................
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.....................K...............................
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.....................K...............................
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FIGURE 22-30

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AJJ92967 A/silkie chicken/Jiangxi/9469/2014(H7N9)                  .....................K.....................................
AJJ97925 A/silkie chicken/Shantou/2050/2014(H7N9)                  .....................................V.....................
AJJ90538 A/silkie chicken/Shenzhen/918/2013(H7N9)                  ..........................................................
AEK84765 A/spot-billed duck/Korea/447/2011(H7N9)                   .......N............E..M.........P.......................
AGW82588 A/tree sparrow/Shanghai/01/2013(H7N9)                     ..........................................................
ABI84694 A/turkey/Minnesota/1/1988(H7N9)                           S......NP.T.........P........R....N..T.....................
AHA11483 A/turkey/Minnesota/14135-2/2009(H7N9)                     S......NP.T.........T........R....K..T.....................
AHA57050 A/turkey/Minnesota/14659/2009(H7N9)                       S......NP.T.........T........R....K..T.....................
AHA57072 A/turkey/Minnesota/18421/2009(H7N9)                       S......NP.T.........T........R....K..T.....................
AHA11472 A/turkey/Minnesota/31676/2009(H7N9)                       S......NP.T.........T........R....K..T.....................
AHA11461 A/turkey/Minnesota/31900/2011(H7N9)                       S......NP.T.........T........R....N..T.....................
AHA11452 A/turkey/Minnesota/32710/2011(H7N9)                       S......NP.T.........T........R....N..T.....................
AAM19228 A/turkey/Minnesota/38429/1988(H7N9)                       S......NP.T.........P........R....N..T.....------------
ACZ48625 A/turkey/Minnesota/38429/1988(H7N9)                       S......NP.T.........P........R....N..T.....................
AEK84760 A/wild bird/Korea/A14/2011(H7N9)                          .......N............E..M.........L.......................
AEK84761 A/wild bird/Korea/A3/2011(H7N9)                           .......N............E..M.........L...K...................
AEK84763 A/wild bird/Korea/A9/2011(H7N9)                           .......N............E..M.........L.......................
AGG53377 A/wild duck/Korea/CSM42-1/2011(H7N9)                      .......N............E..M.........L.......................
AGG53366 A/wild duck/Korea/CSM42-34/2011(H7N9)                     .......N............E..M.........L.......................
AGG53432 A/wild duck/Korea/MHC35-41/2011(H7N9)                     .......N............E..M.........P.......................
AGG53399 A/wild duck/Korea/MHC39-26/2011(H7N9)                     .......N............E..M.........P.......................
AGG53476 A/wild duck/Korea/SH19-27/2010(H7N9)                      .......N............E..M.........L.......................
AGG53487 A/wild duck/Korea/SH19-50/2010(H7N9)                      .......N............E..M.........L.......................
AGG53520 A/wild duck/Korea/SH20-27/2008(H7N9)                      .......N............E..M.........L.......................
AEM98291 A/wild duck/Mongolia/1-241/2008(H7N9)                     .......N............E..M.........L.......................
AGR49770 A/wild pigeon/Jiangsu/SD001/2013(H7N9)                    ..........................................................

Positions from 361 till 420
Consensus sequence                                                 WYGFRHQNAQGEGTAADYKSTQSAIDQITGKLNRLIEKTNQQFELIDNEFNEVEKQIGNV
ACX53683 A/goose/Czech Republic/1848-K9/2009(H7N9)                 ..........................................................T.....
ADN34727 A/goose/Czech Republic/1848-T14/2009(H7N9)                ..........................................................T.....
CAY39406 A/Anas crecca/Spain/1460/2008(H7N9)                       ..........................................................T.....
AHZ39686 A/Anhui/DEWH72-01/2013(H7N9)                              ..........................................................
AHZ39710 A/Anhui/DEWH72-03/2013(H7N9)                              ..........................................................
AHZ39746 A/Anhui/DEWH72-06/2013(H7N9)                              ..........................................................
AHM24224 A/Beijing/3/2013(H7N9)                                    ..........................................................
AHD25003 A/Guangdong/02/2013(H7N9)                                 ..........................................................
AHH25185 A/Guangdong/04/2013(H7N9)                                 ..........................................................
AGI60301 A/Hangzhou/1/2013(H7N9)                                   ..........................................................
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FIGURE 22-31

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AGK84857 A/Hangzhou/2/2013(H7N9)
AHF20528 A/Hong Kong/47029/2013(H7N9)
AGY41893 A/Huizhou/01/2013(H7N9)
AGO51387 A/Jiangsu/2/2013(H7N9)
AGJ73503 A/Nanjing/1/2013(H7N9)
AGR85026 A/Nanjing/2/2013(H7N9)
AGR84954 A/Nanjing/6/2013(H7N9)
AHK10800 A/Shanghai/01/2014(H7N9)
AGL44438 A/Shanghai/02/2013(H7N9)
AGL33692 A/Shanghai/4655T/2013(H7N9)
AGL33693 A/Shanghai/4659T/2013(H7N9)
AGI60292 A/Shanghai/4664T/2013(H7N9)
AGM53887 A/Shanghai/4701T/2013(H7N9)
AGM53886 A/Shanghai/4842T/2013(H7N9)
AGM53883 A/Shanghai/5083T/2013(H7N9)
AGM53884 A/Shanghai/5180T/2013(H7N9)
AGM53885 A/Shanghai/5240T/2013(H7N9)
AGW82600 A/Shanghai/CN01/2013(H7N9)
AHF20568 A/Shanghai/CN02/2013(H7N9)
AGW82612 A/Shanghai/JS01/2013(H7N9)
AID70634 A/Shanghai/Mix1/2014(H7N9)
AHJ57411 A/Shanghai/PD-01/2014(H7N9)
AHJ57418 A/Shanghai/PD-02/2014(H7N9)
AJJ96841 A/Shenzhen/SP139/2014(H7N9)
AJJ91909 A/Shenzhen/SP26/2014(H7N9)
AJJ91945 A/Shenzhen/SP38/2014(H7N9)
AJJ91885 A/Shenzhen/SP4/2014(H7N9)
AJJ91957 A/Shenzhen/SP44/2014(H7N9)
AJJ91969 A/Shenzhen/SP48/2014(H7N9)
AJJ95596 A/Shenzhen/SP58/2014(H7N9)
AJJ95632 A/Shenzhen/SP62/2014(H7N9)
AJJ95620 A/Shenzhen/SP75/2014(H7N9)
AGR84942 A/Suzhou/5/2013(H7N9)
AGL43637 A/Taiwan/1/2013(H7N9)
AGL95088 A/Taiwan/S02076/2013(H7N9)
AGL95098 A/Taiwan/T02081/2013(H7N9)
AGN69474 A/Wuxi/1/2013(H7N9)
AGN69462 A/Wuxi/2/2013(H7N9)
AGR84990 A/Wuxi/3/2013(H7N9)
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AGR84978 A/Wuxi/4/2013(H7N9)                                    ..................................................
AGO02477 A/Xuzhou/1/2013(H7N9)                                  ..................................................
AGU02230 A/Zhejiang/DTID-ZJU05/2013(H7N9)                       ..................................................
AGU02233 A/Zhejiang/DTID-ZJU08/2013(H7N9)                       ..................................................
AHA11500 A/Zhejiang/DTID-ZJU10/2013(H7N9)                       ..................................................
AIN76383 A/Zhejiang/LS01/2014(H7N9)                             ..................................................
AGR85002 A/Zhenjiang/1/2013(H7N9)                               ..................................................
ADK71137 A/blue-winged teal/Guatemala/CIP049-01/2008(H7N9)      ..................................D....H........S.I.Q.....
ADK71148 A/blue-winged teal/Guatemala/CIP049-02/2008(H7N9)      ..................................D.............S.I.Q.....
ABS89409 A/blue-winged teal/Ohio/566/2006(H7N9)                 ..................................D.............S.I.Q.....
AJJ95048 A/chicken/Dongguan/1009/2014(H7N9)                     ..................................................
AJJ94134 A/chicken/Dongguan/1051/2014(H7N9)                     ..................................................
AJJ94158 A/chicken/Dongguan/1075/2014(H7N9)                     ....K.............................................T.....
AJJ95572 A/chicken/Dongguan/1100/2014(H7N9)                     ..................................................
AJJ94182 A/chicken/Dongguan/1177/2014(H7N9)                     ..................................................
AJJ95171 A/chicken/Dongguan/1314/2014(H7N9)                     ....K.............................................T.....
AJJ94254 A/chicken/Dongguan/1374/2014(H7N9)                     ....K.............................................T.....
AJJ95227 A/chicken/Dongguan/1382/2014(H7N9)                     ..................................................
AJJ95251 A/chicken/Dongguan/1401/2014(H7N9)                     ..................................................T.....
AJJ94356 A/chicken/Dongguan/1456/2014(H7N9)                     ..................................................T.....
AJJ94396 A/chicken/Dongguan/1494/2014(H7N9)                     ..................................................
AJJ95346 A/chicken/Dongguan/1548/2014(H7N9)                     ..................................................T.....
AJJ93857 A/chicken/Dongguan/169/2014(H7N9)                      ..................................................
AJJ95382 A/chicken/Dongguan/1690/2014(H7N9)                     ..................................................
AJJ93869 A/chicken/Dongguan/173/2014(H7N9)                      ..................................................
AJJ93881 A/chicken/Dongguan/189/2014(H7N9)                      ..................................................
AJJ94606 A/chicken/Dongguan/191/2014(H7N9)                      ..................................................
AJJ90661 A/chicken/Dongguan/2912/2013(H7N9)                     ..................................................
AJJ90951 A/chicken/Dongguan/3544/2013(H7N9)                     ..................................................
AJJ91155 A/chicken/Dongguan/4037/2013(H7N9)                     ..................................................
AJJ92031 A/chicken/Dongguan/4064/2013(H7N9)                     ..................................................
AJJ91993 A/chicken/Dongguan/4119/2013(H7N9)                     ..................................................
AJJ93907 A/chicken/Dongguan/449/2014(H7N9)                      ..................................................
AJJ93931 A/chicken/Dongguan/536/2014(H7N9)                      ..................................................
AJJ93943 A/chicken/Dongguan/568/2014(H7N9)                      ..................................................
AJJ94754 A/chicken/Dongguan/748/2014(H7N9)                      ..................................................
AJJ94838 A/chicken/Dongguan/835/2014(H7N9)                      ..................................................
AJJ94862 A/chicken/Dongguan/843/2014(H7N9)                      ..................................................
AJJ94886 A/chicken/Dongguan/851/2014(H7N9)                      ..................................................
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FIGURE 22-33

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AJJ94910 A/chicken/Dongguan/874/2014(H7N9)
AHK10585 A/chicken/Guangdong/G1/2013(H7N9)
AJJ91402 A/chicken/Huzhou/4045/2013(H7N9)
AJJ91476 A/chicken/Huzhou/4076/2013(H7N9)
AJJ93051 A/chicken/Jiangxi/10573/2014(H7N9)
AJJ96552 A/chicken/Jiangxi/12206/2014(H7N9)
AJJ96684 A/chicken/Jiangxi/13207/2014(H7N9)
AJJ96720 A/chicken/Jiangxi/13220/2014(H7N9)
AJJ96732 A/chicken/Jiangxi/13223/2014(H7N9)
AJJ97267 A/chicken/Jiangxi/13252/2014(H7N9)
AJJ97291 A/chicken/Jiangxi/13493/2014(H7N9)
AJJ96889 A/chicken/Jiangxi/13496/2014(H7N9)
AJJ96901 A/chicken/Jiangxi/13502/2014(H7N9)
AJJ97331 A/chicken/Jiangxi/13512/2014(H7N9)
AJJ96925 A/chicken/Jiangxi/13513/2014(H7N9)
AJJ97373 A/chicken/Jiangxi/13521/2014(H7N9)
AJJ97443 A/chicken/Jiangxi/13530/2014(H7N9)
AJJ97582 A/chicken/Jiangxi/14023/2014(H7N9)
AJJ97697 A/chicken/Jiangxi/14517/2014(H7N9)
AJJ97709 A/chicken/Jiangxi/14518/2014(H7N9)
AJJ97745 A/chicken/Jiangxi/14554/2014(H7N9)
AJJ97899 A/chicken/Jiangxi/15524/2014(H7N9)
AJJ96817 A/chicken/Jiangxi/9513/2014(H7N9)
AJJ93027 A/chicken/Jiangxi/9558/2014(H7N9)
AGR49399 A/chicken/Jiangxi/SD001/2013(H7N9)
AGQ81043 A/chicken/Rizhao/515/2013(H7N9)
AGR33894 A/chicken/Rizhao/719b/2013(H7N9)
AHN96472 A/chicken/Shanghai/PD-CN-02/2014(H7N9)
AGR49495 A/chicken/Shanghai/S1358/2013(H7N9)
AGR49506 A/chicken/Shanghai/S1410/2013(H7N9)
AJJ97757 A/chicken/Shantou/2537/2014(H7N9)
AJJ97973 A/chicken/Shantou/4325/2014(H7N9)
AJJ97998 A/chicken/Shantou/4816/2014(H7N9)
AJJ91314 A/chicken/Shaoxing/2417/2013(H7N9)
AJJ91725 A/chicken/Shaoxing/5201/2013(H7N9)
AJJ95464 A/chicken/Shenzhen/138/2014(H7N9)
AJJ90576 A/chicken/Shenzhen/1665/2013(H7N9)
AJJ90588 A/chicken/Shenzhen/2110/2013(H7N9)
AJJ91035 A/chicken/Shenzhen/3780/2013(H7N9)

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AJJ90490 A/chicken/Shenzhen/742/2013(H7N9)
AJJ92005 A/chicken/Shenzhen/801/2013(H7N9)
AJJ90526 A/chicken/Shenzhen/898/2013(H7N9)
AIU47013 A/chicken/Suzhou/0402O1H/2013(H7N9)
AGJ72861 A/chicken/Zhejiang/DTID-ZJU01/2013(H7N9)
AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013(H7N9)
AGR49554 A/chicken/Zhejiang/SD033/2013(H7N9)
AGR49566 A/duck/Anhui/SC702/2013(H7N9)
BAN16711 A/duck/Gunma/466/2011(H7N9)
AJJ97841 A/duck/Jiangxi/15044/2014(H7N9)
AGQ80952 A/duck/Jiangxi/3096/2009(H7N9)
AGQ80989 A/duck/Jiangxi/3257/2009(H7N9)
BAH22785 A/duck/Mongolia/119/2008(H7N9)
BAN59727 A/duck/Mongolia/129/2010(H7N9)
BAN59726 A/duck/Mongolia/147/2008(H7N9)
AJK00354 A/duck/Zhejiang/LS02/2014(H7N9)
AFM09439 A/emperor goose/Alaska/44063-061/2006(H7N9)
AFV33947 A/goose/Nebraska/17097-4/2011(H7N9)
AFV33945 A/guinea fowl/Nebraska/17096-1/2011(H7N9)
AHA11441 A/guinea fowl/Nebraska/17096/2011(H7N9)
AGR49722 A/homing pigeon/Jiangsu/SD184/2013(H7N9)
AGL97639 A/mallard/Minnesota/AI09-3770/2009(H7N9)
AHZ42537 A/mallard/Minnesota/AI09-3770/2009(H7N9)
ADC29485 A/mallard/Spain/08.00991.3/2005(H7N9)
AHZ41929 A/mallard/Sweden/1621/2002(H7N9)
AAY46211 A/mallard/Sweden/91/2002(H7N9)
AGY42258 A/mallard/Sweden/91/2002(H7N9)
AGE08098 A/northern shoveler/Mississippi/11OS145/2011(H7N9)
AGR49734 A/pigeon/Shanghai/S1069/2013(H7N9)
ACD03594 A/ruddy turnstone/DE/1538/2000(H7N9)
AFX85260 A/ruddy turnstone/Delaware Bay/220/1995(H7N9)
AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000(H7N9)
AJJ94194 A/silkie chicken/Dongguan/1264/2014(H7N9)
AJJ94206 A/silkie chicken/Dongguan/1268/2014(H7N9)
AJJ94344 A/silkie chicken/Dongguan/1451/2014(H7N9)
AJJ95584 A/silkie chicken/Dongguan/1519/2014(H7N9)
AJJ93845 A/silkie chicken/Dongguan/157/2014(H7N9)
AJJ90673 A/silkie chicken/Dongguan/3049/2013(H7N9)
AJJ90795 A/silkie chicken/Dongguan/3281/2013(H7N9)
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FIGURE 22-35

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AJJ90891 A/silkie chicken/Dongguan/3520/2013(H7N9)  ..................................................
AJJ91264 A/silkie chicken/Dongguan/4129/2013(H7N9)  ..................................M...............
AJJ93979 A/silkie chicken/Dongguan/656/2014(H7N9)   ..........................................G.......
AJJ94959 A/silkie chicken/Dongguan/967/2014(H7N9)   ..................................................
AJJ92967 A/silkie chicken/Jiangxi/9469/2014(H7N9)   ..................................................
AJJ97925 A/silkie chicken/Shantou/2050/2014(H7N9)   ..................................................
AJJ90538 A/silkie chicken/Shenzhen/918/2013(H7N9)   ..................................................
AEK84765 A/spot-billed duck/Korea/447/2011(H7N9)    ..............................................T.....
AGW82588 A/tree sparrow/Shanghai/01/2013(H7N9)     ..................................................
ABI84694 A/turkey/Minnesota/1/1988(H7N9)           ....K.............................D.....S.I.Q.....
AHA11483 A/turkey/Minnesota/14135-2/2009(H7N9)     ........................S.....D.....S.I.Q.....
AHA57050 A/turkey/Minnesota/14659/2009(H7N9)       ........................S.....D.....S.I.Q.....
AHA57072 A/turkey/Minnesota/18421/2009(H7N9)       ..............................D.....S.I.Q.....
AHA11472 A/turkey/Minnesota/31676/2009(H7N9)       ........................N.....D.....S.I.Q.....
AHA11461 A/turkey/Minnesota/31900/2011(H7N9)       ..............................D.....S.I.Q.....
AHA11452 A/turkey/Minnesota/32710/2011(H7N9)       ..............................D.....M.....S.I.Q.....
AAM19228 A/turkey/Minnesota/38429/1988(H7N9)       --------------------------------------------------
ACZ48625 A/turkey/Minnesota/38429/1988(H7N9)       ....K.............................D.....----------
AEK84760 A/wild bird/Korea/A14/2011(H7N9)          ..............................................T.....
AEK84761 A/wild bird/Korea/A3/2011(H7N9)           ..............................................T.....
AEK84763 A/wild bird/Korea/A9/2011(H7N9)           ..............................................T.....
AGG53377 A/wild duck/Korea/CSM42-1/2011(H7N9)      ..............................................T.....
AGG53366 A/wild duck/Korea/CSM42-34/2011(H7N9)     ..............................................T.....
AGG53432 A/wild duck/Korea/MHC35-41/2011(H7N9)     ..............................................T.....
AGG53399 A/wild duck/Korea/MHC39-26/2011(H7N9)     ..............................................T.....
AGG53476 A/wild duck/Korea/SH19-27/2010(H7N9)      ..............................................T.....
AGG53487 A/wild duck/Korea/SH19-50/2010(H7N9)      ..............................................T.....
AGG53520 A/wild duck/Korea/SH20-27/2008(H7N9)      ..............................................T.....
AEM98291 A/wild duck/Mongolia/1-241/2008(H7N9)     ..............................................T.....
AGR49770 A/wild pigeon/Jiangsu/SD001/2013(H7N9)    ..................................................

