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(54) Titre : COMBINAISONS D'ANTAGONISTES DE PD-1 ET D'AGONISTES DE STING DINUCLEOTIDIQUES
CYCLIQUES POUR LE TRAITEMENT DU CANCER
(54) Title: COMBINATIONS OF PD-1 ANTAGONISTS AND CYCLIC DINUCLEOTIDE STING AGONISTS FOR CANCER
TREATMENT

SEQ ID NO 1: Pembrolizumab, Variable Light chain, amino acid

Glu Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser
Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Lys Gly
Val Ser Thr Ser Gly Tyr Ser Tyr Leu His Trp Tyr Gln Gln
Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr Leu Ala Ser
Tyr Leu Glu Ser Gly Val Pro Ala Arg Phe Ser Gly Ser Gly
Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro
Glu Asp Phe Ala Val Tyr Tyr Cys Gln His Ser Arg Asp Leu
Pro Leu Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys

SEQ ID NO 2: Pembrolizumab, Variable Heavy chain, amino acid

Gln Val Gln Leu Val Gln Ser Gly Val Glu Val Lys Lys Pro
Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr
Phe Thr Asn Tyr Tyr Met Tyr Trp Val Arg Gln Ala Pro Gly
Gln Gly Leu Glu Trp Met Gly Gly Ile Asn Pro Ser Asn Gly
Gly Thr Asn Phe Asn Glu Lys Phe Lys Asn Arg Val Thr Leu
Thr Thr Asp Ser Ser Thr Thr Thr Ala Tyr Met Glu Leu Lys
Ser Leu Gln Phe Asp Asp Thr Ala Val Tyr Tyr Cys Ala Arg
Arg Asp Tyr Arg Phe Asp Met Gly Phe Asp Tyr Trp Gly Gln
Gly Thr Thr Val Thr Val Ser Ser

FIG.1

(57) Abrégé/Abstract:

Therapeutic combinations that comprise at least one antagonist of the Programmed Death 1 receptor (PD-1) and at least one cyclic dinucleotide compound that activates the Stimulator of Interferon Genes (STING) pathway are disclosed herein. Also disclosed is the use of such therapeutic combinations for the treatment of cancers.

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(54) Title: COMBINATIONS OF PD-1 ANTAGONISTS AND CYCLIC DINUCLEOTIDE STING AGONISTS FOR CANCER
TREATMENT

SEQ ID NO 1: Pembrolizumab, Variable Light chain, amino acid
Glu Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser
Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Lys Gly
Val Ser Thr Ser Gly Tyr Ser Tyr Leu His Trp Tyr Gln Gln
Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr Leu Ala Ser
Tyr Leu Glu Ser Gly Val Pro Ala Arg Phe Ser Gly Ser Gly
Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro
Glu Asp Phe Ala Val Tyr Tyr Cys Gln His Ser Arg Asp Leu
Pro Leu Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys

SEQ ID NO 2: Pembrolizumab, Variable Heavy chain, amino acid
Gln Val Gln Leu Val Gln Ser Gly Val Glu Val Lys Lys Pro
Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr
Phe Thr Asn Tyr Tyr Met Tyr Trp Val Arg Gln Ala Pro Gly
Gln Gly Leu Glu Trp Met Gly Gly Ile Asn Pro Ser Asn Gly
Gly Thr Asn Phe Asn Glu Lys Phe Lys Asn Arg Val Thr Leu
Thr Thr Asp Ser Ser Thr Thr Thr Ala Tyr Met Glu Leu Lys
Ser Leu Gln Phe Asp Asp Thr Ala Val Tyr Tyr Cys Ala Arg
Arg Asp Tyr Arg Phe Asp Met Gly Phe Asp Tyr Trp Gly Gln
Gly Thr Thr Val Thr Val Ser Ser

FIG.1

(57) Abstract: Therapeutic combinations that comprise at least one antagonist of the Programmed Death 1 receptor (PD-1) and at least one cyclic dinucleotide compound that activates the Stimulator of Interferon Genes (STING) pathway are disclosed herein. Also disclosed is the use of such therapeutic combinations for the treatment of cancers.

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TITLE OF THE APPLICATION

COMBINATIONS OF PD-1 ANTAGONISTS AND CYCLIC DINUCLEOTIDE STING
AGONISTS FOR CANCER TREATMENT

5 FIELD OF THE INVENTION

[0001] The present disclosure relates to combinations of therapeutic compounds that are
useful to treat cancer. In particular, this disclosure relates to combination therapies comprising at
least one antagonist of a Programmed Death 1 protein (PD-1) and at least one cyclic dinucleotide
compound (CDN) that is useful as a STING (Stimulator of Interferon Genes) agonist and
10 activates the STING pathway.

BACKGROUND OF THE INVENTION

[0002] The cytotoxic T-lymphocyte-associated antigen 4 (CTLA-4) and PD-1 pathways
are important negative regulators of immune response. Activated T-cells up-regulate CTLA-4,
15 which binds on antigen-presenting cells and inhibits T-cell stimulation, IL-2 gene expression,
and T-cell proliferation. These anti-tumor effects have been observed in mouse models of colon
carcinoma, metastatic prostate cancer, and metastatic melanoma. PD-1 binds to active T-cells
and suppresses T-cell activation. PD-1 antagonists have demonstrated anti-tumor effects as well.
PD-1 is moderately expressed on naïve T-, B- and natural killer (NK) T-cells and is upregulated
20 by T/B cell receptor signaling on lymphocytes, monocytes, and myeloid cells.

[0003] Two known ligands for PD-1, PD-L1 (B7-H1) and PD-L2 (B7-DC), are expressed
in human cancers that arise in various tissues. In large sample sets of, for example, ovarian,
renal, colorectal, pancreatic, and liver cancers, and of melanoma, it was shown that PD-L1
expression correlated with poor prognosis and reduced overall patient survival irrespective of
25 subsequent treatment. Similarly, PD-1 expression on tumor infiltrating lymphocytes was found
to mark dysfunctional T-cells in breast cancer and melanoma and to correlate with poor
prognosis in renal cancer patients. Thus, it has been proposed that PD-L1 expressing tumor cells
interact with PD-1 expressing T-cells to attenuate T-cell activation and evasion of immune
surveillance, thereby contributing to an impaired immune response against the tumor.

30 [0004] Several monoclonal antibodies that inhibit the interaction between PD-1 and one
or both of its ligands PD-L1 and PD-L2 are in clinical development for treating cancer. It has
been proposed that the efficacy of such antibodies might be enhanced if administered in

combination with other approved or experimental cancer therapies, e.g., radiation, surgery, chemotherapeutic agents, targeted therapies, agents that inhibit other signaling pathways that are dysregulated in tumors, and other immune enhancing agents. *See Morrissey et al.*, Clinical and Translational Science 9(2): 89-104 (2016).

5 [0005] Another potential immune therapy for cancers and for other cell-proliferation disorders is related to the immune system response to certain danger signals associated with cellular or tissue damage. The innate immune system has no antigen specificity but does respond to a variety of effector mechanisms, such as the damage-associated molecular patterns (DAMPs) or pathogen-associated molecular patterns (PAMPs), such as those associated with opsonization,
10 phagocytosis, activation of the complement system, and production of soluble bioactive molecules such as cytokines or chemokines. These are all mechanisms by which the innate immune system mediates its response. In this way, the innate immune system is able to provide broad protection against a wide range of threats to the host.

[0006] Free cytosolic DNA and RNA are among these PAMPs and DAMPs. It has
15 recently been demonstrated that the main sensor for cytosolic DNA is cGAS (cyclic GMP-AMP synthase). Upon recognition of cytosolic DNA, cGAS catalyzes the generation of the cyclic-dinucleotide 2'-3' cGAMP, an atypical second messenger that strongly binds to the ER-transmembrane adaptor protein STING. A conformational change is undergone by cGAMP-bound STING, which translocates to a perinuclear compartment and induces the activation of
20 critical transcription factors IRF-3 and NF- κ B. This leads to a strong induction of type I interferons and production of pro-inflammatory cytokines such as IL-6, TNF- α and IFN- γ .

[0007] The importance of type I interferons and pro-inflammatory cytokines on various cells of the immune system has been very well established. In particular, these molecules strongly potentiate T-cell activation by enhancing the ability of dendritic cells and macrophages
25 to uptake, process, present and cross-present antigens to T-cells. The T-cell stimulatory capacity of these antigen-presenting cells is augmented by the up-regulation of critical co-stimulatory molecules, such as CD80 or CD86. Finally, type I interferons can rapidly engage their cognate receptors and trigger the activation of interferon-responsive genes that can significantly contribute to adaptive immune cell activation.