Positions from 421 till 480
        Consensus sequence                         INWTRDSITEVWSYNAELLVAMENQHTIDLADSEMDKLYERVKRQLRENAEEDGTGCFEI
ACX53683  A/goose/Czech Republic/1848-K9/2009(H7N9)  .......M.............................N.....................
ADN34727  A/goose/Czech Republic/1848-T14/2009(H7N9) .......M.............................N.....................
CAY39406  A/Anas crecca/Spain/1460/2008(H7N9)        ..................................N.....................
AHZ39686  A/Anhui/DEWH72-01/2013(H7N9)              ..................................................
AHZ39710  A/Anhui/DEWH72-03/2013(H7N9)              ..................................................
AHZ39746  A/Anhui/DEWH72-06/2013(H7N9)              ..................................................
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FIGURE 22-36

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AHM24224 A/Beijing/3/2013(H7N9)
AHD25003 A/Guangdong/02/2013(H7N9)
AHH25185 A/Guangdong/04/2013(H7N9)
AGI60301 A/Hangzhou/1/2013(H7N9)
AGK84857 A/Hangzhou/2/2013(H7N9)
AHF20528 A/Hong Kong/470129/2013(H7N9)
AGY41893 A/Huizhou/01/2013(H7N9)
AGO51387 A/Jiangsu/2/2013(H7N9)
AGJ73503 A/Nanjing/1/2013(H7N9)
AGR85026 A/Nanjing/2/2013(H7N9)
AGR84954 A/Nanjing/6/2013(H7N9)
AHK10800 A/Shanghai/01/2014(H7N9)
AGL44438 A/Shanghai/02/2013(H7N9)
AGL33692 A/Shanghai/4655T/2013(H7N9)
AGL33693 A/Shanghai/4659T/2013(H7N9)
AGI60292 A/Shanghai/4664T/2013(H7N9)
AGM53887 A/Shanghai/4701T/2013(H7N9)
AGM53886 A/Shanghai/4842T/2013(H7N9)
AGM53883 A/Shanghai/5083T/2013(H7N9)
AGM53884 A/Shanghai/5180T/2013(H7N9)
AGM53885 A/Shanghai/5240T/2013(H7N9)
AGW82600 A/Shanghai/CN01/2013(H7N9)
AHF20568 A/Shanghai/CN02/2013(H7N9)
AGW82612 A/Shanghai/JS01/2013(H7N9)
AID70634 A/Shanghai/Mix1/2014(H7N9)
AHJ57411 A/Shanghai/PD-01/2014(H7N9)
AHJ57418 A/Shanghai/PD-02/2014(H7N9)
AJJ96841 A/Shenzhen/SP139/2014(H7N9)
AJJ91909 A/Shenzhen/SP26/2014(H7N9)
AJJ91945 A/Shenzhen/SP38/2014(H7N9)
AJJ91885 A/Shenzhen/SP4/2014(H7N9)
AJJ91957 A/Shenzhen/SP44/2014(H7N9)
AJJ91969 A/Shenzhen/SP48/2014(H7N9)
AJJ95596 A/Shenzhen/SP58/2014(H7N9)
AJJ95632 A/Shenzhen/SP62/2014(H7N9)
AJJ95620 A/Shenzhen/SP75/2014(H7N9)
AGR84942 A/Suzhou/5/2013(H7N9)
AGL43637 A/Taiwan/1/2013(H7N9)
AGL95088 A/Taiwan/S02076/2013(H7N9)
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FIGURE 22-37

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AGL95098 A/Taiwan/T02081/2013(H7N9)                                                
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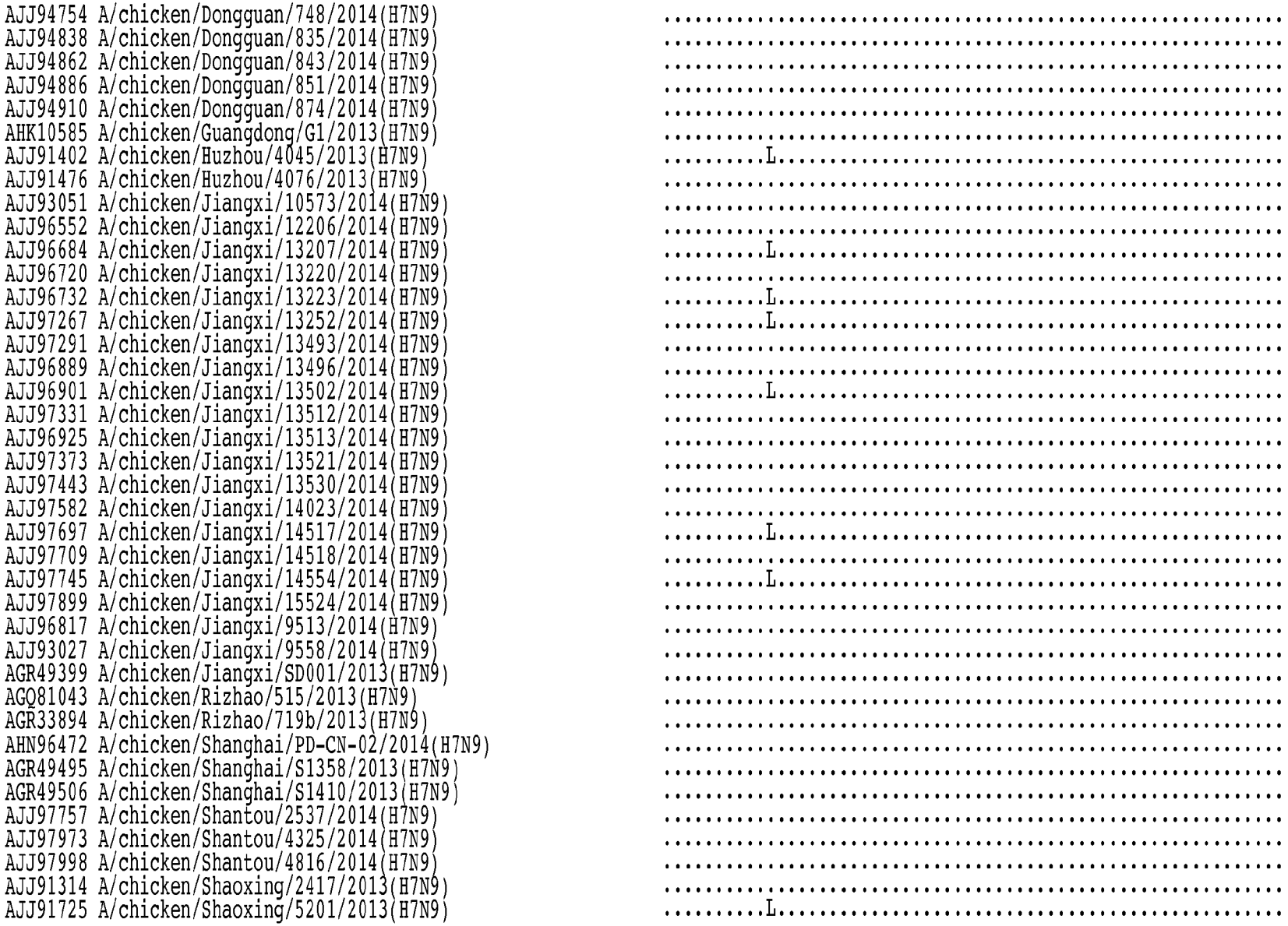

AJJ94754 A/chicken/Dongguan/748/2014(H7N9)
AJJ94838 A/chicken/Dongguan/835/2014(H7N9)
AJJ94862 A/chicken/Dongguan/843/2014(H7N9)
AJJ94886 A/chicken/Dongguan/851/2014(H7N9)
AJJ94910 A/chicken/Dongguan/874/2014(H7N9)
AHK10585 A/chicken/Guangdong/G1/2013(H7N9)
AJJ91402 A/chicken/Huzhou/4045/2013(H7N9)
AJJ91476 A/chicken/Huzhou/4076/2013(H7N9)
AJJ93051 A/chicken/Jiangxi/10573/2014(H7N9)
AJJ96552 A/chicken/Jiangxi/12206/2014(H7N9)
AJJ96684 A/chicken/Jiangxi/13207/2014(H7N9)
AJJ96720 A/chicken/Jiangxi/13220/2014(H7N9)
AJJ96732 A/chicken/Jiangxi/13223/2014(H7N9)
AJJ97267 A/chicken/Jiangxi/13252/2014(H7N9)
AJJ97291 A/chicken/Jiangxi/13493/2014(H7N9)
AJJ96889 A/chicken/Jiangxi/13496/2014(H7N9)
AJJ96901 A/chicken/Jiangxi/13502/2014(H7N9)
AJJ97331 A/chicken/Jiangxi/13512/2014(H7N9)
AJJ96925 A/chicken/Jiangxi/13513/2014(H7N9)
AJJ97373 A/chicken/Jiangxi/13521/2014(H7N9)
AJJ97443 A/chicken/Jiangxi/13530/2014(H7N9)
AJJ97582 A/chicken/Jiangxi/14023/2014(H7N9)
AJJ97697 A/chicken/Jiangxi/14517/2014(H7N9)
AJJ97709 A/chicken/Jiangxi/14518/2014(H7N9)
AJJ97745 A/chicken/Jiangxi/14554/2014(H7N9)
AJJ97899 A/chicken/Jiangxi/15524/2014(H7N9)
AJJ96817 A/chicken/Jiangxi/9513/2014(H7N9)
AJJ93027 A/chicken/Jiangxi/9558/2014(H7N9)
AGR49399 A/chicken/Jiangxi/SD001/2013(H7N9)
AGQ81043 A/chicken/Rizhao/515/2013(H7N9)
AGR33894 A/chicken/Rizhao/719b/2013(H7N9)
AHN96472 A/chicken/Shanghai/PD-CN-02/2014(H7N9)
AGR49495 A/chicken/Shanghai/S1358/2013(H7N9)
AGR49506 A/chicken/Shanghai/S1410/2013(H7N9)
AJJ97757 A/chicken/Shantou/2537/2014(H7N9)
AJJ97973 A/chicken/Shantou/4325/2014(H7N9)
AJJ97998 A/chicken/Shantou/4816/2014(H7N9)
AJJ91314 A/chicken/Shaoxing/2417/2013(H7N9)
AJJ91725 A/chicken/Shaoxing/5201/2013(H7N9)
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FIGURE 22-39

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AJJ95464 A/chicken/Shenzhen/138/2014(H7N9)                          ..................................................
AJJ90576 A/chicken/Shenzhen/1665/2013(H7N9)                         ..................................................
AJJ90588 A/chicken/Shenzhen/2110/2013(H7N9)                         ..................................................
AJJ91035 A/chicken/Shenzhen/3780/2013(H7N9)                         ..................................................
AJJ90490 A/chicken/Shenzhen/742/2013(H7N9)                          ..................................................
AJJ92005 A/chicken/Shenzhen/801/2013(H7N9)                          ..................................................
AJJ90526 A/chicken/Shenzhen/898/2013(H7N9)                          ..................................................
AIU47013 A/chicken/Suzhou/040201H/2013(H7N9)                        ..................................................
AGJ72861 A/chicken/Zhejiang/DTID-ZJU01/2013(H7N9)                   ..................................................
AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013(H7N9)                   ..................................................
AGR49554 A/chicken/Zhejiang/SD033/2013(H7N9)                        ..........................................R.......
AGR49566 A/duck/Anhui/SC702/2013(H7N9)                              ..................................................
BAN16711 A/duck/Gunma/466/2011(H7N9)                                .......M..........................N......R.........D.....
AJJ97841 A/duck/Jiangxi/15044/2014(H7N9)                            ..................................................
AGQ80952 A/duck/Jiangxi/3096/2009(H7N9)                             .......M..........................N......R.......
AGQ80989 A/duck/Jiangxi/3257/2009(H7N9)                             .......M..........................N......R.......
BAH22785 A/duck/Mongolia/119/2008(H7N9)                             .......M..........................N.......
BAN59727 A/duck/Mongolia/129/2010(H7N9)                             .......M..........................N.......
BAN59726 A/duck/Mongolia/147/2008(H7N9)                             .......M..........................N.......
AJK00354 A/duck/Zhejiang/LS02/2014(H7N9)                            .......M..........................N.......
AFM09439 A/emperor goose/Alaska/44063-061/2006(H7N9)                .......M..........................N......RK.....
AFV33947 A/goose/Nebraska/17097-4/2011(H7N9)                        .......M..I.......................N......RK.....
AFV33945 A/guinea fowl/Nebraska/17096-1/2011(H7N9)                  .......M..I.......................N......RK.....
AHA11441 A/guinea fowl/Nebraska/17096/2011(H7N9)                    .......M..I.......................N......RK.....
AGR49722 A/homing pigeon/Jiangsu/SD184/2013(H7N9)                   ..................................................
AGL97639 A/mallard/Minnesota/AI09-3770/2009(H7N9)                   .......M..L.......................N......RK.....
AHZ42537 A/mallard/Minnesota/AI09-3770/2009(H7N9)                   .......M..L.......................N......RK.....
ADC29485 A/mallard/Spain/08.00991.3/2005(H7N9)                      .......M..........................N.......
AHZ41929 A/mallard/Sweden/1621/2002(H7N9)                           .......M..........................N......R.......
AAY46211 A/mallard/Sweden/91/2002(H7N9)                             .......M..........................N......R.......
AGY42258 A/mallard/Sweden/91/2002(H7N9)                             .......M..........................N......R.......
AGE08098 A/northern shoverl/Mississippi/11OS145/2011(H7N9)          .......M..........................N......RK.....
AGR49734 A/pigeon/Shanghai/S1069/2013(H7N9)                         ..................................................
ACD03594 A/ruddy turnstone/DE/1538/2000(H7N9)                       .......M..........................N......RK.....
AFX85260 A/ruddy turnstone/Delaware Bay/220/1995(H7N9)              .......M..........................N......RK.....
AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000(H7N9)            .......M..........................N......RK.....
AJJ94194 A/silkie chicken/Dongguan/1264/2014(H7N9)                  ..................................................
AJJ94206 A/silkie chicken/Dongguan/1268/2014(H7N9)                  ..................................................
AJJ94344 A/silkie chicken/Dongguan/1451/2014(H7N9)                  ..................................................
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FIGURE 22-40

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AJJ95584 A/silkie chicken/Dongguan/1519/2014(H7N9)  ..................................................
AJJ93845 A/silkie chicken/Dongguan/157/2014(H7N9)   ..................................................
AJJ90673 A/silkie chicken/Dongguan/3049/2013(H7N9)  ..................................................
AJJ90795 A/silkie chicken/Dongguan/3281/2013(H7N9)  ..................................................
AJJ90891 A/silkie chicken/Dongguan/3520/2013(H7N9)  ..................................................
AJJ91264 A/silkie chicken/Dongguan/4129/2013(H7N9)  ..................................................
AJJ93979 A/silkie chicken/Dongguan/656/2014(H7N9)   ..................................................
AJJ94959 A/silkie chicken/Dongguan/967/2014(H7N9)   ..................................................
AJJ92967 A/silkie chicken/Jiangxi/9469/2014(H7N9)   ..................................................
AJJ97925 A/silkie chicken/Shantou/2050/2014(H7N9)   ..................................................
AJJ90538 A/silkie chicken/Shenzhen/918/2013(H7N9)   ..................................................
AEK84765 A/spot-billed duck/Korea/447/2011(H7N9)    .......M..........................N.....R.....
AGW82588 A/tree sparrow/Shanghai/01/2013(H7N9)     ..................................................
ABI84694 A/turkey/Minnesota/1/1988(H7N9)           .......M..........................N.....R.....
AHA11483 A/turkey/Minnesota/14135-2/2009(H7N9)     .......M..........................N.....RK....
AHA57050 A/turkey/Minnesota/14659/2009(H7N9)       .......M..........................N.....RK....
AHA57072 A/turkey/Minnesota/18421/2009(H7N9)       .......M..........................N.....RK....
AHA11472 A/turkey/Minnesota/31676/2009(H7N9)       .......M..........................N.....RK....
AHA11461 A/turkey/Minnesota/31900/2011(H7N9)       .......M..I.......................N.....RK....
AHA11452 A/turkey/Minnesota/32710/2011(H7N9)       .......M..I.......................N.....RK....
AAM19228 A/turkey/Minnesota/38429/1988(H7N9)       ----------------------------------------------.K..GSHYG-
ACZ48625 A/turkey/Minnesota/38429/1988(H7N9)       --------------------------------------------------------
AEK84760 A/wild bird/Korea/A14/2011(H7N9)          .......M..........................N.....R.....
AEK84761 A/wild bird/Korea/A3/2011(H7N9)           .......M..........................N.....R.....
AEK84763 A/wild bird/Korea/A9/2011(H7N9)           .......M..........................N.....R.....
AGG53377 A/wild duck/Korea/CSM42-1/2011(H7N9)      .......M..........................N.....R.....
AGG53366 A/wild duck/Korea/CSM42-34/2011(H7N9)     .......M..........................N.....R.....
AGG53432 A/wild duck/Korea/MHC35-41/2011(H7N9)     .......M..........................N.....R.....
AGG53399 A/wild duck/Korea/MHC39-26/2011(H7N9)     .......M..........................N.....R.....
AGG53476 A/wild duck/Korea/SH19-27/2010(H7N9)      .......M..........................N.....R.....
AGG53487 A/wild duck/Korea/SH19-50/2010(H7N9)      .......M..........................N.....R.....
AGG53520 A/wild duck/Korea/SH20-27/2008(H7N9)      .......M..........................N..........
AEM98291 A/wild duck/Mongolia/1-241/2008(H7N9)     .......M..........................N..........
AGR49770 A/wild pigeon/Jiangsu/SD001/2013(H7N9)    ..................................................

Positions from 481 till 540
        Consensus sequence                         FHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVKLSSGYKDVILWFSFGASCFILLA
ACX53683 A/goose/Czech Republic/1848-K9/2009(H7N9) ..............................N...................
ADN34727 A/goose/Czech Republic/1848-T14/2009(H7N9) ..............................N...................
```

FIGURE 22-41

```
CAY39406  A/Anas crecca/Spain/1460/2008(H7N9)  ..................................................
AHZ39686  A/Anhui/DEWH72-01/2013(H7N9)         ..................................................
AHZ39710  A/Anhui/DEWH72-03/2013(H7N9)         ..................................................
AHZ39746  A/Anhui/DEWH72-06/2013(H7N9)         ..................................................
AHM24224  A/Beijing/3/2013(H7N9)               ..................................................
AHD25003  A/Guangdong/02/2013(H7N9)            ..................................................
AHH25185  A/Guangdong/04/2013(H7N9)            ..................................................
AGI60301  A/Hangzhou/1/2013(H7N9)              ..................................................
AGK84857  A/Hangzhou/2/2013(H7N9)              ..................................................
AHF20528  A/Hong Kong/470129/2013(H7N9)        ..................................................
AGY41893  A/Huizhou/01/2013(H7N9)              ..................................................
AGO51387  A/Jiangsu/2/2013(H7N9)               ...............X..XBX.....................XX.
AGJ73503  A/Nanjing/1/2013(H7N9)               ..................................................
AGR85026  A/Nanjing/2/2013(H7N9)               ..................................................
AGR84954  A/Nanjing/6/2013(H7N9)               ..................................................
AHK10800  A/Shanghai/01/2014(H7N9)             ..................................................
AGL44438  A/Shanghai/02/2013(H7N9)             ..................................................
AGL33692  A/Shanghai/4655T/2013(H7N9)          ..................................................
AGL33693  A/Shanghai/4659T/2013(H7N9)          ..................................................
AGI60292  A/Shanghai/4664T/2013(H7N9)          ..................................................
AGM53887  A/Shanghai/4701T/2013(H7N9)          ..................................................
AGM53886  A/Shanghai/4842T/2013(H7N9)          ..................................................
AGM53883  A/Shanghai/5083T/2013(H7N9)          ..................................................
AGM53884  A/Shanghai/5180T/2013(H7N9)          ..................................................
AGM53885  A/Shanghai/5240T/2013(H7N9)          ..................................................
AGW82600  A/Shanghai/CN01/2013(H7N9)           ..................................................
AHF20568  A/Shanghai/CN02/2013(H7N9)           ..................................................
AGW82612  A/Shanghai/JS01/2013(H7N9)           ..................................................
AID70634  A/Shanghai/Mix1/2014(H7N9)           ..................................................
AHJ57411  A/Shanghai/PD-01/2014(H7N9)          ..................................................
AHJ57418  A/Shanghai/PD-02/2014(H7N9)          ..................................................
AJU96841  A/Shenzhen/SP139/2014(H7N9)          ..................................................
AJU91909  A/Shenzhen/SP26/2014(H7N9)           .....G.....................R.................
AJU91945  A/Shenzhen/SP38/2014(H7N9)           ..................................................
AJU91885  A/Shenzhen/SP4/2014(H7N9)            ..................................R...............
AJU91957  A/Shenzhen/SP44/2014(H7N9)           ..................................................
AJU91969  A/Shenzhen/SP48/2014(H7N9)           ..................................................
AJU95596  A/Shenzhen/SP58/2014(H7N9)           ..................................................
AJU95632  A/Shenzhen/SP62/2014(H7N9)           ..................................................
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FIGURE 22-42

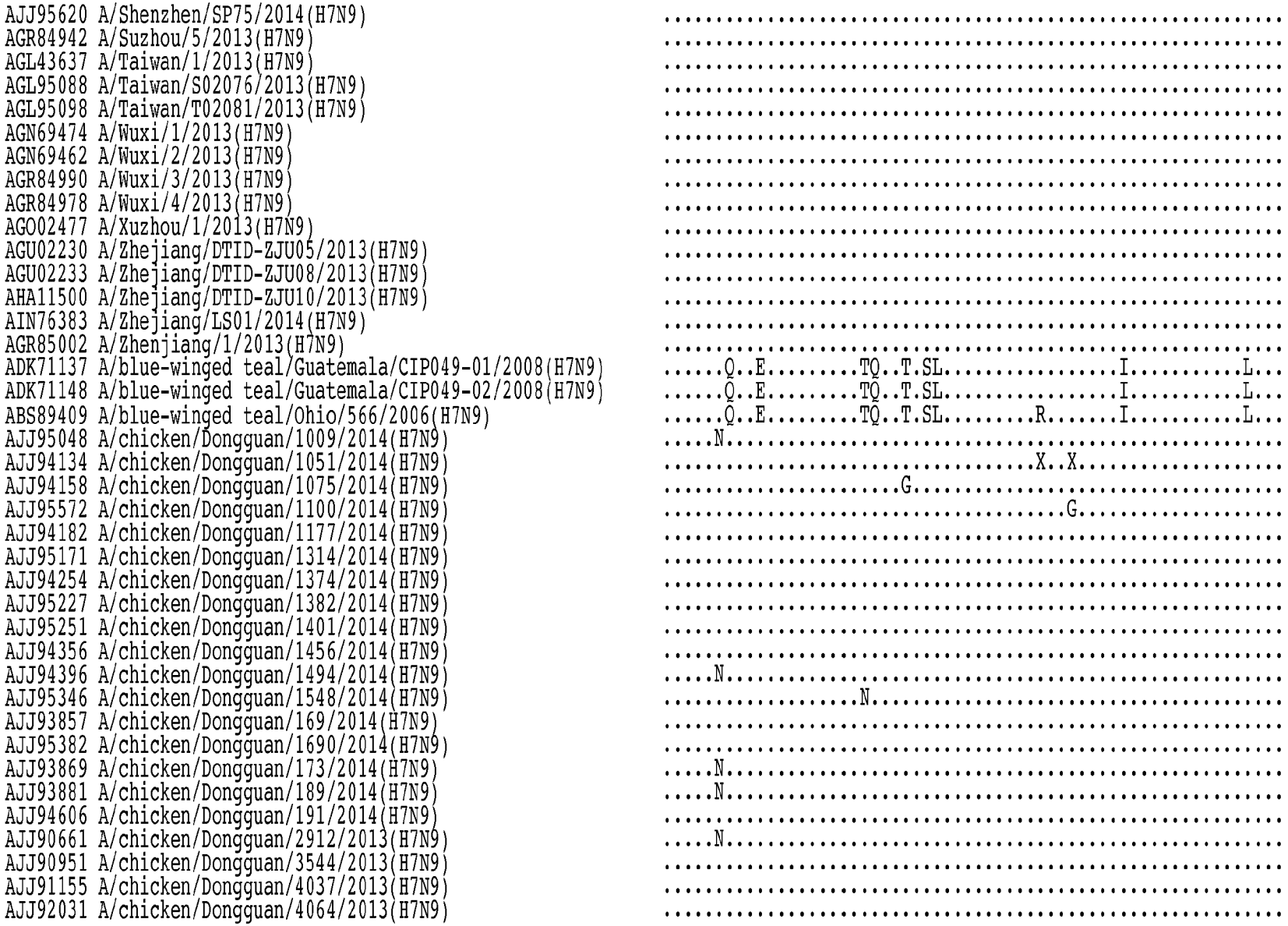

AJJ95620 A/Shenzhen/SP75/2014(H7N9)
AGR84942 A/Suzhou/5/2013(H7N9)
AGL43637 A/Taiwan/1/2013(H7N9)
AGL95088 A/Taiwan/S02076/2013(H7N9)
AGL95098 A/Taiwan/T02081/2013(H7N9)
AGN69474 A/Wuxi/1/2013(H7N9)
AGN69462 A/Wuxi/2/2013(H7N9)
AGR84990 A/Wuxi/3/2013(H7N9)
AGR84978 A/Wuxi/4/2013(H7N9)
AGO02477 A/Xuzhou/1/2013(H7N9)
AGU02230 A/Zhejiang/DTID-ZJU05/2013(H7N9)
AGU02233 A/Zhejiang/DTID-ZJU08/2013(H7N9)
AHA11500 A/Zhejiang/DTID-ZJU10/2013(H7N9)
AIN76383 A/Zhejiang/LS01/2014(H7N9)
AGR85002 A/Zhenjiang/1/2013(H7N9)
ADK71137 A/blue-winged teal/Guatemala/CIP049-01/2008(H7N9)
ADK71148 A/blue-winged teal/Guatemala/CIP049-02/2008(H7N9)
ABS89409 A/blue-winged teal/Ohio/566/2006(H7N9)
AJJ95048 A/chicken/Dongguan/1009/2014(H7N9)
AJJ94134 A/chicken/Dongguan/1051/2014(H7N9)
AJJ94158 A/chicken/Dongguan/1075/2014(H7N9)
AJJ95572 A/chicken/Dongguan/1100/2014(H7N9)
AJJ94182 A/chicken/Dongguan/1177/2014(H7N9)
AJJ95171 A/chicken/Dongguan/1314/2014(H7N9)
AJJ94254 A/chicken/Dongguan/1374/2014(H7N9)
AJJ95227 A/chicken/Dongguan/1382/2014(H7N9)
AJJ95251 A/chicken/Dongguan/1401/2014(H7N9)
AJJ94356 A/chicken/Dongguan/1456/2014(H7N9)
AJJ94396 A/chicken/Dongguan/1494/2014(H7N9)
AJJ95346 A/chicken/Dongguan/1548/2014(H7N9)
AJJ93857 A/chicken/Dongguan/169/2014(H7N9)
AJJ95382 A/chicken/Dongguan/1690/2014(H7N9)
AJJ93869 A/chicken/Dongguan/173/2014(H7N9)
AJJ93881 A/chicken/Dongguan/189/2014(H7N9)
AJJ94606 A/chicken/Dongguan/191/2014(H7N9)
AJJ90661 A/chicken/Dongguan/2912/2013(H7N9)
AJJ90951 A/chicken/Dongguan/3544/2013(H7N9)
AJJ91155 A/chicken/Dongguan/4037/2013(H7N9)
AJJ92031 A/chicken/Dongguan/4064/2013(H7N9)

FIGURE 22-43

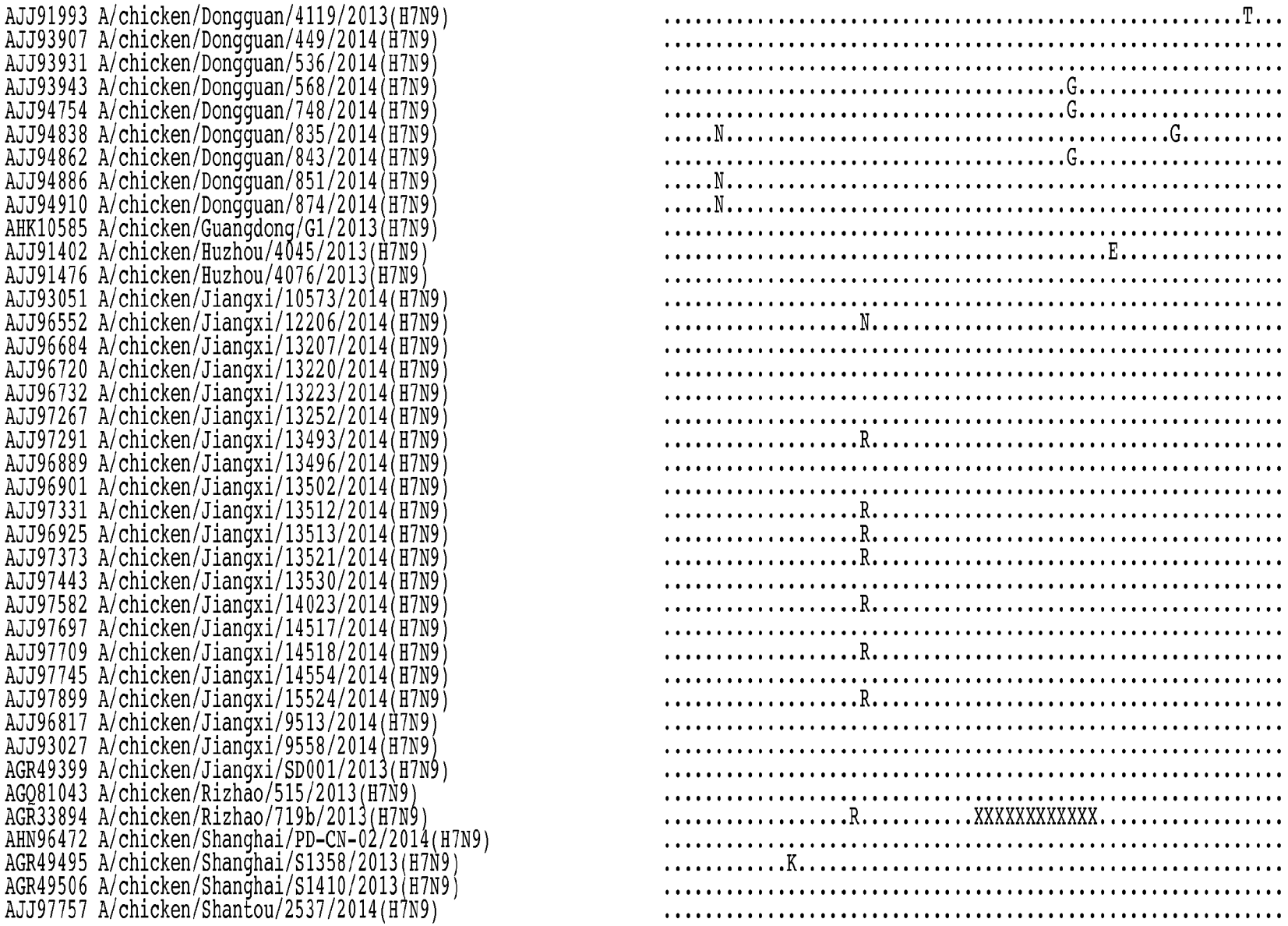

AJJ91993 A/chicken/Dongguan/4119/2013(H7N9)
AJJ93907 A/chicken/Dongguan/449/2014(H7N9)
AJJ93931 A/chicken/Dongguan/536/2014(H7N9)
AJJ93943 A/chicken/Dongguan/568/2014(H7N9)
AJJ94754 A/chicken/Dongguan/748/2014(H7N9)
AJJ94838 A/chicken/Dongguan/835/2014(H7N9)
AJJ94862 A/chicken/Dongguan/843/2014(H7N9)
AJJ94886 A/chicken/Dongguan/851/2014(H7N9)
AJJ94910 A/chicken/Dongguan/874/2014(H7N9)
AHK10585 A/chicken/Guangdong/G1/2013(H7N9)
AJJ91402 A/chicken/Huzhou/4045/2013(H7N9)
AJJ91476 A/chicken/Huzhou/4076/2013(H7N9)
AJJ93051 A/chicken/Jiangxi/10573/2014(H7N9)
AJJ96552 A/chicken/Jiangxi/12206/2014(H7N9)
AJJ96684 A/chicken/Jiangxi/13207/2014(H7N9)
AJJ96720 A/chicken/Jiangxi/13220/2014(H7N9)
AJJ96732 A/chicken/Jiangxi/13223/2014(H7N9)
AJJ97267 A/chicken/Jiangxi/13252/2014(H7N9)
AJJ97291 A/chicken/Jiangxi/13493/2014(H7N9)
AJJ96889 A/chicken/Jiangxi/13496/2014(H7N9)
AJJ96901 A/chicken/Jiangxi/13502/2014(H7N9)
AJJ97331 A/chicken/Jiangxi/13512/2014(H7N9)
AJJ96925 A/chicken/Jiangxi/13513/2014(H7N9)
AJJ97373 A/chicken/Jiangxi/13521/2014(H7N9)
AJJ97443 A/chicken/Jiangxi/13530/2014(H7N9)
AJJ97582 A/chicken/Jiangxi/14023/2014(H7N9)
AJJ97697 A/chicken/Jiangxi/14517/2014(H7N9)
AJJ97709 A/chicken/Jiangxi/14518/2014(H7N9)
AJJ97745 A/chicken/Jiangxi/14554/2014(H7N9)
AJJ97899 A/chicken/Jiangxi/15524/2014(H7N9)
AJJ96817 A/chicken/Jiangxi/9513/2014(H7N9)
AJJ93027 A/chicken/Jiangxi/9558/2014(H7N9)
AGR49399 A/chicken/Jiangxi/SD001/2013(H7N9)
AGQ81043 A/chicken/Rizhao/515/2013(H7N9)
AGR33894 A/chicken/Rizhao/719b/2013(H7N9)
AHN96472 A/chicken/Shanghai/PD-CN-02/2014(H7N9)
AGR49495 A/chicken/Shanghai/S1358/2013(H7N9)
AGR49506 A/chicken/Shanghai/S1410/2013(H7N9)
AJJ97757 A/chicken/Shantou/2537/2014(H7N9)
.....T...
.....G.....
.....G.....
.N.....G.....
.....G.....
.N.....
.N.....
.....E.....
.....N.....
.....R.....
.....R.....
.....R.....
.....R.....
.....R.....
.....R.....
.....R.....
.....R.....XXXXXXXXXXX.....
.K.....