30 [0008] From a therapeutic perspective, interferons, and compounds that can induce interferon production, have potential use in the treatment of human cancers. Such molecules are potentially useful as anti-cancer agents with multiple pathways of activity. Interferons can

inhibit human tumor cell-proliferation directly and may be synergistic with various approved chemotherapeutic agents. Type I interferons can significantly enhance anti-tumor immune responses by inducing activation of both the adaptive and innate immune cells. Finally, tumor invasiveness may be inhibited by interferons by modulating enzyme expression related to tissue remodeling.

[0009] In view of the potential of type I interferons and type I interferon-inducing compounds as anti-viral and anti-cancer agents, there remains a need for new agents that can induce potent type I interferon production. With the growing body of data demonstrating that the cGAS-STING cytosolic DNA sensory pathway has a significant capacity to induce type I interferons, cyclic dinucleotide STING activating agents are rapidly taking an important place in today's anti-tumor therapy landscape.

SUMMARY OF THE INVENTION

[0010] Embodiments of the disclosure include combination therapies, or therapeutic combinations, comprising at least one PD-1 antagonist and at least one cyclic dinucleotide STING agonist.

[0011] Another embodiment includes a method of treating a cell-proliferation disorder in a subject in need thereof, comprising administering a combination therapy comprising at least one PD-1 antagonist and at least one cyclic dinucleotide STING agonist.

[0012] Other embodiments, aspects and features of the present invention are either further described in or will be apparent from the ensuing description, examples, and appended claims.

BRIEF DESCRIPTION OF THE DRAWINGS

[0013] FIGURE 1 shows the amino acid sequences of the light chain and heavy chain variable regions for pembrolizumab that may be used in the combinations disclosed herein.

[0014] FIGURE 2 shows the amino acid sequence of the light chain for pembrolizumab.

[0015] FIGURE 3 shows the amino acid sequence of the heavy chain for pembrolizumab.

[0016] FIGURE 4 shows the amino acid sequences of the CDRs 1, 2, and 3 of the light chain variable region (CDRL1, CDRL2, and CDRL3) and of the CDRs 1, 2, and 3 of the heavy chain variable region (CDRH1, CDRH2, and CDRH3) for pembrolizumab.

[0017] FIGURE 5 shows the amino acid sequences of the light chain and heavy chain variable regions for nivolumab that may be used in the combinations disclosed herein.

[0018] FIGURE 6 shows the amino acid sequence of the light chain for nivolumab.

[0019] FIGURE 7 shows the amino acid sequence of the heavy chain for nivolumab.

5 [0020] FIGURE 8 shows the amino acid sequences of the CDRs 1, 2, and 3 of the light chain variable region (CDRL1, CDRL2, and CDRL3) and of the CDRs 1, 2, and 3 of the heavy chain variable region (CDRH1, CDRH2, and CDRH3) nivolumab.

[0021] FIGURE 9 shows the amino acid sequence for the human PD-L1 molecule (amino acids 19-290).

10

DETAILED DESCRIPTION OF THE INVENTION

ABBREVIATIONS

μg, ug	Microgram
Anti-PD-1	Antagonist of a Programmed Death 1 protein
BID	One dose twice daily
C57B1/6	Common inbred strain of laboratory mouse, also “C57 black 6”, “C57”, “black 6”, or “B6”
CDR	Complementary determining region
CR	Complete regression
Ctrl	Control
DFS	Disease free survival
DLT	Dose limiting toxicity
FFPE	Formalin-fixed, paraffin-embedded
FR	Framework region
IgG	Immunoglobulin G
IgG1	Immunoglobulin G subclass 1
IHC	Immunohistochemistry or immunohistochemical
IP	Intraperitoneal
IT	Intratumoral
kg	Kilogram
mAb	Monoclonal antibody
MC38	Murine Carcinoma-38 Mouse colon adenocarcinoma cell line

mg	Milligram
mIgG1	Murine immunoglobulin G subclass 1, Isotype control mAb for anti-PD-1 antibody muDX400
mL	Milliliter
mm	Millimeter
mm ³	Cubic millimeter, 0.001 mL
MPK	Milligram per kilogram
MTD	Maximum tolerated dose
n	Number of subjects in a treatment group
NCI	National Cancer Institute
OR	Overall response
OS	Overall survival
PBS	Phosphate-buffered saline, vehicle control for cyclic dinucleotide STING agonists
PD-1	Programmed cell death protein 1
PFS	Progression free survival
PR	Partial response
<i>p</i> -values	Calculated probability
QD	One dose per day
RECIST	Response Evaluation Criteria in Solid Tumors
SD	Stable disease
SEM	Standard error of the mean
TGI	Tumor growth inhibition
T/C	Median tumor volume of the treated animal/Median tumor volume of the control animal

[0022] Additional abbreviations may be defined throughout this disclosure.

DEFINITIONS

[0023] Certain technical and scientific terms are specifically defined below. Unless
5 specifically defined elsewhere in this document, all other technical and scientific terms used

herein have the meaning commonly understood by one of ordinary skill in the art to which this disclosure relates.

[0024] “About” when used to modify a numerically defined parameter (e.g., the dose of a PD-1 antagonist or CDN STING agonist, or the length of treatment time with a combination therapy described herein) means that the parameter may vary by as much as 10% below or above the stated numerical value for that parameter; where appropriate, the stated parameter may be rounded to the nearest whole number. For example, a dose of about 5mg/kg may vary between 4.5mg/kg and 5.5mg/kg.

[0025] As used herein, including the appended claims, the singular forms of words such as “a,” “an,” and “the,” include their corresponding plural references unless the context clearly dictates otherwise.

[0026] The terms “administration of” and or “administering” a compound should be understood to include providing a compound described herein, or a pharmaceutically acceptable salt thereof, and compositions of the foregoing to a subject.

[0027] As used herein, the term “antibody” refers to any form of immunoglobulin molecule that exhibits the desired biological or binding activity. Thus, it is used in the broadest sense and specifically covers, but is not limited to, monoclonal antibodies (including full length monoclonal antibodies), polyclonal antibodies, multispecific antibodies (e.g., bispecific antibodies), humanized, fully human antibodies, chimeric antibodies, and camelized single domain antibodies. “Parental antibodies” are antibodies obtained by exposure of an immune system to an antigen prior to modification of the antibodies for an intended use, such as humanization of an antibody for use as a human therapeutic. As used herein, the term “antibody” encompasses not only intact polyclonal or monoclonal antibodies, but also, unless otherwise specified, any antigen binding portion thereof that competes with the intact antibody for specific binding, fusion proteins comprising an antigen binding portion, and any other modified configuration of the immunoglobulin molecule that comprises an antigen recognition site.

[0028] As used herein, unless otherwise indicated, “antibody fragment” or “antigen binding fragment” refers to a fragment of an antibody that retains the ability to bind specifically to the antigen, e.g. fragments that retain one or more CDR regions. An antibody that “specifically binds to” PD-1 or PD-L1 is an antibody that exhibits preferential binding to PD-1 or PD-L1 (as appropriate) as compared to other proteins, but this specificity does not require

absolute binding specificity. An antibody is considered “specific” for its intended target if its binding is determinative of the presence of the target protein in a sample, e.g. without producing undesired results such as false positives. Antibodies, or binding fragments thereof, will bind to the target protein with an affinity that is at least two fold greater, preferably at least ten times greater, more preferably at least 20-times greater, and most preferably at least 100-times greater than the affinity with non-target proteins.

[0029] Antigen binding portions include, for example, Fab, Fab', F(ab')₂, Fd, Fv, domain antibodies (dAbs, e.g., shark and camelid antibodies), fragments including complementarity determining regions (CDRs), single chain variable fragment antibodies (scFv), maxibodies, minibodies, intrabodies, diabodies, triabodies, tetrabodies, v-NAR, and bis-scFv, and polypeptides that contain at least a portion of an immunoglobulin that is sufficient to confer specific antigen binding to the PD-1 or PD-L1. An antibody includes an antibody of any class, such as IgG, IgA, or IgM (or sub-class thereof), and the antibody need not be of any particular class. Depending on the antibody amino acid sequence of the constant region of its heavy chains, immunoglobulins can be assigned to different classes. There are five major classes of immunoglobulins: IgA, IgD, IgE, IgG, and IgM, and several of these may be further divided into subclasses (isotypes), e.g., IgG1, IgG2, IgG3, IgG4, IgA1, and IgA2. The heavy-chain constant regions that correspond to the different classes of immunoglobulins are called alpha, delta, epsilon, gamma, and mu, respectively. The subunit structures and three-dimensional configurations of different classes of immunoglobulins are well known.

[0030] As used herein, the terms “at least one” item or “one or more” item each include a single item selected from the list as well as mixtures of two or more items selected from the list.