FIGURE 22-44

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AJJ97973 A/chicken/Shantou/4325/2014(H7N9)
AJJ97998 A/chicken/Shantou/4816/2014(H7N9)
AJJ91314 A/chicken/Shaoxing/2417/2013(H7N9)
AJJ91725 A/chicken/Shaoxing/5201/2013(H7N9)
AJJ95464 A/chicken/Shenzhen/138/2014(H7N9)
AJJ90576 A/chicken/Shenzhen/1665/2013(H7N9)
AJJ90588 A/chicken/Shenzhen/2110/2013(H7N9)
AJJ91035 A/chicken/Shenzhen/3780/2013(H7N9)
AJJ90490 A/chicken/Shenzhen/742/2013(H7N9)
AJJ92005 A/chicken/Shenzhen/801/2013(H7N9)
AJJ90526 A/chicken/Shenzhen/898/2013(H7N9)
AIU47013 A/chicken/Suzhou/040201H/2013(H7N9)
AGJ72861 A/chicken/Zhejiang/DTID-ZJU01/2013(H7N9)
AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013(H7N9)
AGR49554 A/chicken/Zhejiang/SD033/2013(H7N9)
AGR49566 A/duck/Anhui/SC702/2013(H7N9)
BAN16711 A/duck/Gunma/466/2011(H7N9)
AJJ97841 A/duck/Jiangxi/15044/2014(H7N9)
AGQ80952 A/duck/Jiangxi/3096/2009(H7N9)
AGQ80989 A/duck/Jiangxi/3257/2009(H7N9)
BAH22785 A/duck/Mongolia/119/2008(H7N9)
BAN59727 A/duck/Mongolia/129/2010(H7N9)
BAN59726 A/duck/Mongolia/147/2008(H7N9)
AJK00354 A/duck/Zhejiang/LS02/2014(H7N9)
AFM09439 A/emperor goose/Alaska/44063-061/2006(H7N9)
AFV33947 A/goose/Nebraska/17097-4/2011(H7N9)
AFV33945 A/guinea fowl/Nebraska/17096-1/2011(H7N9)
AHA11441 A/guinea fowl/Nebraska/17096/2011(H7N9)
AGR49722 A/homing pigeon/Jiangsu/SD184/2013(H7N9)
AGL97639 A/mallard/Minnesota/AI09-3770/2009(H7N9)
AHZ42537 A/mallard/Minnesota/AI09-3770/2009(H7N9)
ADC29485 A/mallard/Spain/08.00991.3/2005(H7N9)
AHZ41929 A/mallard/Sweden/1621/2002(H7N9)
AAY46211 A/mallard/Sweden/91/2002(H7N9)
AGY42258 A/mallard/Sweden/91/2002(H7N9)
AGE08098 A/northern shoverl/Mississippi/11OS145/2011(H7N9)
AGR49734 A/pigeon/Shanghai/S1069/2013(H7N9)
ACD03594 A/ruddy turnstone/DE/1538/2000(H7N9)
AFX85260 A/ruddy turnstone/Delaware Bay/220/1995(H7N9)
```

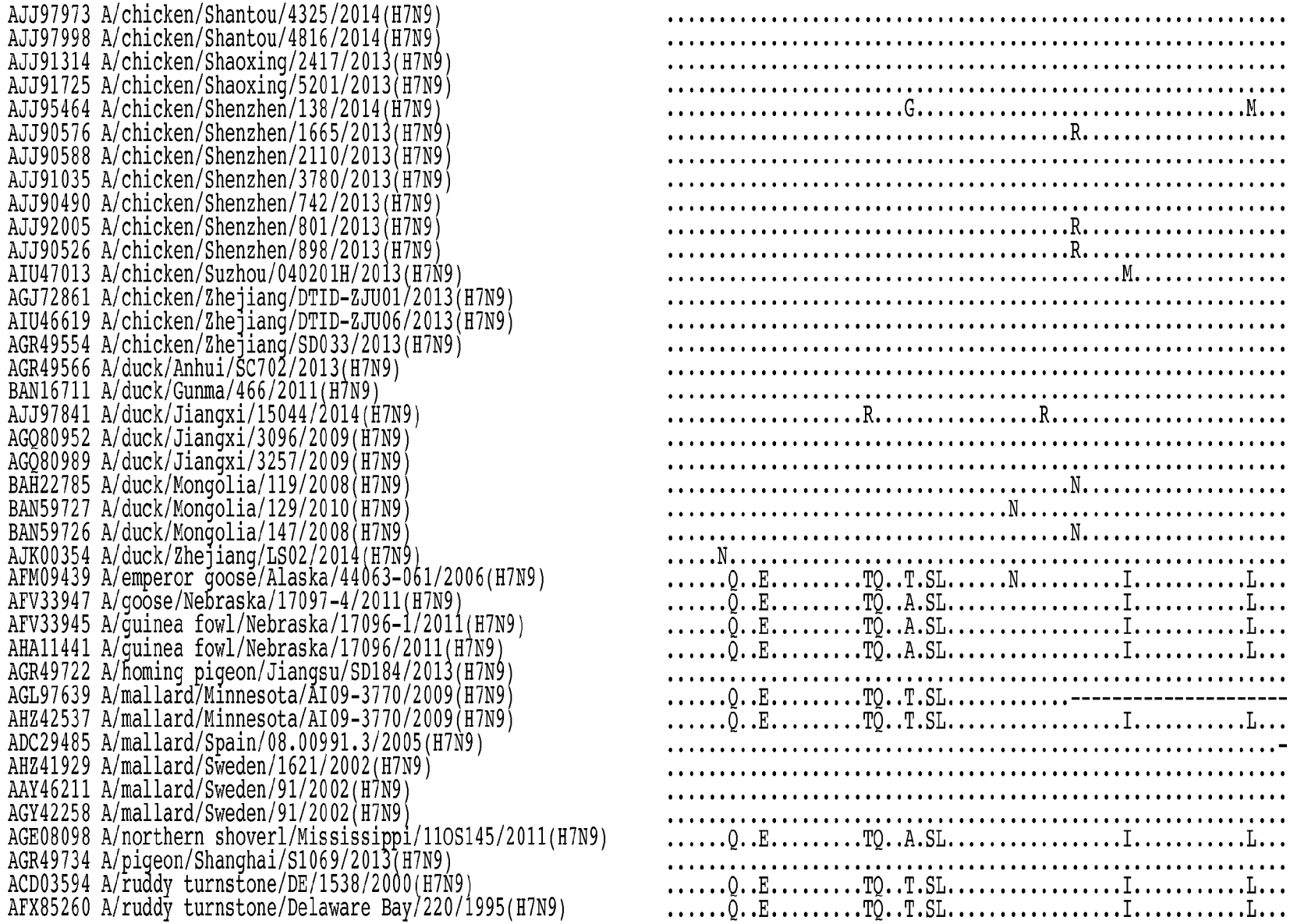


FIGURE 22-45

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AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000(H7N9)      .....Q.E.....TQ..T.SL.....I.....L...
AJJ94194 A/silkie chicken/Dongguan/1264/2014(H7N9)            .....G.....M...
AJJ94206 A/silkie chicken/Dongguan/1268/2014(H7N9)            ..........
AJJ94344 A/silkie chicken/Dongguan/1451/2014(H7N9)            ..........
AJJ95584 A/silkie chicken/Dongguan/1519/2014(H7N9)            .....G.....M...
AJJ93845 A/silkie chicken/Dongguan/157/2014(H7N9)             ....N.....
AJJ90673 A/silkie chicken/Dongguan/3049/2013(H7N9)            ....N.....
AJJ90795 A/silkie chicken/Dongguan/3281/2013(H7N9)            ..........
AJJ90891 A/silkie chicken/Dongguan/3520/2013(H7N9)            ....N.....
AJJ91264 A/silkie chicken/Dongguan/4129/2013(H7N9)            ..........
AJJ93979 A/silkie chicken/Dongguan/656/2014(H7N9)             ....N.....
AJJ94959 A/silkie chicken/Dongguan/967/2014(H7N9)             ....N.....
AJJ92967 A/silkie chicken/Jiangxi/9469/2014(H7N9)             ..........
AJJ97925 A/silkie chicken/Shantou/2050/2014(H7N9)             ..........
AJJ90538 A/silkie chicken/Shenzhen/918/2013(H7N9)             ..........
AEK84765 A/spot-billed duck/Korea/447/2011(H7N9)              .....R.....
AGW82588 A/tree sparrow/Shanghai/01/2013(H7N9)                ..........
ABI84694 A/turkey/Minnesota/1/1988(H7N9)                      .....Q.E.....AQ..A.SL.....I.....L...
AHA11483 A/turkey/Minnesota/14135-2/2009(H7N9)                .....Q.E.....TQ..K.SL.....I.....L...
AHA57050 A/turkey/Minnesota/14659/2009(H7N9)                  .N...Q.E.....TQ..K.SL.....I.....L...
AHA57072 A/turkey/Minnesota/18421/2009(H7N9)                  .....Q.E.....TQ..K.SL.....I.....L...
AHA11472 A/turkey/Minnesota/31676/2009(H7N9)                  .....Q.E.....TQ..K.SL.....I.....L...
AHA11461 A/turkey/Minnesota/31900/2011(H7N9)                  .....Q.E.....TQ..A.SL.....I.....L...
AHA11452 A/turkey/Minnesota/32710/2011(H7N9)                  .....Q.E.....TQ..A.SL.....I.....L...
AAM19228 A/turkey/Minnesota/38429/1988(H7N9)                  ----------
ACZ48625 A/turkey/Minnesota/38429/1988(H7N9)                  ----------
AEK84760 A/wild bird/Korea/A14/2011(H7N9)                     ..........
AEK84761 A/wild bird/Korea/A3/2011(H7N9)                      ..........
AEK84763 A/wild bird/Korea/A9/2011(H7N9)                      ..........
AGG53377 A/wild duck/Korea/CSM42-1/2011(H7N9)                 .....R.....
AGG53366 A/wild duck/Korea/CSM42-34/2011(H7N9)                .....R.....
AGG53432 A/wild duck/Korea/MHC35-41/2011(H7N9)                ..........
AGG53399 A/wild duck/Korea/MHC39-26/2011(H7N9)                ..........
AGG53476 A/wild duck/Korea/SH19-27/2010(H7N9)                 ..........
AGG53487 A/wild duck/Korea/SH19-50/2010(H7N9)                 ..........
AGG53520 A/wild duck/Korea/SH20-27/2008(H7N9)                 .....N.....
AEM98291 A/wild duck/Mongolia/1-241/2008(H7N9)                .....N.....
AGR49770 A/wild pigeon/Jiangsu/SD001/2013(H7N9)               ..........
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FIGURE 22-46

Positions from 541 till 561

	Consensus sequence	**IVMGLVFICVKNGNMRCTICI**
ACX53683	A/goose/Czech Republic/1848-K9/2009(H7N9)	.A.....
ADN34727	A/goose/Czech Republic/1848-T14/2009(H7N9)	.A.....
CAY39406	A/Anas crecca/Spain/1460/2008(H7N9)	.A.....
AHZ39686	A/Anhui/DEWH72-01/2013(H7N9)	.....
AHZ39710	A/Anhui/DEWH72-03/2013(H7N9)	.....
AHZ39746	A/Anhui/DEWH72-06/2013(H7N9)	.....
AHM24224	A/Beijing/3/2013(H7N9)	.....
AHD25003	A/Guangdong/02/2013(H7N9)	.....------
AHH25185	A/Guangdong/04/2013(H7N9)	.....
AGI60301	A/Hangzhou/1/2013(H7N9)	.....
AGK84857	A/Hangzhou/2/2013(H7N9)	.....
AHF20528	A/Hong Kong/470129/2013(H7N9)	.....
AGY41893	A/Huizhou/01/2013(H7N9)	.....
AGO51387	A/Jiangsu/2/2013(H7N9)	.....
AGJ73503	A/Nanjing/1/2013(H7N9)	.....
AGR85026	A/Nanjing/2/2013(H7N9)	.....SR.....
AGR84954	A/Nanjing/6/2013(H7N9)	.....R.....
AHK10800	A/Shanghai/01/2014(H7N9)	.....
AGL44438	A/Shanghai/02/2013(H7N9)	.....
AGL33692	A/Shanghai/4655T/2013(H7N9)	.A.....
AGL33693	A/Shanghai/4659T/2013(H7N9)	.....
AGI60292	A/Shanghai/4664T/2013(H7N9)	.....
AGM53887	A/Shanghai/4701T/2013(H7N9)	.....-
AGM53886	A/Shanghai/4842T/2013(H7N9)	.....---
AGM53883	A/Shanghai/5083T/2013(H7N9)	.....---
AGM53884	A/Shanghai/5180T/2013(H7N9)	.....
AGM53885	A/Shanghai/5240T/2013(H7N9)	.....---
AGW82600	A/Shanghai/CN01/2013(H7N9)	.....
AHF20568	A/Shanghai/CN02/2013(H7N9)	.....
AGW82612	A/Shanghai/JS01/2013(H7N9)	.A.....
AID70634	A/Shanghai/Mix1/2014(H7N9)	.....
AHJ57411	A/Shanghai/PD-01/2014(H7N9)	.....
AHJ57418	A/Shanghai/PD-02/2014(H7N9)	.....
AJJ96841	A/Shenzhen/SP139/2014(H7N9)	.....
AJJ91909	A/Shenzhen/SP26/2014(H7N9)	.....
AJJ91945	A/Shenzhen/SP38/2014(H7N9)	.....
AJJ91885	A/Shenzhen/SP4/2014(H7N9)	.....

FIGURE 22-47

```
AJU91957  A/Shenzhen/SP44/2014(H7N9)                  ..............................
AJU91969  A/Shenzhen/SP48/2014(H7N9)                  ..............................
AJU95596  A/Shenzhen/SP58/2014(H7N9)                  ..............................
AJU95632  A/Shenzhen/SP62/2014(H7N9)                  ..............................
AJU95620  A/Shenzhen/SP75/2014(H7N9)                  V.............................
AGR84942  A/Suzhou/5/2013(H7N9)                       ..............................
AGL43637  A/Taiwan/1/2013(H7N9)                       ..............................
AGL95088  A/Taiwan/S02076/2013(H7N9)                  ..............-----.............
AGL95098  A/Taiwan/T02081/2013(H7N9)                  ..............---.............
AGN69474  A/Wuxi/1/2013(H7N9)                        ..............................
AGN69462  A/Wuxi/2/2013(H7N9)                        ..............................
AGR84990  A/Wuxi/3/2013(H7N9)                        ..............SR..............
AGR84978  A/Wuxi/4/2013(H7N9)                        ..............SR..............
AGO02477  A/Xuzhou/1/2013(H7N9)                      ..............SR..............
AGU02230  A/Zhejiang/DTID-ZJU05/2013(H7N9)           ..............---.............
AGU02233  A/Zhejiang/DTID-ZJU08/2013(H7N9)           ..............---.............
AHA11500  A/Zhejiang/DTID-ZJU10/2013(H7N9)           ..............--------........
AIN76383  A/Zhejiang/LS01/2014(H7N9)                 ..............................
AGR85002  A/Zhejiang/1/2013(H7N9)                    ..............SR.K............
ADK71137  A/blue-winged teal/Guatemala/CIP049-01/2008(H7N9)  A.....I.....................
ADK71148  A/blue-winged teal/Guatemala/CIP049-02/2008(H7N9)  A.....I.....................
ABS89409  A/blue-winged teal/Ohio/566/2006(H7N9)    A.....I.....................
AJU95048  A/chicken/Dongguan/1009/2014(H7N9)        ..............................
AJU94134  A/chicken/Dongguan/1051/2014(H7N9)        ..............................
AJU94158  A/chicken/Dongguan/1075/2014(H7N9)        ..............................
AJU95572  A/chicken/Dongguan/1100/2014(H7N9)        ..............................
AJU94182  A/chicken/Dongguan/1177/2014(H7N9)        ..............................
AJU95171  A/chicken/Dongguan/1314/2014(H7N9)        ..............................
AJU94254  A/chicken/Dongguan/1374/2014(H7N9)        ..............................
AJU95227  A/chicken/Dongguan/1382/2014(H7N9)        ..............................
AJU95251  A/chicken/Dongguan/1401/2014(H7N9)        ..............................
AJU94356  A/chicken/Dongguan/1456/2014(H7N9)        ..............................
AJU94396  A/chicken/Dongguan/1494/2014(H7N9)        ..............................
AJU95346  A/chicken/Dongguan/1548/2014(H7N9)        ..............................
AJU93857  A/chicken/Dongguan/169/2014(H7N9)         ..............................
AJU95382  A/chicken/Dongguan/1690/2014(H7N9)        ..............................
AJU93869  A/chicken/Dongguan/173/2014(H7N9)         ..............................
AJU93881  A/chicken/Dongguan/189/2014(H7N9)         ..............................
AJU94606  A/chicken/Dongguan/191/2014(H7N9)         ..............................
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FIGURE 22-48