[0031] As used herein, the term “immune response” relates to any one or more of the following: specific immune response, non-specific immune response, both specific and non-specific response, innate response, primary immune response, adaptive immunity, secondary immune response, memory immune response, immune cell activation, immune cell-proliferation, immune cell differentiation, and cytokine expression.

[0032] The term “pharmaceutically acceptable carrier” refers to any inactive substance that is suitable for use in a formulation for the delivery of a therapeutic agent. A carrier may be an antiadherent, binder, coating, disintegrant, filler or diluent, preservative (such as antioxidant, antibacterial, or antifungal agent), sweetener, absorption delaying agent, wetting agent, emulsifying agent, buffer, and the like. Examples of suitable pharmaceutically acceptable

carriers include water, ethanol, polyols (such as glycerol, propylene glycol, polyethylene glycol, and the like), dextrose, vegetable oils (such as olive oil), saline, buffer, buffered saline, and isotonic agents such as sugars, polyalcohols, sorbitol, and sodium chloride.

[0033] The term “subject” (alternatively “patient”) as used herein refers to a mammal that has been the object of treatment, observation, or experiment. The mammal may be male or female. The mammal may be one or more selected from the group consisting of humans, bovine (e.g., cows), porcine (e.g., pigs), ovine (e.g., sheep), capra (e.g., goats), equine (e.g., horses), canine (e.g., domestic dogs), feline (e.g., house cats), Lagomorpha (rabbits), rodents (e.g., rats or mice), *Procyon lotor* (e.g., raccoons). In particular embodiments, the subject is human.

[0034] The term “subject in need thereof” as used herein refers to a subject diagnosed with, or suspected of having a diagnosis of a cell-proliferation disorder, such as a cancer, as defined herein.

[0035] As used herein, the terms “treatment” and “treating” refer to all processes in which there may be a slowing, interrupting, arresting, controlling, or stopping of the progression of a disease or disorder described herein. The terms do not necessarily indicate a total elimination of all disease or disorder symptoms.

[0036] “Variable regions” or “V region” or “V chain” as used herein means the segment of IgG chains which is variable in sequence between different antibodies. A “variable region” of an antibody refers to the variable region of the antibody light chain or the variable region of the antibody heavy chain, either alone or in combination. Typically, the variable regions of both the heavy and light chains comprise three hypervariable regions, also called complementarity determining regions (CDRs), which are located within relatively conserved framework regions (FR). The CDRs are usually aligned by the framework regions, enabling binding to a specific epitope. In general, from N-terminal to C-terminal, both light and heavy chains variable domains comprise FR1, CDR1, FR2, CDR2, FR3, CDR3, and FR4. The assignment of amino acids to each domain is, generally, in accordance with the definitions of Sequences of Proteins of Immunological Interest, Kabat, *et al.*; National Institutes of Health, Bethesda, Md.; 5th ed.; NIH Publ. No. 91-3242 (1991); Kabat (1978) *Adv. Prot. Chem.* 32:1-75; Kabat, *et al.*, (1977) *J. Biol. Chem.* 252:6609-6616; Chothia, *et al.*, (1987) *J Mol. Biol.* 196:901-917 or Chothia, *et al.*, (1989) *Nature* 342:878-883.

[0037] “Chimeric antibody” refers to an antibody in which a portion of the heavy and/or light chain contains sequences derived from a particular species (e.g., human) or belonging to a

particular antibody class or subclass, while the remainder of the chain(s) is derived from another species (e.g., mouse) or belonging to another antibody class or subclass, as well as fragments of such antibodies, so long as they exhibit the desired biological activity.

[0038] “Human antibody” refers to an antibody that comprises human immunoglobulin

5 protein sequences or derivatives thereof. A human antibody may contain murine carbohydrate chains if produced in a mouse, in a mouse cell, or in a hybridoma derived from a mouse cell.

Similarly, “mouse antibody” or “rat antibody” refer to an antibody that comprises only mouse or rat immunoglobulin sequences or derivatives thereof, respectively.

[0039] “Humanized antibody” refers to forms of antibodies that contain sequences from

10 non-human (e.g., murine) antibodies as well as human antibodies. Such antibodies contain minimal sequence derived from non-human immunoglobulin. In general, the humanized antibody will comprise substantially all of at least one, and typically two, variable domains, in which all or substantially all of the hypervariable loops correspond to those of a non-human immunoglobulin and all or substantially all of the FR regions are those of a human

15 immunoglobulin sequence. The humanized antibody optionally also will comprise at least a portion of an immunoglobulin constant region (Fc), typically that of a human immunoglobulin.

The prefix “hum”, “hu” or “h” may be added to antibody clone designations when necessary to distinguish humanized antibodies from parental rodent antibodies. The humanized forms of rodent antibodies will generally comprise the same CDR sequences of the parental rodent

20 antibodies, although certain amino acid substitutions may be included to increase affinity, increase stability of the humanized antibody, or for other reasons.

[0040] “Biotherapeutic agent” means a biological molecule, such as an antibody or fusion protein, that blocks ligand/receptor signaling in any biological pathway that supports tumor maintenance and/or growth or suppresses the anti-tumor immune response.

25 [0041] “Chemotherapeutic agent” refers to a chemical or biological substance that can cause death of cancer cells, or interfere with growth, division, repair, and/or function of cancer cells. Examples of chemotherapeutic agents include those that are disclosed in WO2006/129163, and US20060153808, the disclosures of which are incorporated herein by reference. Classes of chemotherapeutic agents include, but are not limited to: hypomethylating agents, alkylating

30 agents, antimetabolites, spindle poison, plant alkaloids, cytotoxic/antitumor antibiotics, topoisomerase inhibitors, photosensitizers, hormonal therapies such as anti-estrogens and selective estrogen receptor modulators (SERMs), anti-progesterones, estrogen receptor down-regulators

(ERDs), estrogen receptor antagonists, leutinizing hormone-releasing hormone agonists, anti-androgens, aromatase inhibitors, and targeted therapies such as kinase inhibitors, EGFR inhibitors, VEGF inhibitors, and anti-sense oligonucleotides that inhibit expression of genes implicated in abnormal cell-proliferation or tumor growth. Chemotherapeutic agents useful in the treatment methods of the present disclosure include cytostatic and/or cytotoxic agents.

[0042] The therapeutic agents and compositions provided by the present disclosure can be administered via any suitable enteral route or parenteral route of administration. The term “enteral route” of administration refers to the administration via any part of the gastrointestinal tract. Examples of enteral routes include oral, mucosal, buccal, and rectal route, or intragastric route. “Parenteral route” of administration refers to a route of administration other than enteral route. Examples of parenteral routes of administration include intravenous, intramuscular, intradermal, intraperitoneal, intratumor, intravesical, intraarterial, intrathecal, intracapsular, intraorbital, intracardiac, transtracheal, intraarticular, subcapsular, subarachnoid, intraspinal, epidural and intrasternal, subcutaneous, or topical administration. The therapeutic agents and compositions of the disclosure can be administered using any suitable method, such as by oral ingestion, nasogastric tube, gastrostomy tube, injection, infusion, implantable infusion pump, and osmotic pump. The suitable route and method of administration may vary depending on a number of factors such as the specific antibody being used, the rate of absorption desired, specific formulation or dosage form used, type or severity of the disorder being treated, the specific site of action, and conditions of the patient, and can be readily selected by a person skilled in the art.

[0043] The term “simultaneous administration” as used herein in relation to the administration of medicaments refers to the administration of medicaments such that the individual medicaments are present within a subject at the same time. In addition to the concomitant administration of medicaments (via the same or alternative routes), simultaneous administration may include the administration of the medicaments (via the same or an alternative route) at different times.

[0044] “Chothia” as used herein means an antibody numbering system described in Al-Lazikani *et al.*, *JMB* **273**:927-948 (1997).

[0045] “Conservatively modified variants” or “conservative substitution” refers to substitutions of amino acids in a protein with other amino acids having similar characteristics (e.g., charge, side-chain size, hydrophobicity/hydrophilicity, backbone conformation and

rigidity, etc.), such that the changes can frequently be made without altering the biological activity or other desired property of the protein, such as antigen affinity and/or specificity. Those of skill in this art recognize that, in general, single amino acid substitutions in non-essential regions of a polypeptide do not substantially alter biological activity (*see, e.g., Watson et al. (1987) Molecular Biology of the Gene*, The Benjamin/Cummings Pub. Co., p. 224 (4th Ed.)). In addition, substitutions of structurally or functionally similar amino acids are less likely to disrupt biological activity. Exemplary conservative substitutions are set forth in Table 1 below.

Table 1. Exemplary Conservative Amino Acid Substitutions

Original residue	Conservative substitution
Ala (A)	Gly; Ser
Arg (R)	Lys; His
Asn (N)	Gln; His
Asp (D)	Glu; Asn
Cys (C)	Ser; Ala
Gln (Q)	Asn
Glu (E)	Asp; Gln
Gly (G)	Ala
His (H)	Asn; Gln
Ile (I)	Leu; Val
Leu (L)	Ile; Val
Lys (K)	Arg; His
Met (M)	Leu; Ile; Tyr
Phe (F)	Tyr; Met; Leu
Pro (P)	Ala
Ser (S)	Thr
Thr (T)	Ser
Trp (W)	Tyr; Phe
Tyr (Y)	Trp; Phe
Val (V)	Ile; Leu

[0046] “Consists essentially of,” and variations such as “consist essentially of” or “consisting essentially of,” as used throughout the specification and claims, indicate the inclusion of any recited elements or group of elements, and the optional inclusion of other elements, of similar or different nature than the recited elements, that do not materially change the basic or novel properties of the specified dosage regimen, method, or composition.

[0047] “Diagnostic anti-PD-L monoclonal antibody” means a mAb that specifically binds to the mature form of the designated PD-L (PD-L1 or PDL2) expressed on the surface of certain mammalian cells. A mature PD-L lacks the presecretory leader sequence, also referred to as leader peptide. The terms “PD-L” and “mature PD-L” are used interchangeably herein, and shall be understood to mean the same molecule unless otherwise indicated or readily apparent from the context.

[0048] As used herein, a diagnostic anti-human PD-L1 mAb or an anti-hPD-L1 mAb refers to a monoclonal antibody that specifically binds to mature human PD-L1. A mature human PD-L1 molecule consists of amino acids 19-290 set forth in SEQ ID NO 21.

[0049] Specific examples of diagnostic anti-human PD-L1 mAbs useful as diagnostic mAbs for IHC detection of PD-L1 expression in FFPE tumor tissue sections are antibodies 20C3 and 22C3, which are described in PCT International Patent Application Publication No. WO2014/100079. Another anti-human PD-L1 mAb that has been reported to be useful for IHC detection of PD-L1 expression in FFPE tissue sections (Chen, B.J. *et al.*, *Clin Cancer Res* 19: 3462-3473 (2013)) is a rabbit anti-human PD-L1 mAb publicly available from Sino Biological, Inc. (Beijing, P.R. China; Catalog number 10084-R015).

[0050] “Homology” refers to sequence similarity between two polypeptide sequences when they are optimally aligned. When a position in both of the two compared sequences is occupied by the same amino acid monomer subunit, e.g., if a position in a light chain CDR of two different Abs is occupied by alanine, then the two Abs are homologous at that position. The percent of homology is the number of homologous positions shared by the two sequences divided by the total number of positions compared $\times 100$. For example, if 8 of 10 of the positions in two sequences are matched when the sequences are optimally aligned then the two sequences are 80% homologous. Generally, the comparison is made when two sequences are aligned to give maximum percent homology. For example, the comparison can be performed by a BLAST algorithm wherein the parameters of the algorithm are selected to give the largest

match between the respective sequences over the entire length of the respective reference sequences.

[0051] The following references relate to BLAST algorithms often used for sequence analysis: BLAST ALGORITHMS: Altschul, S.F., *et al.*, (1990) J. Mol. Biol. 215:403-410; Gish, W., *et al.*, (1993) Nature Genet. 3:266-272; Madden, T.L., *et al.*, (1996) Meth. Enzymol. 266:131-141; Altschul, S.F., *et al.*, (1997) Nucleic Acids Res. 25:3389-3402; Zhang, J., *et al.*, (1997) Genome Res. 7:649-656; Wootton, J.C., *et al.*, (1993) Comput. Chem. 17:149-163; Hancock, J.M. *et al.*, (1994) Comput. Appl. Biosci. 10:67-70; ALIGNMENT SCORING SYSTEMS: Dayhoff, M.O., *et al.*, "A model of evolutionary change in proteins." in Atlas of Protein Sequence and Structure, (1978) vol. 5, suppl. 3. M.O. Dayhoff (ed.), pp. 345-352, Natl. Biomed. Res. Found., Washington, DC; Schwartz, R.M., *et al.*, "Matrices for detecting distant relationships." in Atlas of Protein Sequence and Structure, (1978) vol. 5, suppl. 3." M.O. Dayhoff (ed.), pp. 353-358, Natl. Biomed. Res. Found., Washington, DC; Altschul, S.F., (1991) J. Mol. Biol. 219:555-565; States, D.J., *et al.*, (1991) Methods 3:66-70; Henikoff, S., *et al.*, (1992) Proc. Natl. Acad. Sci. USA 89:10915-10919; Altschul, S.F., *et al.*, (1993) J. Mol. Evol. 36:290-300; ALIGNMENT STATISTICS: Karlin, S., *et al.*, (1990) Proc. Natl. Acad. Sci. USA 87:2264-2268; Karlin, S., *et al.*, (1993) Proc. Natl. Acad. Sci. USA 90:5873-5877; Dembo, A., *et al.*, (1994) Ann. Prob. 22:2022-2039; and Altschul, S.F. "Evaluating the statistical significance of multiple distinct local alignments." in Theoretical and Computational Methods in Genome Research (S. Suhai, ed.), (1997) pp. 1-14, Plenum, New York.

[0052] The term "isolated" as used in reference to an antibody or fragment thereof refers to the purification status and, in such context, means the named molecule is substantially free of other biological molecules such as nucleic acids, proteins, lipids, carbohydrates, or other material such as cellular debris and growth media. Generally, the term "isolated" is not intended to refer to a complete absence of such material or to an absence of water, buffers, or salts, unless they are present in amounts that substantially interfere with experimental or therapeutic use of the binding compound as described herein.

[0053] "Kabat" as used herein means an immunoglobulin alignment and numbering system pioneered by Elvin A. Kabat ((1991) Sequences of Proteins of Immunological Interest, 5th Ed. Public Health Service, National Institutes of Health, Bethesda, Md.).

[0054] "Monoclonal antibody" or "mAb" or "Mab", as used herein, refers to a population of substantially homogeneous antibodies, *i.e.*, the antibody molecules comprising the population

are identical in amino acid sequence except for possible naturally occurring mutations that may be present in minor amounts. In contrast, conventional (polyclonal) antibody preparations typically include a multitude of different antibodies having different amino acid sequences in their variable domains, particularly their CDRs, which are often specific for different epitopes.

5 The modifier “monoclonal” indicates the character of the antibody as being obtained from a substantially homogeneous population of antibodies, and is not to be construed as requiring production of the antibody by any particular method. For example, the monoclonal antibodies to be used in accordance with the present disclosure may be made by the hybridoma method first described by Kohler *et al.* (1975) *Nature* 256: 495, or may be made by recombinant DNA
10 methods (*see, e.g.,* U.S. Pat. No. 4,816,567). The “monoclonal antibodies” may also be isolated from phage antibody libraries using the techniques described in Clackson *et al.* (1991) *Nature* 352: 624-628 and Marks *et al.* (1991) *J. Mol. Biol.* 222: 581-597, for example. *See also* Presta (2005) *J. Allergy Clin. Immunol.* 116:731.

[0055] “RECIST 1.1 Response Criteria” as used herein means the definitions set forth in
15 Eisenhauer, E.A. *et al.*, *Eur. J. Cancer* 45:228-247 (2009) for target lesions or nontarget lesions, as appropriate based on the context in which response is being measured.

[0056] “Sustained response” means a sustained therapeutic effect after cessation of treatment as described herein. In some embodiments, the sustained response has a duration that is at least the same as the treatment duration, or at least 1.5, 2.0, 2.5 or 3 times longer than the
20 treatment duration.

[0057] “Tissue Section” refers to a single part or piece of a tissue, e.g., a thin slice of tissue cut from a sample of a normal tissue or of a tumor.