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AUJ90661 A/chicken/Dongguan/2912/2013(H7N9)  ..............................
AUJ90951 A/chicken/Dongguan/3544/2013(H7N9)  ..............................
AUJ91155 A/chicken/Dongguan/4037/2013(H7N9)  ..............................
AUJ92031 A/chicken/Dongguan/4064/2013(H7N9)  ..............................
AUJ91993 A/chicken/Dongguan/4119/2013(H7N9)  ..............................
AUJ93907 A/chicken/Dongguan/449/2014(H7N9)   ..............................
AUJ93931 A/chicken/Dongguan/536/2014(H7N9)   ..............................
AUJ93943 A/chicken/Dongguan/568/2014(H7N9)   ..............................
AUJ94754 A/chicken/Dongguan/748/2014(H7N9)   ..............................
AUJ94838 A/chicken/Dongguan/835/2014(H7N9)   ..............................
AUJ94862 A/chicken/Dongguan/843/2014(H7N9)   ..............................
AUJ94886 A/chicken/Dongguan/851/2014(H7N9)   ..............................
AUJ94910 A/chicken/Dongguan/874/2014(H7N9)   ..............................
AHK10585 A/chicken/Guangdong/G1/2013(H7N9)   ..............................
AUJ91402 A/chicken/Huzhou/4045/2013(H7N9)    ..............................
AUJ91476 A/chicken/Huzhou/4076/2013(H7N9)    ..............................
AUJ93051 A/chicken/Jiangxi/10573/2014(H7N9)  ..............................
AUJ96552 A/chicken/Jiangxi/12206/2014(H7N9)  ..............................
AUJ96684 A/chicken/Jiangxi/13207/2014(H7N9)  ..............................
AUJ96720 A/chicken/Jiangxi/13220/2014(H7N9)  ..............................
AUJ96732 A/chicken/Jiangxi/13223/2014(H7N9)  ..............................
AUJ97267 A/chicken/Jiangxi/13252/2014(H7N9)  ..............................
AUJ97291 A/chicken/Jiangxi/13493/2014(H7N9)  ..............................
AUJ96889 A/chicken/Jiangxi/13496/2014(H7N9)  ..............................
AUJ96901 A/chicken/Jiangxi/13502/2014(H7N9)  ..............................
AUJ97331 A/chicken/Jiangxi/13512/2014(H7N9)  ..............................
AUJ96925 A/chicken/Jiangxi/13513/2014(H7N9)  ..............................
AUJ97373 A/chicken/Jiangxi/13521/2014(H7N9)  ..............................
AUJ97443 A/chicken/Jiangxi/13530/2014(H7N9)  ..............................
AUJ97582 A/chicken/Jiangxi/14023/2014(H7N9)  ..............................
AUJ97697 A/chicken/Jiangxi/14517/2014(H7N9)  ..............................
AUJ97709 A/chicken/Jiangxi/14518/2014(H7N9)  ..............................
AUJ97745 A/chicken/Jiangxi/14554/2014(H7N9)  ..............................
AUJ97899 A/chicken/Jiangxi/15524/2014(H7N9)  .....M........................
AUJ96817 A/chicken/Jiangxi/9513/2014(H7N9)   ..............................
AUJ93027 A/chicken/Jiangxi/9558/2014(H7N9)   ..............................
AGR49399 A/chicken/Jiangxi/SD001/2013(H7N9)  ..............................
AGQ81043 A/chicken/Rizhao/515/2013(H7N9)     ..............................
AGR33894 A/chicken/Rizhao/719b/2013(H7N9)    ..............................
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FIGURE 22-49

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AHN96472 A/chicken/Shanghai/PD-CN-02/2014(H7N9)  ..............................
AGR49495 A/chicken/Shanghai/S1358/2013(H7N9)     ..............................
AGR49506 A/chicken/Shanghai/S1410/2013(H7N9)     ..............................
AJJ97757 A/chicken/Shantou/2537/2014(H7N9)       ..............................
AJJ97973 A/chicken/Shantou/4325/2014(H7N9)       ..............................
AJJ97998 A/chicken/Shantou/4816/2014(H7N9)       ..............................
AJJ91314 A/chicken/Shaoxing/2417/2013(H7N9)      ..............................
AJJ91725 A/chicken/Shaoxing/5201/2013(H7N9)      ..............................
AJJ95464 A/chicken/Shenzhen/138/2014(H7N9)       ..............................
AJJ90576 A/chicken/Shenzhen/1665/2013(H7N9)      ..............................
AJJ90588 A/chicken/Shenzhen/2110/2013(H7N9)      ..............................
AJJ91035 A/chicken/Shenzhen/3780/2013(H7N9)      ..............................
AJJ90490 A/chicken/Shenzhen/742/2013(H7N9)       ..............................
AJJ92005 A/chicken/Shenzhen/801/2013(H7N9)       ..............................
AJJ90526 A/chicken/Shenzhen/898/2013(H7N9)       ..............................
AIU47013 A/chicken/Suzhou/040201H/2013(H7N9)     ..............................
AGJ72861 A/chicken/Zhejiang/DTID-ZJU01/2013(H7N9) ..............................
AIU46619 A/chicken/Zhejiang/DTID-ZJU06/2013(H7N9) ..............................
AGR49554 A/chicken/Zhejiang/SD033/2013(H7N9)     ..............................
AGR49566 A/duck/Anhui/SC702/2013(H7N9)           ..............................
BAN16711 A/duck/Gunma/466/2011(H7N9)             .A............................
AJJ97841 A/duck/Jiangxi/15044/2014(H7N9)         ..............................
AGQ80952 A/duck/Jiangxi/3096/2009(H7N9)          .A............................
AGQ80989 A/duck/Jiangxi/3257/2009(H7N9)          .A............................
BAH22785 A/duck/Mongolia/119/2008(H7N9)          .A............................
BAN59727 A/duck/Mongolia/129/2010(H7N9)          .A............................
BAN59726 A/duck/Mongolia/147/2008(H7N9)          .A............................
AJK00354 A/duck/Zhejiang/LS02/2014(H7N9)         ..............................
AFM09439 A/emperor goose/Alaska/44063-061/2006(H7N9) .A.........I..............
AFV33947 A/goose/Nebraska/17097-4/2011(H7N9)     .A.........I..............
AFV33945 A/guinea fowl/Nebraska/17096-1/2011(H7N9) .A.........I..............
AHA11441 A/guinea fowl/Nebraska/17096/2011(H7N9) .A.........I..............
AGR49722 A/homing pigeon/Jiangsu/SD184/2013(H7N9) ..............................
AGL97639 A/mallard/Minnesota/AI09-3770/2009(H7N9) ------------------------------
AHZ42537 A/mallard/Minnesota/AI09-3770/2009(H7N9) .A.......I..................
ADC29485 A/mallard/Spain/08.00991.3/2005(H7N9)   ------------------------------
AHZ41929 A/mallard/Sweden/1621/2002(H7N9)        .A.....M.....................
AAY46211 A/mallard/Sweden/91/2002(H7N9)          .A.....M.....................
AGY42258 A/mallard/Sweden/91/2002(H7N9)          .A.....M.....................
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FIGURE 22-50

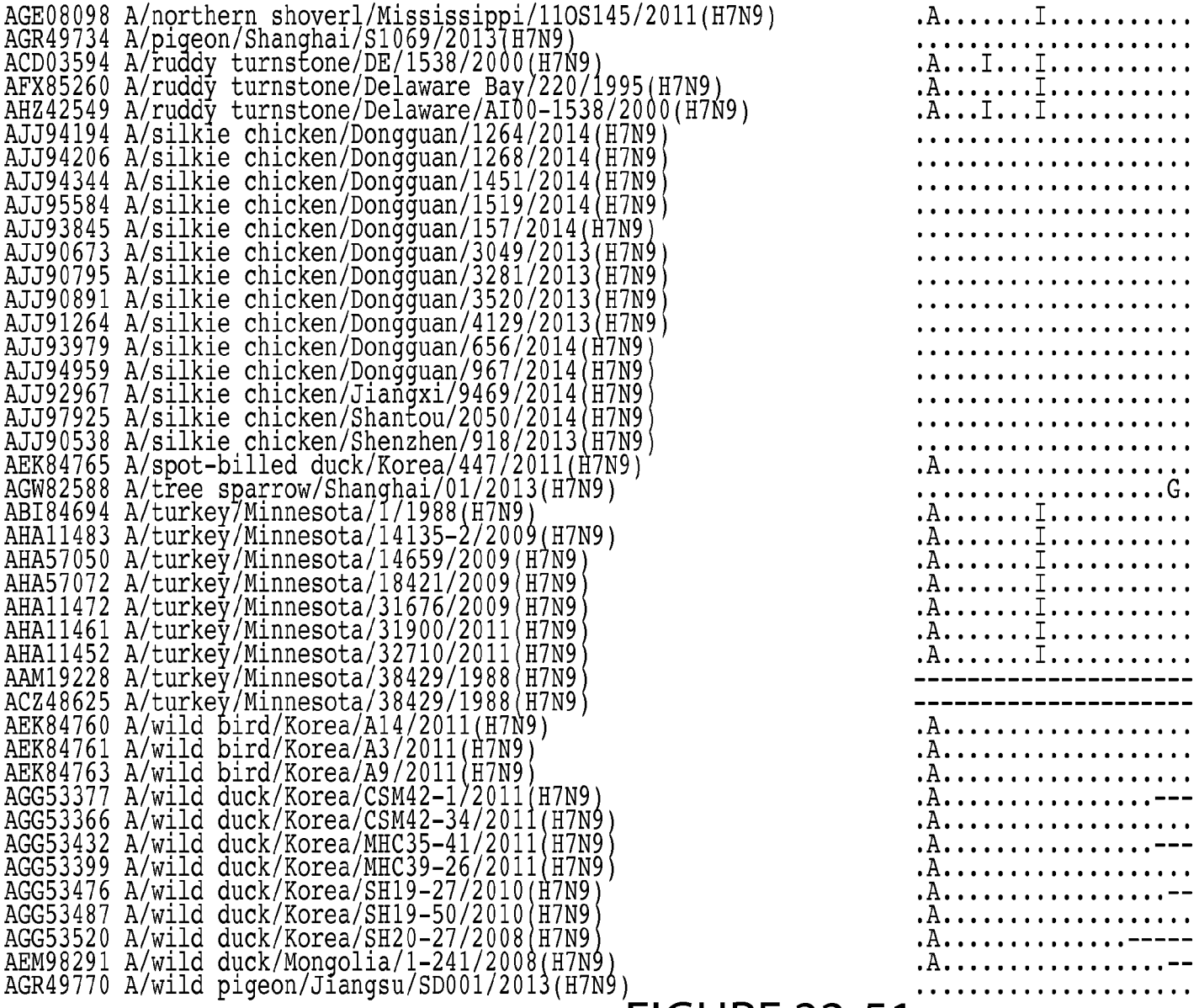

AGE08098 A/northern shoverl/Mississippi/11OS145/2011(H7N9) .A.....I.....
AGR49734 A/pigeon/Shanghai/S1069/2013(H7N9) .....
ACD03594 A/ruddy turnstone/DE/1538/2000(H7N9) .A...I...I.....
AFX85260 A/ruddy turnstone/Delaware Bay/220/1995(H7N9) .A.....I.....
AHZ42549 A/ruddy turnstone/Delaware/AI00-1538/2000(H7N9) .A...I...I.....
AJJ94194 A/silkie chicken/Dongguan/1264/2014(H7N9) .....
AJJ94206 A/silkie chicken/Dongguan/1268/2014(H7N9) .....
AJJ94344 A/silkie chicken/Dongguan/1451/2014(H7N9) .....
AJJ95584 A/silkie chicken/Dongguan/1519/2014(H7N9) .....
AJJ93845 A/silkie chicken/Dongguan/157/2014(H7N9) .....
AJJ90673 A/silkie chicken/Dongguan/3049/2013(H7N9) .....
AJJ90795 A/silkie chicken/Dongguan/3281/2013(H7N9) .....
AJJ90891 A/silkie chicken/Dongguan/3520/2013(H7N9) .....
AJJ91264 A/silkie chicken/Dongguan/4129/2013(H7N9) .....
AJJ93979 A/silkie chicken/Dongguan/656/2014(H7N9) .....
AJJ94959 A/silkie chicken/Dongguan/967/2014(H7N9) .....
AJJ92967 A/silkie chicken/Jiangxi/9469/2014(H7N9) .....
AJJ97925 A/silkie chicken/Shantou/2050/2014(H7N9) .....
AJJ90538 A/silkie chicken/Shenzhen/918/2013(H7N9) .....
AEK84765 A/spot-billed duck/Korea/447/2011(H7N9) .A.....
AGW82588 A/tree sparrow/Shanghai/01/2013(H7N9) .....G.
ABI84694 A/turkey/Minnesota/1/1988(H7N9) .A.....I.....
AHA11483 A/turkey/Minnesota/14135-2/2009(H7N9) .A.....I.....
AHA57050 A/turkey/Minnesota/14659/2009(H7N9) .A.....I.....
AHA57072 A/turkey/Minnesota/18421/2009(H7N9) .A.....I.....
AHA11472 A/turkey/Minnesota/31676/2009(H7N9) .A.....I.....
AHA11461 A/turkey/Minnesota/31900/2011(H7N9) .A.....I.....
AHA11452 A/turkey/Minnesota/32710/2011(H7N9) .A.....I.....
AAM19228 A/turkey/Minnesota/38429/1988(H7N9) ----------
ACZ48625 A/turkey/Minnesota/38429/1988(H7N9) ----------
AEK84760 A/wild bird/Korea/A14/2011(H7N9) .A.....
AEK84761 A/wild bird/Korea/A3/2011(H7N9) .A.....
AEK84763 A/wild bird/Korea/A9/2011(H7N9) .A.....
AGG53377 A/wild duck/Korea/CSM42-1/2011(H7N9) .A.....---
AGG53366 A/wild duck/Korea/CSM42-34/2011(H7N9) .A.....
AGG53432 A/wild duck/Korea/MHC35-41/2011(H7N9) .A.....---
AGG53399 A/wild duck/Korea/MHC39-26/2011(H7N9) .A.....
AGG53476 A/wild duck/Korea/SH19-27/2010(H7N9) .A.....--
AGG53487 A/wild duck/Korea/SH19-50/2010(H7N9) .A.....
AGG53520 A/wild duck/Korea/SH20-27/2008(H7N9) .A.....-----
AEM98291 A/wild duck/Mongolia/1-241/2008(H7N9) .A.....--
AGR49770 A/wild pigeon/Jiangsu/SD001/2013(H7N9) .....

FIGURE 22-51

Positions from 1 till 60

	Consensus sequence	**MYKIVVIIALLGAVKGLDKICLGHHAVANGTIVKTLTNEQEEVTNATETVESTGLNRLCM**
AHN97094	A/American green-winged teal/Ohio/13OS1869/2013(H10N8)	.....LVL......H..........P..........K..........KS.DK...
AHM24205	A/Eurasian coot/Germany/R411/2010(H10N8)	.......VT.....R.............T.................S..K...
AHK10761	A/Jiangxi/IPB13/2013(H10N8)	....I.....
AHK10763	A/Jiangxi/IPB13b/2013(H10N8)	....I.....
AJD10539	A/chicken/Jiangxi/1204/2014(H10N8)	....I.....
AJD11559	A/chicken/Jiangxi/1232/2014(H10N8)	...V...I.....
AJD11571	A/chicken/Jiangxi/1270/2014(H10N8)	....I.....
AJD11583	A/chicken/Jiangxi/1279/2014(H10N8)	....I.....
AJD09903	A/chicken/Jiangxi/18399/2013(H10N8)	....
AJD09915	A/chicken/Jiangxi/18933/2013(H10N8)	....
AJD09927	A/chicken/Jiangxi/19924/2013(H10N8)	....
AJD09939	A/chicken/Jiangxi/19960/2013(H10N8)	....
AJD10551	A/chicken/Jiangxi/23252/2013(H10N8)	....K...............
AJD10563	A/chicken/Jiangxi/23259/2013(H10N8)	....K...............
AJD10575	A/chicken/Jiangxi/23274/2013(H10N8)	....I.....
AJD10587	A/chicken/Jiangxi/27600/2013(H10N8)	....I...I.....
AJD10611	A/chicken/Jiangxi/28113/2013(H10N8)	....I.....
AJD11595	A/chicken/Jiangxi/3152/2014(H10N8)	....I.....
AJD10683	A/chicken/Jiangxi/34609/2013(H10N8)	....I.....
AJD11271	A/chicken/Jiangxi/36229/2013(H10N8)	...V...I.....
AIK26172	A/chicken/Jiangxi/JXA132717/2014(H10N8)	-----...I.....
AFJ19053	A/duck/Guangdong/E1/2012(H10N8)	..........R.....................................S.....
BAG16719	A/duck/Hokkaido/W95/2006(H10N8)	----------
AJD14343	A/duck/Jiangxi/2039/2005(H10N8)	....S.....
AJD14367	A/duck/Jiangxi/22960/2008(H10N8)	....AS.....
AJD14379	A/duck/Jiangxi/4759/2009(H10N8)	..............................I...........AS.....
AJD10887	A/duck/Jiangxi/6450/2013(H10N8)	....
AJD10911	A/duck/Jiangxi/6474/2013(H10N8)	....
AJD10923	A/duck/Jiangxi/6487/2013(H10N8)	................I...
AJD10971	A/duck/Jiangxi/6526/2013(H10N8)	....
AJD10995	A/duck/Jiangxi/6544/2013(H10N8)	........................V.................................
AJD11055	A/duck/Jiangxi/6602/2013(H10N8)	....
AJD11091	A/duck/Jiangxi/6645/2013(H10N8)	....X.....
AIK26225	A/duck/Jiangxi/JXA132712/2014(H10N8)	------...I.....
AIK26196	A/duck/Jiangxi/JXA132727/2014(H10N8)	-------..I.....
ABO52082	A/longtail duck/Maryland/295/2005(H10N8)	.....LVL......H..........P..........K..........KS.DK...
ADU16029	A/mallard/Interior Alaska/6MP0758/2006(H10N8)	.....LVL..F...H..........P..........K..........KS.DK...