[0058] “Treat” or “treating” a cell-proliferation disorder as used herein means to administer a combination therapy of a PD-1 antagonist and a CDN STING agonist to a subject
25 having a cell-proliferation disorder, such as cancer, or diagnosed with a cell-proliferation disorder, such as cancer, to achieve at least one positive therapeutic effect, such as for example, reduced number of cancer cells, reduced tumor size, reduced rate of cancer cell infiltration into peripheral organs, or reduced rate of tumor metastasis or tumor growth. Such “treatment” may result in a slowing, interrupting, arresting, controlling, or stopping of the progression of a cell-
30 proliferation disorder as described herein but does not necessarily indicate a total elimination of the cell-proliferation disorder or the symptoms of the cell-proliferation disorder. Positive therapeutic effects in cancer can be measured in a number of ways (*See, W. A. Weber, J. Nucl.*

Med. 50:1S-10S (2009)). For example, with respect to tumor growth inhibition, according to NCI standards, a $T/C \leq 42\%$ is the minimum level of anti-tumor activity. A $T/C < 10\%$ is considered a high anti-tumor activity level, with $T/C (\%) = \text{Median tumor volume of the treated} / \text{Median tumor volume of the control} \times 100$. In some embodiments, the treatment achieved by a combination therapy of the disclosure is any of PR, CR, OR, PFS, DFS, and OS. PFS, also referred to as “Time to Tumor Progression” indicates the length of time during and after treatment that the cancer does not grow, and includes the amount of time patients have experienced a CR or PR, as well as the amount of time patients have experienced SD. DFS refers to the length of time during and after treatment that the patient remains free of disease. OS refers to a prolongation in life expectancy as compared to naive or untreated individuals or patients. In some embodiments, response to a combination therapy of the disclosure is any of PR, CR, OR, PFS, DFS, or OS that is assessed using RECIST 1.1 response criteria. The treatment regimen for a combination therapy of the disclosure that is effective to treat a cancer patient may vary according to factors such as the disease state, age, and weight of the patient, and the ability of the therapy to elicit an anti-cancer response in the subject. While an embodiment of any of the aspects of the disclosure may not be effective in achieving a positive therapeutic effect in every subject, it should do so in a statistically significant number of subjects as determined by any statistical test known in the art such as the Student’s t-test, the chi²-test, the U-test according to Mann and Whitney, the Kruskal-Wallis test (H-test), Jonckheere-Terpstra-test and the Wilcoxon-test.

[0059] As used herein, the terms “combination therapy” and “therapeutic combination” refer to treatments in which at least one PD-1 antagonist and at least one CDN STING agonist, and optionally additional therapeutic agents, each are administered to a patient in a coordinated manner, over an overlapping period of time. The period of treatment with the at least one PD-1 antagonist (the “anti-PD-1 treatment”) is the period of time that a patient undergoes treatment with the PD-1 antagonist; that is, the period of time from the initial dosing with the PD-1 antagonist through the final day of a treatment cycle. Similarly, the period of treatment with the at least one CDN STING agonist (the “CDN STING agonist treatment”) is the period of time that a patient undergoes treatment with the CDN STING agonist; that is, the period of time from the initial dosing with the CDN STING agonist through the final day of a treatment cycle. In the therapeutic combinations described herein, the anti-PD-1 treatment overlaps by at least one day the CDN STING agonist treatment. In certain embodiments, the anti-PD-1 treatment and the

CDN STING agonist treatment are coextensive. In embodiments, the anti-PD-1 treatment begins prior to the CDN STING agonist treatment. In embodiments, the CDN STING agonist treatment begins prior to the anti-PD-1 treatment. In embodiments, the anti-PD-1 treatment is terminated prior to termination of the CDN STING agonist treatment. In embodiments, the CDN STING agonist treatment is terminated prior to termination of the anti-PD-1 treatment.

[0060] The terms “treatment regimen”, “dosing protocol”, and “dosing regimen” are used interchangeably to refer to the dose and timing of administration of each therapeutic agent in a combination therapy of the disclosure.

[0061] “Tumor” as it applies to a subject diagnosed with, or suspected of having, a cancer refers to a malignant or potentially malignant neoplasm or tissue mass of any size, and includes primary tumors and secondary neoplasms. A solid tumor is an abnormal growth or mass of tissue that usually does not contain cysts or liquid areas. Different types of solid tumors are named for the type of cells that form them. Examples of solid tumors are sarcomas, carcinomas, and lymphomas. Leukemias (cancers of the blood) generally do not form solid tumors (National Cancer Institute, Dictionary of Cancer Terms).

[0062] “Advanced solid tumor malignancy” and “advanced solid tumor” are used interchangeably to refer to a tumor for which curative resection is not possible. Advanced solid tumors include, but are not limited to, metastatic tumors in bone, brain, breast, liver, lungs, lymph node, pancreas, prostate, and soft tissue (sarcoma).

[0063] “Tumor burden” also referred to as “tumor load”, refers to the total amount of tumor material distributed throughout the body. Tumor burden refers to the total number of cancer cells or the total size of tumor(s), throughout the body, including lymph nodes and bone marrow. Tumor burden can be determined by a variety of methods known in the art, such as, e.g. by measuring the dimensions of tumor(s) upon removal from the subject, e.g., using calipers, or while in the body using imaging techniques, e.g., ultrasound, bone scan, computed tomography (CT) or magnetic resonance imaging (MRI) scans.

[0064] The term “tumor size” refers to the total size of the tumor which can be measured as the length and width of a tumor. Tumor size may be determined by a variety of methods known in the art, such as, e.g. by measuring the dimensions of tumor(s) upon removal from the subject, e.g., using calipers, or while in the body using imaging techniques, e.g., bone scan, ultrasound, CT or MRI scans.

[0065] It is understood that wherever embodiments are described herein with the language “comprising,” otherwise analogous embodiments described in terms of “consisting of” and/or “consisting essentially of” are also provided.

[0066] The term “alkyl” refers to a monovalent straight or branched chain, saturated
 5 aliphatic hydrocarbon radical having a number of carbon atoms in the specified range. Thus, for example, “C₁₋₆ alkyl” (or “C₁-C₆ alkyl”) refers to any of the hexyl alkyl and pentyl alkyl isomers as well as n-, iso-, sec-, and *tert*-butyl, n- and iso-propyl, ethyl, and methyl. As another example, “C₁₋₄ alkyl” refers to n-, iso-, sec-, and *tert*-butyl, n- and isopropyl, ethyl, and methyl.

[0067] As used herein, the term “alkylene” refers to a bivalent straight chain, saturated
 10 aliphatic hydrocarbon radical having a number of carbon atoms in the specified range.

[0068] As used herein, the term “alkenyl” refers to a monovalent straight or branched chain, unsaturated aliphatic hydrocarbon radical having a number of carbon atoms in the specified range and including one or more double bond.

[0069] As used herein, the term “alkenylene” refers to a bivalent straight chain,
 15 unsaturated aliphatic hydrocarbon radical having a number of carbon atoms in the specified range and including one or more double bond.

[0070] As used herein, the term “alkynyl” refers to a monovalent straight or branched chain, unsaturated aliphatic hydrocarbon radical having a number of carbon atoms in the specified range and including one or more triple bond.

[0071] As used herein, the term “alkynylene” refers to a bivalent straight chain,
 20 unsaturated aliphatic hydrocarbon radical having a number of carbon atoms in the specified range and including one or more triple bond.

[0072] The term “halogen” (or “halo”) refers to fluorine, chlorine, bromine, and iodine (alternatively referred to as fluoro, chloro, bromo, and iodo or F, Cl, Br, and I).

[0073] The term “haloalkyl” refers to an alkyl group as defined above in which one or
 25 more of the hydrogen atoms have been replaced with a halogen. Thus, for example, “C₁₋₆ haloalkyl” (or “C₁-C₆ haloalkyl”) refers to a C₁ to C₆ linear or branched alkyl group as defined above with one or more halogen substituents. The term “fluoroalkyl” has an analogous meaning except the halogen substituents are restricted to fluoro. Suitable fluoroalkyls include the series
 30 (CH₂)₀₋₄CF₃ (*i.e.*, trifluoromethyl, 2,2,2-trifluoroethyl, 3,3,3-trifluoro-n-propyl, etc.).

[0074] As used herein, the term “haloalkenyl” refers to an alkenyl group as defined above in which one or more of the hydrogen atoms have been replaced with a halogen.

[0075] As used herein, the term “haloalkynyl” refers to an alkynyl group as defined above in which one or more of the hydrogen atoms have been replaced with a halogen.

[0076] As used herein, the term “alkoxy” as used herein, alone or in combination, includes an alkyl group connected to the oxy connecting atom. The term “alkoxy” also includes
 5 alkyl ether groups, where the term ‘alkyl’ is defined above, and ‘ether’ means two alkyl groups with an oxygen atom between them. Examples of suitable alkoxy groups include methoxy, ethoxy, n-propoxy, i-propoxy, n-butoxy, s-butoxy, t-butoxy, methoxymethane (also referred to as ‘dimethyl ether’), and methoxyethane (also referred to as ‘ethyl methyl ether’).

[0077] As used herein, the term “cycloalkyl” refers to a saturated hydrocarbon containing
 10 one ring having a specified number of carbon atoms. Examples of cycloalkyl include cyclopropyl, cyclobutyl, cyclopentyl, and cyclohexyl.

[0078] As used herein, the term “heterocycle”, “heterocyclyl”, or “heterocyclic”, as used herein, represents a stable 3- to 6-membered monocyclic that is either saturated or unsaturated, and that consists of carbon atoms and from one to two heteroatoms selected from the group
 15 consisting of N, O, and S. The heterocyclic ring may be attached at any heteroatom or carbon atom which results in the creation of a stable structure. The term includes heteroaryl moieties. Examples of such heterocyclic elements include, but are not limited to, azepinyl, benzimidazolyl, benzisoxazolyl, benzofurazanyl, benzopyranyl, benzothiopyranyl, benzofuryl, benzothiazolyl, benzothienyl, benzoxazolyl, chromanyl, cinnoliny, dihydrobenzofuryl, dihydrobenzothienyl,
 20 dihydrobenzothiopyranyl, dihydrobenzothiopyranyl sulfone, 1,3-dioxolanyl, furyl, imidazolidinyl, imidazoliny, imidazolyl, indoliny, indolyl, isochromanyl, isoindoliny, isoquinoliny, isothiazolidinyl, isothiazolyl, isothiazolidinyl, morpholiny, naphthyridinyl, oxadiazolyl, 2-oxoazepinyl, oxazolyl, 2-oxopiperazinyl, 2-oxopiperdiny, 2-oxopyrrolidinyl, piperidyl, piperazinyl, pyridyl, pyrazinyl, pyrazolidinyl, pyrazolyl, pyridazinyl, pyrimidinyl,
 25 pyrrolidinyl, pyrrolyl, quinazolinyl, quinoliny, quinoxaliny, tetrahydrofuryl, tetrahydroisoquinoliny, tetrahydroquinoliny, thiamorpholiny, thiamorpholiny sulfoxide, thiazolyl, thiazoliny, thienofuryl, thienothienyl, triazolyl and thienyl.

[0079] As used herein, the term “fused ring” refers to a cyclic group formed by substituents on separate atoms in a straight or branched alkane or alkene, or to a cyclic group
 30 formed by substituents on separate atoms in another ring.

[0080] As used herein, the term “spirocycle” or “spirocyclic ring” refers to a pendant cyclic group formed by substituents on a single atom.

[0081] Unless expressly stated to the contrary, all ranges cited herein are inclusive; *i.e.*, the range includes the values for the upper and lower limits of the range as well as all values in between. As an example, temperature ranges, percentages, ranges of equivalents, and the like described herein include the upper and lower limits of the range and any value in the continuum there between. Numerical values provided herein, and the use of the term “about”, may include variations of $\pm 1\%$, $\pm 2\%$, $\pm 3\%$, $\pm 4\%$, $\pm 5\%$, $\pm 10\%$, $\pm 15\%$, and $\pm 20\%$ and their numerical equivalents. All ranges also are intended to include all included sub-ranges, although not necessarily explicitly set forth. For example, a range of 3 to 7 days is intended to include 3, 4, 5, 6, and 7 days. In addition, the term “or,” as used herein, denotes alternatives that may, where appropriate, be combined; that is, the term “or” includes each listed alternative separately as well as their combination.

[0082] Where aspects or embodiments of the disclosure are described in terms of a Markush group or other grouping of alternatives, the present disclosure encompasses not only the entire group listed as a whole, but each member of the group individually and all possible subgroups of the main group, but also the main group absent one or more of the group members. The present disclosure also envisages the explicit exclusion of one or more of any of the group members in the claims.

[0083] Unless otherwise defined, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which this disclosure relates. In case of conflict, the present specification, including definitions, will control. Throughout this specification and claims, the word “comprise,” or variations such as “comprises” or “comprising” will be understood to imply the inclusion of a stated integer or group of integers but not the exclusion of any other integer or group of integers. Unless otherwise required by context, singular terms shall include pluralities and plural terms shall include the singular. Any example(s) following the term “e.g.” or “for example” is not meant to be exhaustive or limiting.

[0084] Exemplary methods and materials are described herein, although methods and materials similar or equivalent to those described herein can also be used in the practice or testing of the present disclosure. The materials, methods, and examples are illustrative only and not intended to be limiting.

[0085] The present disclosure relates to methods of treating a cell-proliferation disorder as defined herein, wherein the method comprises administering to a subject in need thereof a

combination therapy that comprises (a) a PD-1 antagonist; and (b) a cyclic dinucleotide STING agonist.

[0086] The present disclosure relates to methods of treating a cell-proliferation disorder, wherein the method comprises administering to a subject in need thereof a combination therapy that comprises (a) a PD-1 antagonist; and (b) a cyclic dinucleotide STING agonist; wherein the cell-proliferation disorder is selected from the group consisting of solid tumors and lymphomas.

PD-1 ANTAGONIST

[0087] “PD-1 antagonist” or “PD-1 pathway antagonist” means any chemical compound or biological molecule that blocks binding of PD-L1 expressed on a cancer cell to PD-1 expressed on an immune cell (T-cell, B-cell, or NKT-cell) and preferably also blocks binding of PD-L2 expressed on a cancer cell to the immune-cell expressed PD-1. Alternative names or synonyms for PD-1 and its ligands include: PDCD1, PD1, CD279, and SLEB2 for PD-1; PDCD1L1, PDL1, B7H1, B7-4, CD274, and B7-H for PD-L1; and PDCD1L2, PDL2, B7-DC, Btdc, and CD273 for PD-L2. In any of the treatment methods, medicaments and uses of the present disclosure in which a human individual is being treated, the PD-1 antagonist blocks binding of human PD-L1 to human PD-1, and preferably blocks binding of both human PD-L1 and PD-L2 to human PD-1. Human PD-1 amino acid sequences can be found in NCBI Locus No.: NP_005009. Human PD-L1 and PD-L2 amino acid sequences can be found in NCBI Locus No.: NP_054862 and NP_079515, respectively, and in SEQ ID NO: 21.

[0088] PD-1 antagonists useful in any of the treatment methods, medicaments and uses of the present disclosure include a mAb, or antigen binding fragment thereof, which specifically binds to PD-1 or PD-L1, and preferably specifically binds to human PD-1 or human PD-L1. The mAb may be a human antibody, a humanized antibody, or a chimeric antibody and may include a human constant region. In some embodiments, the human constant region is selected from the group consisting of IgG1, IgG2, IgG3, and IgG4 constant regions, and in specific embodiments, the human constant region is an IgG1 or IgG4 constant region. In some embodiments, the antigen binding fragment is selected from the group consisting of Fab, Fab'-SH, F(ab')₂, scFv, and Fv fragments.

[0089] Examples of mAbs that bind to human PD-1, and that may be useful in the treatment methods, medicaments, and uses of the present disclosure, are described in U.S. Patent Nos. US7488802, US7521051, US8008449, US8354509, and US8168757, PCT International

Patent Application Publication Nos. WO2004/004771, WO2004/072286, and WO2004/056875, and U.S. Patent Application Publication No. US20110271358.

[0090] Examples of mAbs that bind to human PD-L1, and that may be useful in the treatment methods, medicaments and uses of the present disclosure, are described in PCT

5 International Patent Application Nos. WO2013/019906 and WO2010/077634 and in U.S. Patent No. US8383796. Specific anti-human PD-L1 mAbs useful as the PD-1 antagonist in the treatment methods, medicaments, and uses of the present disclosure include MPDL3280A, BMS-936559, MEDI4736, MSB0010718C, and an antibody that comprises the heavy chain and light chain variable regions of SEQ ID NO:24 and SEQ ID NO:21, respectively, of
10 WO2013/019906. In particular embodiments, the PD-1 antagonist is an antigen binding fragment having variable regions comprising the heavy and light chain CDRs of WO2013/019906.

[0091] Other PD-1 antagonists useful in any of the treatment methods, medicaments, and

uses of the present disclosure include an immune-adhesion molecule that specifically binds to
15 PD-1 or PD-L1, and preferably specifically binds to human PD-1 or human PD-L1, *e.g.*, a fusion protein containing the extracellular or PD-1 binding portion of PD-L1 or PD-L2 fused to a constant region such as an Fc region of an immunoglobulin molecule. Examples of immune-adhesion molecules that specifically bind to PD-1 are described in PCT International Patent Application Publication Nos. WO2010/027827 and WO2011/066342. Specific fusion proteins
20 useful as the PD-1 antagonist in the treatment methods, medicaments, and uses of the present disclosure include AMP-224 (also known as B7-DCIg), which is a PD-L2-FC fusion protein and binds to human PD-1.

[0092] In embodiments, the PD-1 antagonist can be conjugated, *e.g.*, to small drug molecules, enzymes, liposomes, polyethylene glycol (PEG).