FIGURE 23-1

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AER09406  A/mallard/Korea/1041/2010(H10N8)                    ----------------------------------------------------------
AER09416  A/mallard/Korea/1203/2010(H10N8)                    ..........T.....................................KS......
AHZ38995  A/mallard/Sweden/233/2002(H10N8)                    ...V.........R.....................................KS.GK...
ADE75294  A/mallard/Sweden/7/2003(H10N8)                      .........I.....................V.......H..........N.S..K...
AJD14391  A/migratory duck/Jiangxi/21248/2009(H10N8)          ..................................................S......
AJD14355  A/migratory duck/Jiangxi/593/2005(H10N8)           ..................................................S......
ACK43693  A/northern pintail/California/44221-656/2006(H10N8) .....LVL......H...........S...........K...........KS.DK...
AEK49546  A/northern shoveler/California/9235/2008(H10N8)     .....PVL......H...........P...........K...........GS.DK...
ACZ48518  A/quail/Italy/1117/1965(H10N8)                     ------------------------.....T...................S..K...
ABI84499  A/quail/Italy/1117/1965(H10N8)                     ...V.........R.............T...................S..K...
AEA04419  A/quail/Italy/1117/1965(H10N8)                     ...V.........R.............T...................S..K...
AGU01953  A/ruddy shelduck/Mongolia/1602/2010(H10N8)         .D.K..V.....................................................S......
AGL58508  A/ruddy turnstone/New Jersey/1148668/2004(H10N8)  .....LVL......H...........P...........K...........KS.DK...
AGG28633  A/shorebird/Delaware Bay/379/2008(H10N8)          .....LVL......H...........S...........K...........KN.D....
AGU01957  A/surf scoter/Mongolia/878V/2009(H10N8)           ......V.....................................................S......
AGU01970  A/velvet scoter/Mongolia/879V/2009(H10N8)        ......V.....................................................S......

Positions from 61 till 120
          Consensus sequence                                  KGRKHKDLGNCHPIGMLIGTPACDLHLTGTWDTLIERENAIAYCYPGATVNEEALRQKIM
AHN97094  A/American green-winged teal/Ohio/13OS1869/2013(H10N8) .S.NY..........I.....................D.S.....................
AHM24205  A/Eurasian coot/Germany/R411/2010(H10N8)           R..SY....R.....I.L...V.E.....A.....................I.........
AHK10761  A/Jiangxi/IPB13/2013(H10N8)                       ...........................M.....................V.........
AHK10763  A/Jiangxi/IPB13b/2013(H10N8)                      ...........................M.....................V.........
AJD10539  A/chicken/Jiangxi/1204/2014(H10N8)                ..................................................V.........
AJD11559  A/chicken/Jiangxi/1232/2014(H10N8)                ..................................................V.........
AJD11571  A/chicken/Jiangxi/1270/2014(H10N8)                ..............................................A.V.........
AJD11583  A/chicken/Jiangxi/1279/2014(H10N8)                ..................................................V.........
AJD09903  A/chicken/Jiangxi/18399/2013(H10N8)               ..................................................V.........
AJD09915  A/chicken/Jiangxi/18933/2013(H10N8)               ..................................................V.........
AJD09927  A/chicken/Jiangxi/19924/2013(H10N8)               ..................................................V.........
AJD09939  A/chicken/Jiangxi/19960/2013(H10N8)               ..................................................V.........
AJD10551  A/chicken/Jiangxi/23252/2013(H10N8)               ..................S...............................G.........
AJD10563  A/chicken/Jiangxi/23259/2013(H10N8)               ...............................I..................V.........
AJD10575  A/chicken/Jiangxi/23274/2013(H10N8)               ..................................................V.........
AJD10587  A/chicken/Jiangxi/27600/2013(H10N8)               ..................................................V.........
AJD10611  A/chicken/Jiangxi/28113/2013(H10N8)               ..................................................V.........
AJD11595  A/chicken/Jiangxi/3152/2014(H10N8)                ..................................................V.........
AJD10683  A/chicken/Jiangxi/34609/2013(H10N8)               ..............................................S.....V.........
AJD11271  A/chicken/Jiangxi/36229/2013(H10N8)               ..................................................V.........
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FIGURE 23-2

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AIK26172  A/chicken/Jiangxi/JXA132717/2014(H10N8)                    .....................................................V........
AFJ19053  A/duck/Guangdong/E1/2012(H10N8)                            ...N.................T.................V.................R..
BAG16719  A/duck/Hokkaido/W95/2006(H10N8)                            -----------------------------------------------------------
AJD14343  A/duck/Jiangxi/2039/2005(H10N8)                            ...S.....................................................
AJD14367  A/duck/Jiangxi/22960/2008(H10N8)                           ...R.....................................................V.
AJD14379  A/duck/Jiangxi/4759/2009(H10N8)                            ...R.....................................................V.
AJD10887  A/duck/Jiangxi/6450/2013(H10N8)                            ...N.....................................................
AJD10911  A/duck/Jiangxi/6474/2013(H10N8)                            ...N.....................................T...............
AJD10923  A/duck/Jiangxi/6487/2013(H10N8)                            ...N.....................................................
AJD10971  A/duck/Jiangxi/6526/2013(H10N8)                            ...N.....................................................
AJD10995  A/duck/Jiangxi/6544/2013(H10N8)                            ...N.....................................................
AJD11055  A/duck/Jiangxi/6602/2013(H10N8)                            ...N.....................................T...............
AJD11091  A/duck/Jiangxi/6645/2013(H10N8)                            ...N.....................................................
AIK26225  A/duck/Jiangxi/JXA132712/2014(H10N8)                       .....................................................V........
AIK26196  A/duck/Jiangxi/JXA132727/2014(H10N8)                       .....................................................V........
ABO52082  A/longtail duck/Maryland/295/2005(H10N8)                   .N.NY..........VV..................D.S.........S.........
ADU16029  A/mallard/Interior Alaska/6MP0758/2006(H10N8)              .S.NY..........V...................D.S.....................
AER09406  A/mallard/Korea/1041/2010(H10N8)                           -----------------------------------------------------------
AER09416  A/mallard/Korea/1203/2010(H10N8)                           ...N.....................................................I.........
AHZ38995  A/mallard/Sweden/233/2002(H10N8)                           ...SYN.........I.....................V...................
ADE75294  A/mallard/Sweden/7/2003(H10N8)                            ...S.................V...................................
AJD14391  A/migratory duck/Jiangxi/21248/2009(H10N8)                 ...N.....................................................I.........
AJD14355  A/migratory duck/Jiangxi/593/2005(H10N8)                   ...S.....................................................
ACK43693  A/northern pintail/California/44221-656/2006(H10N8)       .S.NY..........V...................D.S.....................
AEK49546  A/northern shoveler/California/9235/2008(H10N8)           .S.NY..........V...................D.S.....................
ACZ48518  A/quail/Italy/1117/1965(H10N8)                           ...RY.................V...................T......V.I.........
ABI84499  A/quail/Italy/1117/1965(H10N8)                           ...RY.................V...................T......V.I.........
AEA04419  A/quail/Italy/1117/1965(H10N8)                           ...RY.................V...................T......V.I.........
AGU01953  A/ruddy shelduck/Mongolia/1602/2010(H10N8)                ...N.....................................................
AGL58508  A/ruddy turnstone/New Jersey/1148668/2004(H10N8)          .S.NY..........V...................D.S.....................
AGG28633  A/shorebird/Delaware Bay/379/2008(H10N8)                  .N.NY....S......V...................D.S.....................
AGU01957  A/surf scoter/Mongolia/878V/2009(H10N8)                   ...N.....................................................I.........
AGU01970  A/velvet scoter/Mongolia/879V/2009(H10N8)                 ...N.....................................................I.........

Positions from 121 till 180
          Consensus sequence                                         ESGGIDKISTGFTYGSSINSAGTTRACMRNGGNSFYAELKWLVSKSKGQNFPQTTNTYRN
AHN97094  A/American green-winged teal/Ohio/13OS1869/2013(H10N8)     ........................K....................N.............
AHM24205  A/Eurasian coot/Germany/R411/2010(H10N8)                  ....VS.............P..V.K..L...QS......R.....TQ.............
AHK10761  A/Jiangxi/IPB13/2013(H10N8)                               .....N...................................................
AHK10763  A/Jiangxi/IPB13b/2013(H10N8)                              .....N...................................................
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FIGURE 23-3

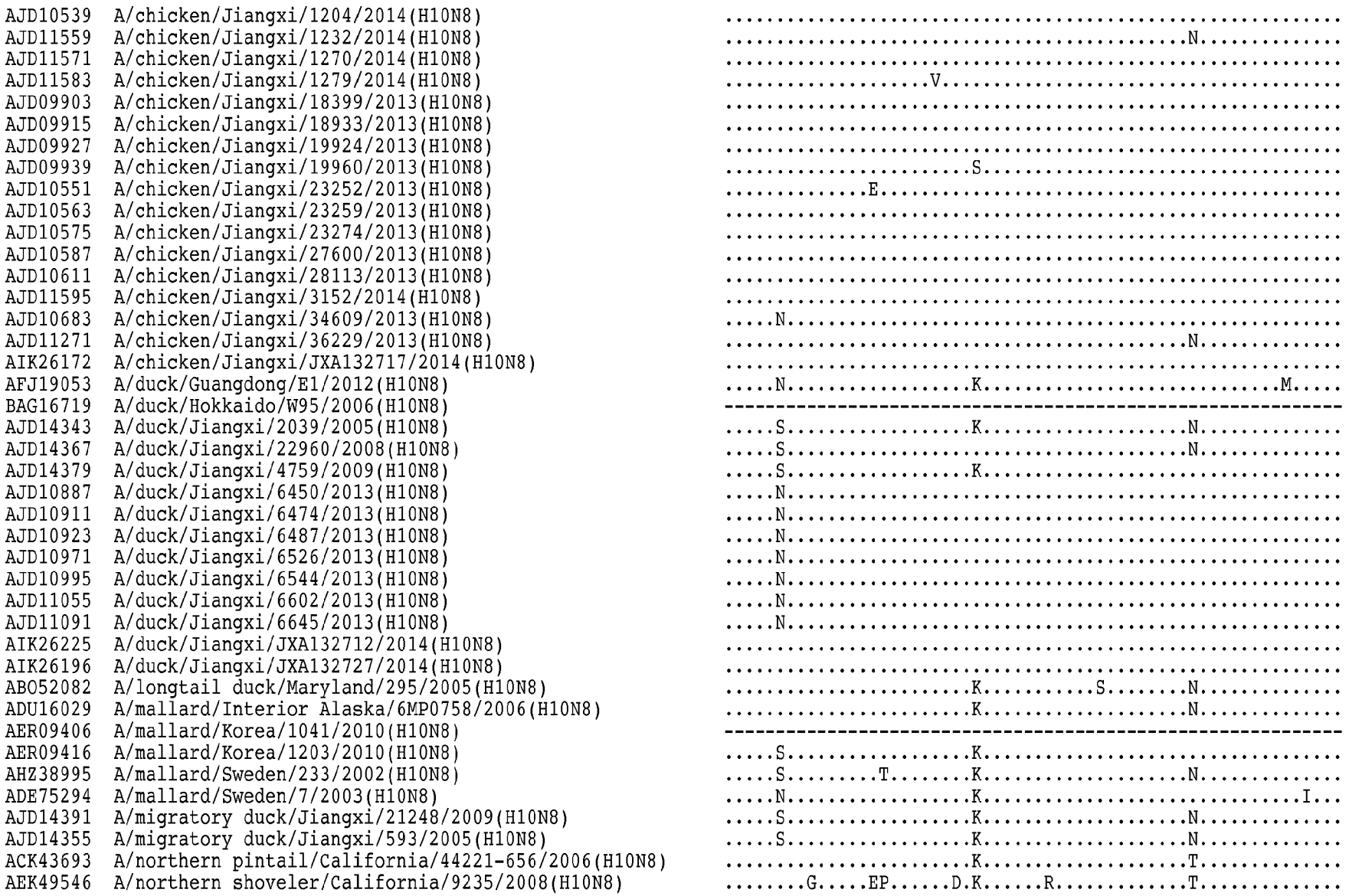

AJD10539 A/chicken/Jiangxi/1204/2014(H10N8)
AJD11559 A/chicken/Jiangxi/1232/2014(H10N8)
AJD11571 A/chicken/Jiangxi/1270/2014(H10N8)
AJD11583 A/chicken/Jiangxi/1279/2014(H10N8)
AJD09903 A/chicken/Jiangxi/18399/2013(H10N8)
AJD09915 A/chicken/Jiangxi/18933/2013(H10N8)
AJD09927 A/chicken/Jiangxi/19924/2013(H10N8)
AJD09939 A/chicken/Jiangxi/19960/2013(H10N8)
AJD10551 A/chicken/Jiangxi/23252/2013(H10N8)
AJD10563 A/chicken/Jiangxi/23259/2013(H10N8)
AJD10575 A/chicken/Jiangxi/23274/2013(H10N8)
AJD10587 A/chicken/Jiangxi/27600/2013(H10N8)
AJD10611 A/chicken/Jiangxi/28113/2013(H10N8)
AJD11595 A/chicken/Jiangxi/3152/2014(H10N8)
AJD10683 A/chicken/Jiangxi/34609/2013(H10N8)
AJD11271 A/chicken/Jiangxi/36229/2013(H10N8)
AIK26172 A/chicken/Jiangxi/JXA132717/2014(H10N8)
AFJ19053 A/duck/Guangdong/E1/2012(H10N8)
BAG16719 A/duck/Hokkaido/W95/2006(H10N8)
AJD14343 A/duck/Jiangxi/2039/2005(H10N8)
AJD14367 A/duck/Jiangxi/22960/2008(H10N8)
AJD14379 A/duck/Jiangxi/4759/2009(H10N8)
AJD10887 A/duck/Jiangxi/6450/2013(H10N8)
AJD10911 A/duck/Jiangxi/6474/2013(H10N8)
AJD10923 A/duck/Jiangxi/6487/2013(H10N8)
AJD10971 A/duck/Jiangxi/6526/2013(H10N8)
AJD10995 A/duck/Jiangxi/6544/2013(H10N8)
AJD11055 A/duck/Jiangxi/6602/2013(H10N8)
AJD11091 A/duck/Jiangxi/6645/2013(H10N8)
AIK26225 A/duck/Jiangxi/JXA132712/2014(H10N8)
AIK26196 A/duck/Jiangxi/JXA132727/2014(H10N8)
ABO52082 A/longtail duck/Maryland/295/2005(H10N8)
ADU16029 A/mallard/Interior Alaska/6MP0758/2006(H10N8)
AER09406 A/mallard/Korea/1041/2010(H10N8)
AER09416 A/mallard/Korea/1203/2010(H10N8)
AHZ38995 A/mallard/Sweden/233/2002(H10N8)
ADE75294 A/mallard/Sweden/7/2003(H10N8)
AJD14391 A/migratory duck/Jiangxi/21248/2009(H10N8)
AJD14355 A/migratory duck/Jiangxi/593/2005(H10N8)
ACK43693 A/northern pintail/California/44221-656/2006(H10N8)
AEK49546 A/northern shoveler/California/9235/2008(H10N8)

FIGURE 23-4

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ACZ48518  A/quail/Italy/1117/1965(H10N8)                          .....S.MR......P.........S..................GT.............
ABI84499  A/quail/Italy/1117/1965(H10N8)                          .....S.MR......P.........S..................GT.............
AEA04419  A/quail/Italy/1117/1965(H10N8)                          .....S.MR......P.........S..................GT.............
AGU01953  A/ruddy shelduck/Mongolia/1602/2010(H10N8)              .....S.TN................K.....................N.............
AGL58508  A/ruddy turnstone/New Jersey/1148668/2004(H10N8)       ..........................S...S..............N........A.....
AGG28633  A/shorebird/Delaware Bay/379/2008(H10N8)               ..............................................................
AGU01957  A/surf scoter/Mongolia/878V/2009(H10N8)                .....S..................K........G...........N.............
AGU01970  A/velvet scoter/Mongolia/879V/2009(H10N8)              .....S..................K........G...........N.............