25 [0093] In some embodiments of the treatment methods, medicaments, and uses of the present disclosure, the PD-1 antagonist is a monoclonal antibody, or antigen binding fragment thereof, which specifically binds to human PD-1 and comprises (a) a heavy chain variable region comprising CDRH1 of SEQ ID NO 8, CDRH2 of SEQ ID NO 9, and CDRH3 of SEQ ID NO 10, and (b) a light chain variable region comprising CDRL1 of SEQ ID NO 5, CDRL2 of SEQ ID
30 NO 6, and CDRL3 of SEQ ID NO 7. In specific embodiments, the PD-1 antagonist is a monoclonal antibody, or antigen binding fragment thereof, which specifically binds to human PD-1 and comprises (a) a heavy chain variable region comprising SEQ ID NO 2, and (b) a light

chain variable region comprising SEQ ID NO 1. In specific embodiments, the PD-1 antagonist is a monoclonal antibody, or antigen binding fragment thereof, which specifically binds to human PD-1 and comprises (a) a heavy chain comprising SEQ ID NO 4, and (b) a light chain comprising SEQ ID NO 3.

5 [0094] In some embodiments of the treatment methods, medicaments, and uses of the present disclosure, the PD-1 antagonist is a monoclonal antibody, or antigen binding fragment thereof, which specifically binds to human PD-1 and comprises (a) a heavy chain variable region comprising CDRH1 of SEQ ID NO 18, CDRH2 of SEQ ID NO 19, and CDRH3 of SEQ ID NO 20, and (b) a light chain variable region comprising CDRL1 of SEQ ID NO 15, CDRL2 of SEQ
10 ID NO 16, and CDRL3 of SEQ ID NO 17. In specific embodiments, the PD-1 antagonist is a monoclonal antibody, or antigen binding fragment thereof, which specifically binds to human PD-1 and comprises (a) a heavy chain variable region comprising SEQ ID NO 12, and (b) a light chain variable region comprising SEQ ID NO 11. In specific embodiments, the PD-1 antagonist is a monoclonal antibody, or antigen binding fragment thereof, which specifically binds to
15 human PD-1 and comprises (a) a heavy chain comprising SEQ ID NO 14, and (b) a light chain comprising SEQ ID NO 13.

[0095] In some embodiments of the treatment methods, medicaments, and uses of the present disclosure, the PD-1 antagonist is an anti-PD-1 monoclonal antibody. In aspects of these embodiments, the PD-1 antagonist is selected from the group consisting of nivolumab,
20 pembrolizumab, pidilizumab, and AMP-224. In specific aspects, the PD-1 antagonist is selected from nivolumab and pembrolizumab. In a more specific aspect, the PD-1 antagonist is nivolumab. In a further specific aspect, the PD-1 antagonist is pembrolizumab.

[0096] The present disclosure relates to PD-1 antagonists that are monoclonal antibodies, or antigen binding fragments thereof, which specifically bind to human PD-1 as described
25 herein. In embodiments, PD-1 antagonists may comprise variant heavy chain variable region sequence and/or variant light chain variable region sequence identical to the reference sequence except having up to 17 conservative amino acid substitutions in the framework region (i.e., outside of the CDRs), and preferably have less than ten, nine, eight, seven, six, or five conservative amino acid substitutions in the framework region.

30 [0097] Table 2 below provides a list of the amino acid sequences of exemplary anti-PD-1 mAbs for use in the treatment methods, medicaments, and uses of the present disclosure, and the sequences are shown in Figures 1-9.

Table 2. Description of Sequences in Sequence Listing

SEQ ID NO:	Description
1	Pembrolizumab, variable light chain, amino acid
2	Pembrolizumab, variable heavy chain, amino acid
3	Pembrolizumab, light chain, amino acid
4	Pembrolizumab, heavy chain, amino acid
5	Pembrolizumab, CDRL1
6	Pembrolizumab, CDRL2
7	Pembrolizumab, CDRL3
8	Pembrolizumab, CDRH1
9	Pembrolizumab, CDRH2
10	Pembrolizumab, CDRH3
11	Nivolumab, variable light chain, amino acid
12	Nivolumab, variable heavy chain, amino acid
13	Nivolumab, light chain, amino acid
14	Nivolumab, heavy chain, amino acid
15	Nivolumab, CDRL1
16	Nivolumab, CDRL2
17	Nivolumab, CDRL3
18	Nivolumab, CDRH1
19	Nivolumab, CDRH2
20	Nivolumab, CDRH3
21	Human PD-L1

[0098] “PD-L1” expression or “PD-L2” expression as used herein means any detectable level of expression of the designated PD-L protein on the cell surface or of the designated PD-L mRNA within a cell or tissue. PD-L protein expression may be detected with a diagnostic PD-L antibody in an IHC assay of a tumor tissue section or by flow cytometry. Alternatively, PD-L protein expression by tumor cells may be detected by PET imaging, using a binding agent (e.g., antibody fragment, affibody, and the like) that specifically binds to the desired PD-L target, e.g.,

PD-L1 or PD-L2. Techniques for detecting and measuring PD-L mRNA expression include RT-PCR and realtime quantitative RT-PCR.

[0099] Several approaches have been described for quantifying PD-L1 protein expression in IHC assays of tumor tissue sections. *See, e.g.,* Thompson, R. H., *et al.*, *PNAS* **101** (49); 17174-17179 (2004); Thompson, R. H. *et al.*, *Cancer Res.* **66**:3381-3385 (2006); Gadiot, J., *et al.*, *Cancer* **117**:2192-2201 (2011); Taube, J. M. *et al.*, *Sci Transl Med* **4**, 127ra37 (2012); and Toplian, S. L. *et al.*, *New Eng. J Med.* **366** (26): 2443-2454 (2012).

[0100] One approach employs a simple binary end-point of positive or negative for PD-L1 expression, with a positive result defined in terms of the percentage of tumor cells that exhibit histologic evidence of cell-surface membrane staining. A tumor tissue section is counted as positive for PD-L1 expression is at least 1%, and preferably 5% of total tumor cells.

[0101] In another approach, PD-L1 expression in the tumor tissue section is quantified in the tumor cells as well as in infiltrating immune cells, which predominantly comprise lymphocytes. The percentage of tumor cells and infiltrating immune cells that exhibit membrane staining are separately quantified as < 5%, 5 to 9%, and then in 10% increments up to 100%. For tumor cells, PD-L1 expression is counted as negative if the score is < 5% score and positive if the score is $\geq$ 5%. PD-L1 expression in the immune infiltrate is reported as a semi-quantitative measurement called the adjusted inflammation score (AIS), which is determined by multiplying the percent of membrane staining cells by the intensity of the infiltrate, which is graded as none (0), mild (score of 1, rare lymphocytes), moderate (score of 2, focal infiltration of tumor by lymphohistiocytic aggregates), or severe (score of 3, diffuse infiltration). A tumor tissue section is counted as positive for PD-L1 expression by immune infiltrates if the AIS is $\geq$ 5.

[0102] The level of PD-L1 mRNA expression may be compared to the mRNA expression levels of one or more reference genes that are frequently used in quantitative RT-PCR, such as ubiquitin C.

[0103] In some embodiments, a level of PD-L1 expression (protein and/or mRNA) by malignant cells and/or by infiltrating immune cells within a tumor is determined to be “overexpressed” or “elevated” based on comparison with the level of PD-L1 expression (protein and/or mRNA) by an appropriate control. For example, a control PD-L1 protein or mRNA expression level may be the level quantified in nonmalignant cells of the same type or in a section from a matched normal tissue. In some embodiments, PD-L1 expression in a tumor