Positions from 181 till 240
          Consensus sequence                                      TDTAEHLIMWGIHHPSSTQEKNDLYGTQSLSISVGSSTYQNNFVPVVGARPQVNGQSGRI
AHN97094  A/American green-winged teal/Ohio/13OS1869/2013(H10N8)  ..S.....I.....................................................
AHM24205  A/Eurasian coot/Germany/R411/2010(H10N8)                ........I.............K...A...Y..........S.T.................
AHK10761  A/Jiangxi/IPB13/2013(H10N8)                            ..........................................R...................
AHK10763  A/Jiangxi/IPB13b/2013(H10N8)                           ..........................................R.........G.........
AJD10539  A/chicken/Jiangxi/1204/2014(H10N8)                     ..........................................R...................
AJD11559  A/chicken/Jiangxi/1232/2014(H10N8)                     ..................I.......................R.S.................
AJD11571  A/chicken/Jiangxi/1270/2014(H10N8)                     ..................I.......................R...................
AJD11583  A/chicken/Jiangxi/1279/2014(H10N8)                     ..................I.......................R...................
AJD09903  A/chicken/Jiangxi/18399/2013(H10N8)                    ..........................................R...................
AJD09915  A/chicken/Jiangxi/18933/2013(H10N8)                    ..........................................R...................
AJD09927  A/chicken/Jiangxi/19924/2013(H10N8)                    ..........................................R...................
AJD09939  A/chicken/Jiangxi/19960/2013(H10N8)                    ..........................................R...................
AJD10551  A/chicken/Jiangxi/23252/2013(H10N8)                    ..........................................R...................
AJD10563  A/chicken/Jiangxi/23259/2013(H10N8)                    ..........................................R...................
AJD10575  A/chicken/Jiangxi/23274/2013(H10N8)                    ..........................................R...................
AJD10587  A/chicken/Jiangxi/27600/2013(H10N8)                    ..........................................R...................
AJD10611  A/chicken/Jiangxi/28113/2013(H10N8)                    ..........................................R...................
AJD11595  A/chicken/Jiangxi/3152/2014(H10N8)                     ..................I.......................R...................
AJD10683  A/chicken/Jiangxi/34609/2013(H10N8)                    ..........................................R...................
AJD11271  A/chicken/Jiangxi/36229/2013(H10N8)                    ..................I.......................R...................
AIK26172  A/chicken/Jiangxi/JXA132717/2014(H10N8)                ..........................................R...................
AFJ19053  A/duck/Guangdong/E1/2012(H10N8)                        ...V.............I.....................................................
BAG16719  A/duck/Hokkaido/W95/2006(H10N8)                        -------------------------------------------------------------
AJD14343  A/duck/Jiangxi/2039/2005(H10N8)                        ..............................................................
AJD14367  A/duck/Jiangxi/22960/2008(H10N8)                       ........I.....................................................
AJD14379  A/duck/Jiangxi/4759/2009(H10N8)                        ........I.....................................................
AJD10887  A/duck/Jiangxi/6450/2013(H10N8)                        ..............................................................
AJD10911  A/duck/Jiangxi/6474/2013(H10N8)                        ..............................................................
AJD10923  A/duck/Jiangxi/6487/2013(H10N8)                        ..............................................................
AJD10971  A/duck/Jiangxi/6526/2013(H10N8)                        ..............................................................
AJD10995  A/duck/Jiangxi/6544/2013(H10N8)                        ..............................................................
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FIGURE 23-5

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AJD11055  A/duck/Jiangxi/6602/2013(H10N8)                                  ..................................................
AJD11091  A/duck/Jiangxi/6645/2013(H10N8)                                  ..................................................
AIK26225  A/duck/Jiangxi/JXA132712/2014(H10N8)                             ..................................R...............
AIK26196  A/duck/Jiangxi/JXA132727/2014(H10N8)                             ..................................R...............
ABO52082  A/longtail duck/Maryland/295/2005(H10N8)                         ..SV....I..........................................
ADU16029  A/mallard/Interior Alaska/6MP0758/2006(H10N8)                    ..S.....I..........................................
AER09406  A/mallard/Korea/1041/2010(H10N8)                                 --.S...............................................
AER09416  A/mallard/Korea/1203/2010(H10N8)                                 ...S..........Y....................................
AHZ38995  A/mallard/Sweden/233/2002(H10N8)                                 ........I..........................................
ADE75294  A/mallard/Sweden/7/2003(H10N8)                                   .......V...........................................
AJD14391  A/migratory duck/Jiangxi/21248/2009(H10N8)                       ...............................I..................
AJD14355  A/migratory duck/Jiangxi/593/2005(H10N8)                         ..................................................
ACK43693  A/northern pintail/California/44221-656/2006(H10N8)              ..S.....I..........................................
AEK49546  A/northern shoveler/California/9235/2008(H10N8)                  ..S.....I.........K................................
ACZ48518  A/quail/Italy/1117/1965(H10N8)                                   ........I.....................................I.....
ABI84499  A/quail/Italy/1117/1965(H10N8)                                   ........I.....................................I.....
AEA04419  A/quail/Italy/1117/1965(H10N8)                                   ........I.....................................I.....
AGU01953  A/ruddy shelduck/Mongolia/1602/2010(H10N8)                       ..................................S...............
AGL58508  A/ruddy turnstone/New Jersey/1148668/2004(H10N8)                 ..S.....I..........................................
AGG28633  A/shorebird/Delaware Bay/379/2008(H10N8)                         ..S.....I..........................................
AGU01957  A/surf scoter/Mongolia/878V/2009(H10N8)                          ..............................................I.....
AGU01970  A/velvet scoter/Mongolia/879V/2009(H10N8)                        ..............................................I.....

Positions from 241 till 300
          Consensus sequence                                               DFHWTLVQPGDNITFSHNGGLIAPSRVSKLIGRGLGIQSDAPIDNNCESKCFWRGGSINT
AHN97094  A/American green-winged teal/Ohio/13OS1869/2013(H10N8)           .....M.....................K........G.SV..D.......K......
AHM24205  A/Eurasian coot/Germany/R411/2010(H10N8)                         E..........................T........E.SV...............
AHK10761  A/Jiangxi/IPB13/2013(H10N8)                                      ..................................................
AHK10763  A/Jiangxi/IPB13b/2013(H10N8)                                     ..................................................
AJD10539  A/chicken/Jiangxi/1204/2014(H10N8)                               ..................................................
AJD11559  A/chicken/Jiangxi/1232/2014(H10N8)                               ..................................................
AJD11571  A/chicken/Jiangxi/1270/2014(H10N8)                               ..................................................
AJD11583  A/chicken/Jiangxi/1279/2014(H10N8)                               ..................................................
AJD09903  A/chicken/Jiangxi/18399/2013(H10N8)                              ..................................................
AJD09915  A/chicken/Jiangxi/18933/2013(H10N8)                              ..................................................
AJD09927  A/chicken/Jiangxi/19924/2013(H10N8)                              ..................................................
AJD09939  A/chicken/Jiangxi/19960/2013(H10N8)                              ..................................................
AJD10551  A/chicken/Jiangxi/23252/2013(H10N8)                              ..................................................
AJD10563  A/chicken/Jiangxi/23259/2013(H10N8)                              ..................................................
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FIGURE 23-6

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AJD10575  A/chicken/Jiangxi/23274/2013(H10N8)                    ..........................................................
AJD10587  A/chicken/Jiangxi/27600/2013(H10N8)                    ..........................................................
AJD10611  A/chicken/Jiangxi/28113/2013(H10N8)                    ..........................................................
AJD11595  A/chicken/Jiangxi/3152/2014(H10N8)                     ..........................................................
AJD10683  A/chicken/Jiangxi/34609/2013(H10N8)                    ..........................................................
AJD11271  A/chicken/Jiangxi/36229/2013(H10N8)                    ..........................................................
AIK26172  A/chicken/Jiangxi/JXA132717/2014(H10N8)                ..........................................................
AFJ19053  A/duck/Guangdong/E1/2012(H10N8)                        ..........................V.......................V.....
BAG16719  A/duck/Hokkaido/W95/2006(H10N8)                        ----------------------------------------------------------
AJD14343  A/duck/Jiangxi/2039/2005(H10N8)                        ......I.....................V.....................V.....
AJD14367  A/duck/Jiangxi/22960/2008(H10N8)                       ..........................V.....T.....................
AJD14379  A/duck/Jiangxi/4759/2009(H10N8)                        ..........................V.....................V.....
AJD10887  A/duck/Jiangxi/6450/2013(H10N8)                        ..........................................................
AJD10911  A/duck/Jiangxi/6474/2013(H10N8)                        ..........................................................
AJD10923  A/duck/Jiangxi/6487/2013(H10N8)                        ..........................................................
AJD10971  A/duck/Jiangxi/6526/2013(H10N8)                        ..........................................Y...............
AJD10995  A/duck/Jiangxi/6544/2013(H10N8)                        ..........................................................
AJD11055  A/duck/Jiangxi/6602/2013(H10N8)                        ..........................................T...............
AJD11091  A/duck/Jiangxi/6645/2013(H10N8)                        ..........................................................
AIK26225  A/duck/Jiangxi/JXA132712/2014(H10N8)                   ..........................................................
AIK26196  A/duck/Jiangxi/JXA132727/2014(H10N8)                   ..........................................................
ABO52082  A/longtail duck/Maryland/295/2005(H10N8)               .....M.................N.....K........G.SV..D.......K......
ADU16029  A/mallard/Interior Alaska/6MP0758/2006(H10N8)          .....M.....................K........GVSV..D.......K......
AER09406  A/mallard/Korea/1041/2010(H10N8)                       ..........................................L...............
AER09416  A/mallard/Korea/1203/2010(H10N8)                       ..........................................S...............
AHZ38995  A/mallard/Sweden/233/2002(H10N8)                      ......L.....................................E.....G.......K......
ADE75294  A/mallard/Sweden/7/2003(H10N8)                        ..........................................D...............
AJD14391  A/migratory duck/Jiangxi/21248/2009(H10N8)            ..........................................................
AJD14355  A/migratory duck/Jiangxi/593/2005(H10N8)              ..........................V.....................V.....
ACK43693  A/northern pintail/California/44221-656/2006(H10N8)   .....M.....................K........G.SV..D.......K......
AEK49546  A/northern shoveler/California/9235/2008(H10N8)       .....M.....................K........E.S...D.......K......
ACZ48518  A/quail/Italy/1117/1965(H10N8)                       E......R.............D.......K.I....G.V..KD............I.
ABI84499  A/quail/Italy/1117/1965(H10N8)                       E......R.............D.......K.I....G.V..KD............I.
AEA04419  A/quail/Italy/1117/1965(H10N8)                       E......R.............D.......K.I....G.V..KD............I.
AGU01953  A/ruddy shelduck/Mongolia/1602/2010(H10N8)           ....................N.....................................
AGL58508  A/ruddy turnstone/New Jersey/1148668/2004(H10N8)     .....M.....................K........G.SV..D.......K......
AGG28633  A/shorebird/Delaware Bay/379/2008(H10N8)            .....M.....................K........G.SV..D.......K....K.
AGU01957  A/surf scoter/Mongolia/878V/2009(H10N8)             ...................M..N.....................................
AGU01970  A/velvet scoter/Mongolia/879V/2009(H10N8)           ...................M..N.....................................
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FIGURE 23-7

Positions from 301 till 360

	Consensus sequence	RLPFQNLSPRTVGQCPKYVNKRSLMLATGMRNVPELMQGRGLFGAIAGFIENGWEGMVDG
AHN97094	A/American green-winged teal/Ohio/13OS1869/2013(H10N8)	K.....L.....VV.....
AHM24205	A/Eurasian coot/Germany/R411/2010(H10N8)	K.....L.....V.....
AHK10761	A/Jiangxi/IPB13/2013(H10N8)	.....R.....I.....L.....
AHK10763	A/Jiangxi/IPB13b/2013(H10N8)	.....R.....I.....L.....
AJD10539	A/chicken/Jiangxi/1204/2014(H10N8)	K.....R.....I.....L.....
AJD11559	A/chicken/Jiangxi/1232/2014(H10N8)	.....R.....I.....L.....
AJD11571	A/chicken/Jiangxi/1270/2014(H10N8)	.....R.....I.....L.....
AJD11583	A/chicken/Jiangxi/1279/2014(H10N8)	.....R.....I.....L.....
AJD09903	A/chicken/Jiangxi/18399/2013(H10N8)	.....L.....
AJD09915	A/chicken/Jiangxi/18933/2013(H10N8)	.....I.....L.....
AJD09927	A/chicken/Jiangxi/19924/2013(H10N8)	.....L.....
AJD09939	A/chicken/Jiangxi/19960/2013(H10N8)	.....L.....
AJD10551	A/chicken/Jiangxi/23252/2013(H10N8)	.....L.....
AJD10563	A/chicken/Jiangxi/23259/2013(H10N8)	.....I.....L.....
AJD10575	A/chicken/Jiangxi/23274/2013(H10N8)	.....R.....I.....L.....
AJD10587	A/chicken/Jiangxi/27600/2013(H10N8)	.....R.....I.....L.....
AJD10611	A/chicken/Jiangxi/28113/2013(H10N8)	.....R.....I.....L.....
AJD11595	A/chicken/Jiangxi/3152/2014(H10N8)	.....R.....I.....L.....
AJD10683	A/chicken/Jiangxi/34609/2013(H10N8)	.....R.....I.....L.....
AJD11271	A/chicken/Jiangxi/36229/2013(H10N8)	.....R.....I.....L.....
AIK26172	A/chicken/Jiangxi/JXA132717/2014(H10N8)	K.....RK.....I.....L.....
AFJ19053	A/duck/Guangdong/E1/2012(H10N8)	K.....K.....IV.E.....
BAG16719	A/duck/Hokkaido/W95/2006(H10N8)	----------
AJD14343	A/duck/Jiangxi/2039/2005(H10N8)	.....K.....I.....
AJD14367	A/duck/Jiangxi/22960/2008(H10N8)	.....I.....
AJD14379	A/duck/Jiangxi/4759/2009(H10N8)	.....I.....
AJD10887	A/duck/Jiangxi/6450/2013(H10N8)	.....
AJD10911	A/duck/Jiangxi/6474/2013(H10N8)	.....
AJD10923	A/duck/Jiangxi/6487/2013(H10N8)	.....
AJD10971	A/duck/Jiangxi/6526/2013(H10N8)	.....
AJD10995	A/duck/Jiangxi/6544/2013(H10N8)	.....
AJD11055	A/duck/Jiangxi/6602/2013(H10N8)	.....
AJD11091	A/duck/Jiangxi/6645/2013(H10N8)	.....
AIK26225	A/duck/Jiangxi/JXA132712/2014(H10N8)	.....R.....I.....L.....
AIK26196	A/duck/Jiangxi/JXA132727/2014(H10N8)	K.....RK.....I.....L.....
ABO52082	A/longtail duck/Maryland/295/2005(H10N8)	K.....K..L.....VA.....
ADU16029	A/mallard/Interior Alaska/6MP0758/2006(H10N8)	K.....K..L.....VV.....
AER09406	A/mallard/Korea/1041/2010(H10N8)	.....K.....I.....
AER09416	A/mallard/Korea/1203/2010(H10N8)	.....K.....I.....

FIGURE 23-8

AHZ38995	A/mallard/Sweden/233/2002(H10N8)	K.....I.H.....
ADE75294	A/mallard/Sweden/7/2003(H10N8)	.....K.....I.....
AJD14391	A/migratory duck/Jiangxi/21248/2009(H10N8)	.....K.....I.....
AJD14355	A/migratory duck/Jiangxi/593/2005(H10N8)	.....K.....I.....
ACK43693	A/northern pintail/California/44221-656/2006(H10N8)	K.....K..L.....VV.....
AEK49546	A/northern shoveler/California/9235/2008(H10N8)	K.....L.....VV.....
ACZ48518	A/quail/Italy/1117/1965(H10N8)	E.....K....L.....----------
ABI84499	A/quail/Italy/1117/1965(H10N8)	E.....K....L.....VV.....
AEA04419	A/quail/Italy/1117/1965(H10N8)	E.....K....L.....VV.....K.....
AGU01953	A/ruddy shelduck/Mongolia/1602/2010(H10N8)	.....K.....I.....
AGL58508	A/ruddy turnstone/New Jersey/1148668/2004(H10N8)	K.....K..L.....VV.....
AGG28633	A/shorebird/Delaware Bay/379/2008(H10N8)	K.....K..L.....VV.....
AGU01957	A/surf scoter/Mongolia/878V/2009(H10N8)	.....K.....I.....
AGU01970	A/velvet scoter/Mongolia/879V/2009(H10N8)	.....K.....I.....

Positions from 361 till 420

	Consensus sequence	**WYGFRHQNAQGTGQAADYKSTQAAIDQITGKLNRLIEKTNTEFESIESEFSEIEHQIGNV**
AHN97094	A/American green-winged teal/Ohio/13OS1869/2013(H10N8)	.....
AHM24205	A/Eurasian coot/Germany/R411/2010(H10N8)	.....N.....
AHK10761	A/Jiangxi/IPB13/2013(H10N8)	.....V.....
AHK10763	A/Jiangxi/IPB13b/2013(H10N8)	.....V.....
AJD10539	A/chicken/Jiangxi/1204/2014(H10N8)	.....V.....
AJD11559	A/chicken/Jiangxi/1232/2014(H10N8)	.....V.....
AJD11571	A/chicken/Jiangxi/1270/2014(H10N8)	.....V.....
AJD11583	A/chicken/Jiangxi/1279/2014(H10N8)	.....V.....
AJD09903	A/chicken/Jiangxi/18399/2013(H10N8)	.....V.....
AJD09915	A/chicken/Jiangxi/18933/2013(H10N8)	.....V.....
AJD09927	A/chicken/Jiangxi/19924/2013(H10N8)	.....V.....
AJD09939	A/chicken/Jiangxi/19960/2013(H10N8)	.....V.....
AJD10551	A/chicken/Jiangxi/23252/2013(H10N8)	.....V.....
AJD10563	A/chicken/Jiangxi/23259/2013(H10N8)	.....V.....
AJD10575	A/chicken/Jiangxi/23274/2013(H10N8)	.....V.....
AJD10587	A/chicken/Jiangxi/27600/2013(H10N8)	.....V.....
AJD10611	A/chicken/Jiangxi/28113/2013(H10N8)	.....V.....
AJD11595	A/chicken/Jiangxi/3152/2014(H10N8)	.....V.....
AJD10683	A/chicken/Jiangxi/34609/2013(H10N8)	.....V.....
AJD11271	A/chicken/Jiangxi/36229/2013(H10N8)	.....V.....
AIK26172	A/chicken/Jiangxi/JXA132717/2014(H10N8)	.....V.....
AFJ19053	A/duck/Guangdong/E1/2012(H10N8)	.....T.....
BAG16719	A/duck/Hokkaido/W95/2006(H10N8)	----------.....
AJD14343	A/duck/Jiangxi/2039/2005(H10N8)	.....
AJD14367	A/duck/Jiangxi/22960/2008(H10N8)	.....

FIGURE 23-9

```
AJD14379  A/duck/Jiangxi/4759/2009(H10N8)                                  ..................................................
AJD10887  A/duck/Jiangxi/6450/2013(H10N8)                                  ..................................................
AJD10911  A/duck/Jiangxi/6474/2013(H10N8)                                  ..................................................
AJD10923  A/duck/Jiangxi/6487/2013(H10N8)                                  ..................................................
AJD10971  A/duck/Jiangxi/6526/2013(H10N8)                                  ..................................................
AJD10995  A/duck/Jiangxi/6544/2013(H10N8)                                  ..................................................
AJD11055  A/duck/Jiangxi/6602/2013(H10N8)                                  ..................................................
AJD11091  A/duck/Jiangxi/6645/2013(H10N8)                                  ..................................................
AIK26225  A/duck/Jiangxi/JXA132712/2014(H10N8)                             ..................................V.................
AIK26225  A/duck/Jiangxi/JXA132712/2014(H10N8)                             ..................................V.................
AIK26196  A/duck/Jiangxi/JXA132727/2014(H10N8)                             ..................................V.................
ABO52082  A/longtail duck/Maryland/295/2005(H10N8)                         ..................................................
ADU16029  A/mallard/Interior Alaska/6MP0758/2006(H10N8)                    ..................................................
AER09406  A/mallard/Korea/1041/2010(H10N8)                                 ....................T.............................
AER09416  A/mallard/Korea/1203/2010(H10N8)                                 ....................T.............................
AHZ38995  A/mallard/Sweden/233/2002(H10N8)                                 ..................................................I
ADE75294  A/mallard/Sweden/7/2003(H10N8)                                   ..................................................
AJD14391  A/migratory duck/Jiangxi/21248/2009(H10N8)                       ..................................................
AJD14355  A/migratory duck/Jiangxi/593/2005(H10N8)                         ..................................................
ACK43693  A/northern pintail/California/44221-656/2006(H10N8)              ..................................................
AEK49546  A/northern shoveler/California/9235/2008(H10N8)                  ......................V...........................
ACZ48518  A/quail/Italy/1117/1965(H10N8)                                   --------------------------------------------------
ABI84499  A/quail/Italy/1117/1965(H10N8)                                   ..........I.........T.............................
AEA04419  A/quail/Italy/1117/1965(H10N8)                                   ..........I.........T.............................
AGU01953  A/ruddy shelduck/Mongolia/1602/2010(H10N8)                       ..................................................D.....
AGL58508  A/ruddy turnstone/New Jersey/1148668/2004(H10N8)                 ..................................................
AGG28633  A/shorebird/Delaware Bay/379/2008(H10N8)                         ..................................................
AGU01957  A/surf scoter/Mongolia/878V/2009(H10N8)                          ..................................................
AGU01970  A/velvet scoter/Mongolia/879V/2009(H10N8)                        ..................................................

Positions from 421 till 480
          Consensus sequence                                               INWTKDSITDIWTYQAELLVAMENQHTIDMADSEMLNLYERVRKQLRQNAEEDGKGCFEI
AHN97094  A/American green-winged teal/Ohio/13OS1869/2013(H10N8)           ..........................V.......................
AHM24205  A/Eurasian coot/Germany/R411/2010(H10N8)                         ......................................D...........
AHK10761  A/Jiangxi/IPB13/2013(H10N8)                                      ..................................................
AHK10763  A/Jiangxi/IPB13b/2013(H10N8)                                     ..................................................
AJD10539  A/chicken/Jiangxi/1204/2014(H10N8)                               ..................................................
AJD11559  A/chicken/Jiangxi/1232/2014(H10N8)                               ....M.............................................
AJD11571  A/chicken/Jiangxi/1270/2014(H10N8)                               ....M.............................................
AJD11583  A/chicken/Jiangxi/1279/2014(H10N8)                               ....M.............................................
```

FIGURE 23-10

AJD09903	A/chicken/Jiangxi/18399/2013(H10N8)	.....
AJD09915	A/chicken/Jiangxi/18933/2013(H10N8)	.....
AJD09927	A/chicken/Jiangxi/19924/2013(H10N8)	.....
AJD09939	A/chicken/Jiangxi/19960/2013(H10N8)	.....I.....
AJD10551	A/chicken/Jiangxi/23252/2013(H10N8)	.....
AJD10563	A/chicken/Jiangxi/23259/2013(H10N8)	.....
AJD10575	A/chicken/Jiangxi/23274/2013(H10N8)	.....
AJD10587	A/chicken/Jiangxi/27600/2013(H10N8)	.....
AJD10611	A/chicken/Jiangxi/28113/2013(H10N8)	.....
AJD11595	A/chicken/Jiangxi/3152/2014(H10N8)	....M.....
AJD10683	A/chicken/Jiangxi/34609/2013(H10N8)	.....
AJD11271	A/chicken/Jiangxi/36229/2013(H10N8)	....M.....
AIK26172	A/chicken/Jiangxi/JXA132717/2014(H10N8)	.....
AFJ19053	A/duck/Guangdong/E1/2012(H10N8)	.....
BAG16719	A/duck/Hokkaido/W95/2006(H10N8)	.....
AJD14343	A/duck/Jiangxi/2039/2005(H10N8)	.....
AJD14367	A/duck/Jiangxi/22960/2008(H10N8)	.....
AJD14379	A/duck/Jiangxi/4759/2009(H10N8)	.....
AJD10887	A/duck/Jiangxi/6450/2013(H10N8)	.....
AJD10911	A/duck/Jiangxi/6474/2013(H10N8)	.....
AJD10923	A/duck/Jiangxi/6487/2013(H10N8)	.....
AJD10971	A/duck/Jiangxi/6526/2013(H10N8)	.....
AJD10995	A/duck/Jiangxi/6544/2013(H10N8)	.....
AJD11055	A/duck/Jiangxi/6602/2013(H10N8)	.....
AJD11091	A/duck/Jiangxi/6645/2013(H10N8)	.....
AIK26225	A/duck/Jiangxi/JXA132712/2014(H10N8)	....M.....
AIK26196	A/duck/Jiangxi/JXA132727/2014(H10N8)	.....
ABO52082	A/longtail duck/Maryland/295/2005(H10N8)	.....
ADU16029	A/mallard/Interior Alaska/6MP0758/2006(H10N8)	.....
AER09406	A/mallard/Korea/1041/2010(H10N8)	.....
AER09416	A/mallard/Korea/1203/2010(H10N8)	.....
AHZ38995	A/mallard/Sweden/233/2002(H10N8)	.....
ADE75294	A/mallard/Sweden/7/2003(H10N8)	.....
AJD14391	A/migratory duck/Jiangxi/21248/2009(H10N8)	.....
AJD14355	A/migratory duck/Jiangxi/593/2005(H10N8)	.....
ACK43693	A/northern pintail/California/44221-656/2006(H10N8)	.....
AEK49546	A/northern shoveler/California/9235/2008(H10N8)	.....
ACZ48518	A/quail/Italy/1117/1965(H10N8)	----------
ABI84499	A/quail/Italy/1117/1965(H10N8)	.....

FIGURE 23-11

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AEA04419  A/quail/Italy/1117/1965(H10N8)                                  ..........
AGU01953  A/ruddy shelduck/Mongolia/1602/2010(H10N8)                      ..........
AGL58508  A/ruddy turnstone/New Jersey/1148668/2004(H10N8)                ..........
AGG28633  A/shorebird/Delaware Bay/379/2008(H10N8)                        ..........
AGU01957  A/surf scoter/Mongolia/878V/2009(H10N8)                         ..........
AGU01970  A/velvet scoter/Mongolia/879V/2009(H10N8)                       ..........

Positions from 481 till 540
          Consensus sequence                                              YHACDDSCMESIRNNTYDHSQYREEALLNRLNINPVTLSSGYKDIILWFSFGASCFVLLA
AHN97094  A/American green-winged teal/Ohio/13OS1869/2013(H10N8)          ..K...N.....T.....K.....V.....
AHM24205  A/Eurasian coot/Germany/R411/2010(H10N8)                        ..T..N.....KF.....
AHK10761  A/Jiangxi/IPB13/2013(H10N8)                                     ..........
AHK10763  A/Jiangxi/IPB13b/2013(H10N8)                                    ..........
AJD10539  A/chicken/Jiangxi/1204/2014(H10N8)                              ..........
AJD11559  A/chicken/Jiangxi/1232/2014(H10N8)                              ..........
AJD11571  A/chicken/Jiangxi/1270/2014(H10N8)                              ..........
AJD11583  A/chicken/Jiangxi/1279/2014(H10N8)                              ..........
AJD09903  A/chicken/Jiangxi/18399/2013(H10N8)                             .....R.....
AJD09915  A/chicken/Jiangxi/18933/2013(H10N8)                             ..........
AJD09927  A/chicken/Jiangxi/19924/2013(H10N8)                             ..........
AJD09939  A/chicken/Jiangxi/19960/2013(H10N8)                             ..........
AJD10551  A/chicken/Jiangxi/23252/2013(H10N8)                             ..........
AJD10563  A/chicken/Jiangxi/23259/2013(H10N8)                             ..........
AJD10575  A/chicken/Jiangxi/23274/2013(H10N8)                             ..........
AJD10587  A/chicken/Jiangxi/27600/2013(H10N8)                             ..........
AJD10611  A/chicken/Jiangxi/28113/2013(H10N8)                             .....N.....
AJD11595  A/chicken/Jiangxi/3152/2014(H10N8)                              ..........
AJD10683  A/chicken/Jiangxi/34609/2013(H10N8)                             ..........
AJD11271  A/chicken/Jiangxi/36229/2013(H10N8)                             ..........
AIK26172  A/chicken/Jiangxi/JXA132717/2014(H10N8)                         ..........
AFJ19053  A/duck/Guangdong/E1/2012(H10N8)                                 ..........
BAG16719  A/duck/Hokkaido/W95/2006(H10N8)                                 .....H.....K.....
AJD14343  A/duck/Jiangxi/2039/2005(H10N8)                                 .....K.....
AJD14367  A/duck/Jiangxi/22960/2008(H10N8)                                .....K.....
AJD14379  A/duck/Jiangxi/4759/2009(H10N8)                                 .....K.....
AJD10887  A/duck/Jiangxi/6450/2013(H10N8)                                 ..........
AJD10911  A/duck/Jiangxi/6474/2013(H10N8)                                 ..........
AJD10923  A/duck/Jiangxi/6487/2013(H10N8)                                 ..........
AJD10971  A/duck/Jiangxi/6526/2013(H10N8)                                 ..........
AJD10995  A/duck/Jiangxi/6544/2013(H10N8)                                 ..........
AJD11055  A/duck/Jiangxi/6602/2013(H10N8)                                 ..........
AJD11091  A/duck/Jiangxi/6645/2013(H10N8)                                 ..........
```

FIGURE 23-12

AIK26225	A/duck/Jiangxi/JXA132712/2014(H10N8)	.....
AIK26196	A/duck/Jiangxi/JXA132727/2014(H10N8)	.....
ABO52082	A/longtail duck/Maryland/295/2005(H10N8)	..K...N.....T.....K.....V.....
ADU16029	A/mallard/Interior Alaska/6MP0758/2006(H10N8)	..K...N.....T.....K.....V.....
AER09406	A/mallard/Korea/1041/2010(H10N8)	.....D.H.....
AER09416	A/mallard/Korea/1203/2010(H10N8)	.....
AHZ38995	A/mallard/Sweden/233/2002(H10N8)	.....K.....LI...
ADE75294	A/mallard/Sweden/7/2003(H10N8)	.....K.....
AJD14391	A/migratory duck/Jiangxi/21248/2009(H10N8)	.....
AJD14355	A/migratory duck/Jiangxi/593/2005(H10N8)	.....K.....
ACK43693	A/northern pintail/California/44221-656/2006(H10N8)	..K...N.....A.....K.....V.....
AEK49546	A/northern shoveler/California/9235/2008(H10N8)	..K.....A.....K.....V.....
ACZ48518	A/quail/Italy/1117/1965(H10N8)	----------|
ABI84499	A/quail/Italy/1117/1965(H10N8)	..T..N.....K.....
AEA04419	A/quail/Italy/1117/1965(H10N8)	..T..N.....K.....
AGU01953	A/ruddy shelduck/Mongolia/1602/2010(H10N8)	.....
AGL58508	A/ruddy turnstone/New Jersey/1148668/2004(H10N8)	..K...N.....T.....K.....V.....
AGG28633	A/shorebird/Delaware Bay/379/2008(H10N8)	..K...N.....T.....S.K.....V.....
AGU01957	A/surf scoter/Mongolia/878V/2009(H10N8)	.....E.....
AGU01970	A/velvet scoter/Mongolia/879V/2009(H10N8)	.....E.....

Positions from 541 till 561

	Consensus sequence	**VVMGLVFFCLKNGNMRCTICI**
AHN97094	A/American green-winged teal/Ohio/13OS1869/2013(H10N8)	.I.....
AHM24205	A/Eurasian coot/Germany/R411/2010(H10N8)	..L.....Q..V..
AHK10761	A/Jiangxi/IPB13/2013(H10N8)	.....F.....
AHK10763	A/Jiangxi/IPB13b/2013(H10N8)	.....F.....
AJD10539	A/chicken/Jiangxi/1204/2014(H10N8)	.....F.....
AJD11559	A/chicken/Jiangxi/1232/2014(H10N8)	.....
AJD11571	A/chicken/Jiangxi/1270/2014(H10N8)	.....
AJD11583	A/chicken/Jiangxi/1279/2014(H10N8)	.....
AJD09903	A/chicken/Jiangxi/18399/2013(H10N8)	.....
AJD09915	A/chicken/Jiangxi/18933/2013(H10N8)	.....
AJD09927	A/chicken/Jiangxi/19924/2013(H10N8)	.....
AJD09939	A/chicken/Jiangxi/19960/2013(H10N8)	.....
AJD10551	A/chicken/Jiangxi/23252/2013(H10N8)	.....
AJD10563	A/chicken/Jiangxi/23259/2013(H10N8)	.....
AJD10575	A/chicken/Jiangxi/23274/2013(H10N8)	.....F.....
AJD10587	A/chicken/Jiangxi/27600/2013(H10N8)	.....F.....
AJD10611	A/chicken/Jiangxi/28113/2013(H10N8)	.....F.....
AJD11595	A/chicken/Jiangxi/3152/2014(H10N8)	.....
AJD10683	A/chicken/Jiangxi/34609/2013(H10N8)	.....F.....

FIGURE 23-13

AJD11271	A/chicken/Jiangxi/36229/2013(H10N8)	.....
AIK26172	A/chicken/Jiangxi/JXA13272717/2014(H10N8)	.....F.....-
AFJ19053	A/duck/Guangdong/E1/2012(H10N8)	I.....
BAG16719	A/duck/Hokkaido/W95/2006(H10N8)	.....
AJD14343	A/duck/Jiangxi/2039/2005(H10N8)	.....
AJD14367	A/duck/Jiangxi/22960/2008(H10N8)	.....
AJD14379	A/duck/Jiangxi/4759/2009(H10N8)	.....
AJD10887	A/duck/Jiangxi/6450/2013(H10N8)	.....
AJD10911	A/duck/Jiangxi/6474/2013(H10N8)	.....
AJD10923	A/duck/Jiangxi/6487/2013(H10N8)	.....
AJD10971	A/duck/Jiangxi/6526/2013(H10N8)	.....
AJD10995	A/duck/Jiangxi/6544/2013(H10N8)	.....
AJD11055	A/duck/Jiangxi/6602/2013(H10N8)	.....
AJD11091	A/duck/Jiangxi/6645/2013(H10N8)	.....
AIK26225	A/duck/Jiangxi/JXA132712/2014(H10N8)	.....
AIK26196	A/duck/Jiangxi/JXA132727/2014(H10N8)	.....F.....---
ABO52082	A/longtail duck/Maryland/295/2005(H10N8)	.I.....
ADU16029	A/mallard/Interior Alaska/6MP0758/2006(H10N8)	.I.....
AER09406	A/mallard/Korea/1041/2010(H10N8)	.....Q.....
AER09416	A/mallard/Korea/1203/2010(H10N8)	.....
AHZ38995	A/mallard/Sweden/233/2002(H10N8)	.....
ADE75294	A/mallard/Sweden/7/2003(H10N8)	.....
AJD14391	A/migratory duck/Jiangxi/21248/2009(H10N8)	.....
AJD14355	A/migratory duck/Jiangxi/593/2005(H10N8)	.....
ACK43693	A/northern pintail/California/44221-656/2006(H10N8)	.I.....
AEK49546	A/northern shoveler/California/9235/2008(H10N8)	.I.....
ACZ48518	A/quail/Italy/1117/1965(H10N8)	----------
ABI84499	A/quail/Italy/1117/1965(H10N8)	.I...G.....
AEA04419	A/quail/Italy/1117/1965(H10N8)	.I...G.....
AGU01953	A/ruddy shelduck/Mongolia/1602/2010(H10N8)	.....
AGL58508	A/ruddy turnstone/New Jersey/1148668/2004(H10N8)	.I.....
AGG28633	A/shorebird/Delaware Bay/379/2008(H10N8)	.I.....
AGU01957	A/surf scoter/Mongolia/878V/2009(H10N8)	.....L.
AGU01970	A/velvet scoter/Mongolia/879V/2009(H10N8)	.A.....

FIGURE 23-14

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