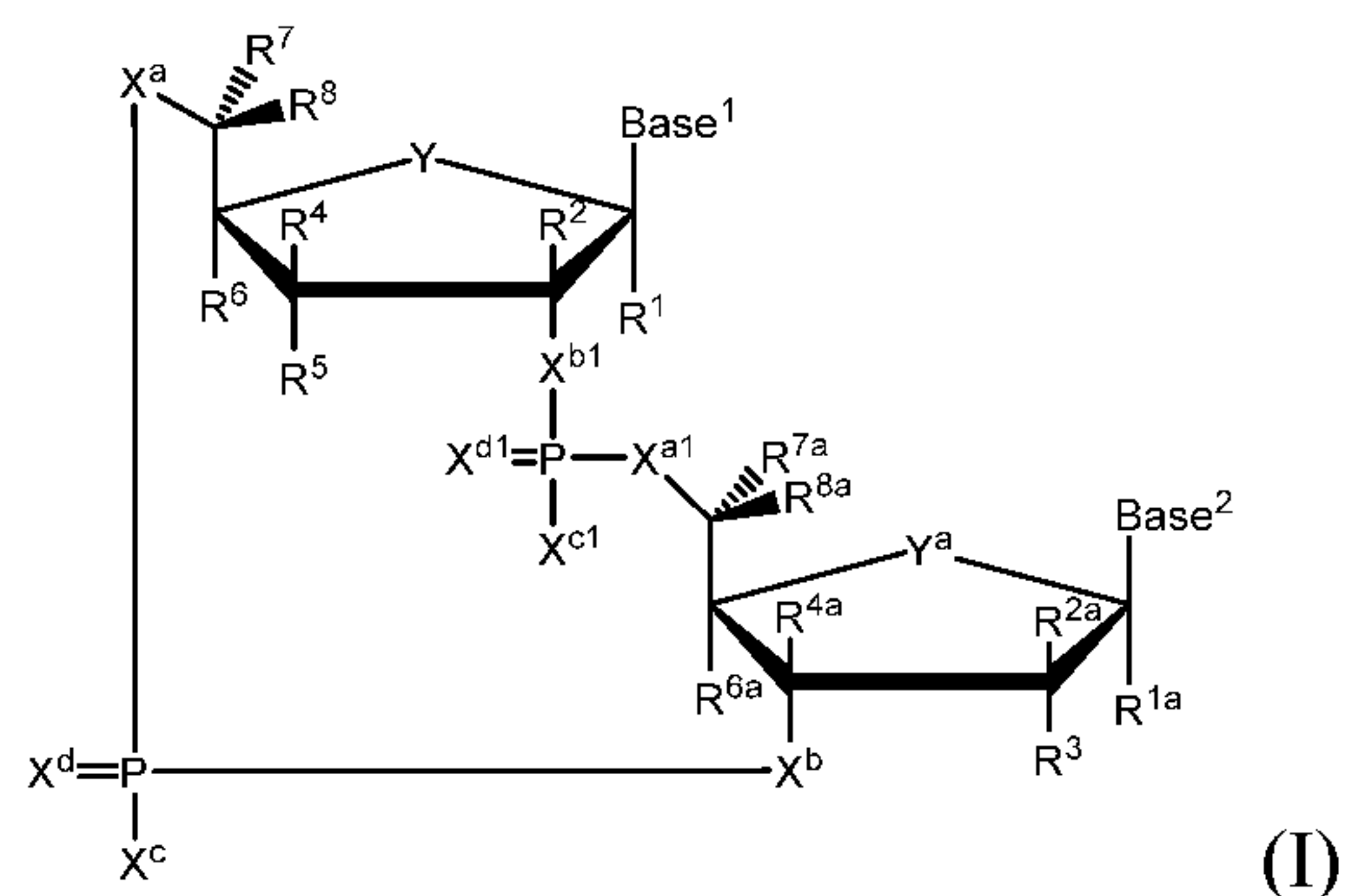
sample is determined to be elevated if PD-L1 protein (and/or PD-L1 mRNA) in the sample is at least 10%, 20%, or 30% greater than in the control.

[0104] In embodiments, PD-1 antagonists disclosed herein may be provided by continuous infusion, or by doses administered, e.g., daily, 1-7 times per week, weekly, bi-weekly, monthly, bimonthly, quarterly, semiannually, annually etc. Doses may be provided, e.g., intravenously, subcutaneously, topically, orally, nasally, rectally, intramuscular, intracerebrally, intraspinally, or by inhalation. A total dose for a treatment interval is generally at least 0.05 µg/kg body weight, more generally at least 0.2 µg/kg, 0.5 µg/kg, 1 µg/kg, 10 µg/kg, 100 µg/kg, 0.25 mg/kg, 1.0 mg/kg, 2.0 mg/kg, 5.0 mg/ml, 10 mg/kg, 25 mg/kg, 50 mg/kg or more (*see, e.g.,* Yang, *et al.* (2003) New Engl. J. Med. 349:427-434; Herold, *et al.* (2002) New Engl. J. Med. 346:1692-1698; Liu, *et al.* (1999) J. Neurol. Neurosurg. Psych. 67:451-456; Portielji, *et al.* (20003) Cancer Immunol. Immunother. 52:133-144). Doses may also be provided to achieve a pre-determined target concentration of PD-1 antagonists in the subject's serum, such as 0.1, 0.3, 1, 3, 10, 30, 100, 300 µg/mL or more. In embodiments, the PD-1 antagonist is administered as a 200mg dose once every 21 days. In other embodiments, PD-1 antagonists are administered subcutaneously or intravenously, on a weekly, biweekly, "every 4 weeks," monthly, bimonthly, or quarterly basis at 10, 20, 50, 80, 100, 200, 500, 1000 or 2500 mg/subject.

CYCLIC DINUCLEOTIDE STING AGONISTS

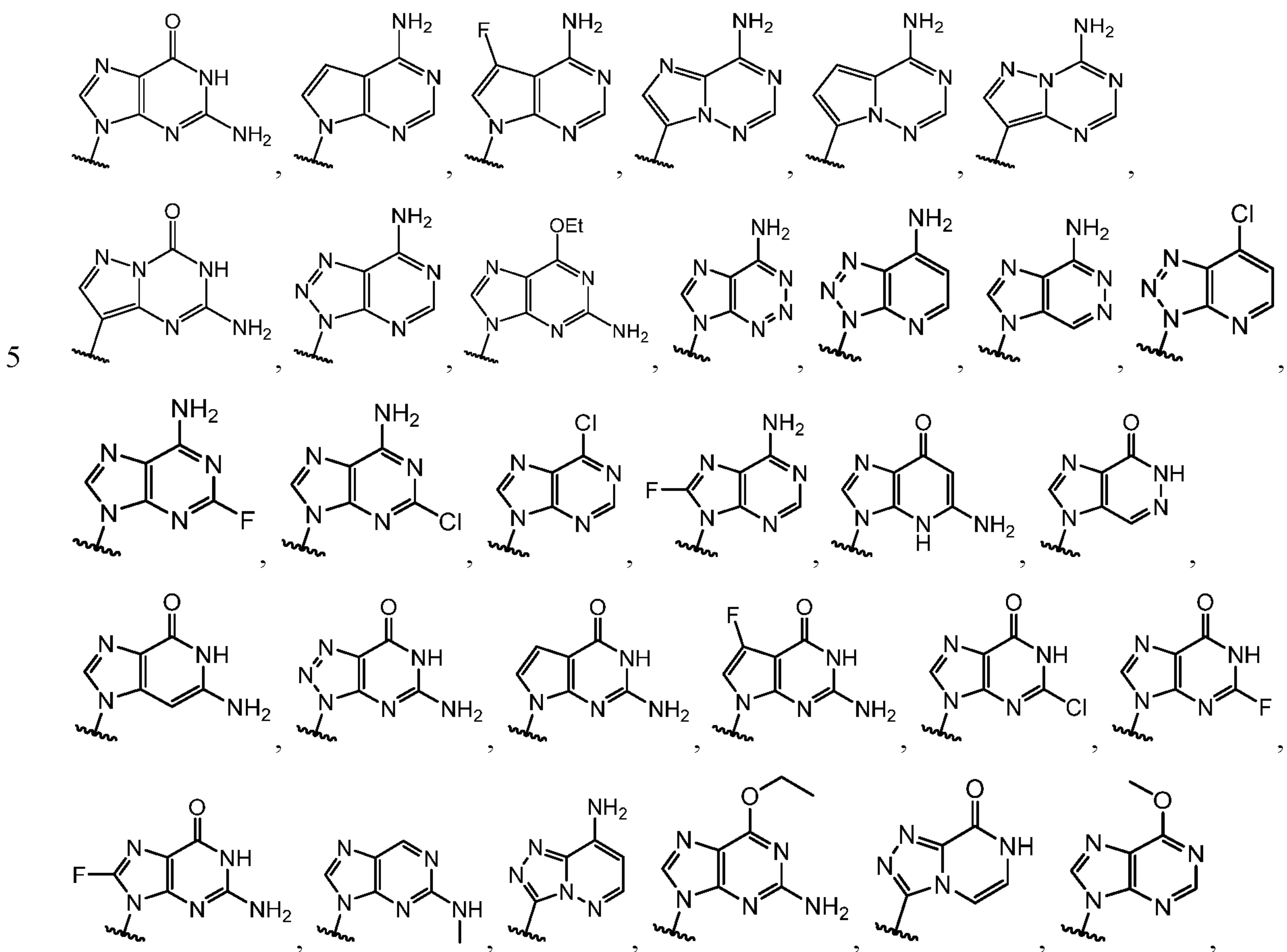
[0105] As used herein, "CDN STING agonist" means any cyclic dinucleotide chemical compound that activates the STING pathway, and in particular, the cyclic dinucleotide STING agonists as disclosed in PCT International Patent Application No. PCT/US2016/046444, which published as PCT International Patent Application Publication No. WO2017/027646, and United States Patent Application No. 15/234,182, which published as U.S. Patent Application Publication No. US2017/0044206, both of which are incorporated herein in their entirety. CDN STING agonists, and particularly the compounds of formulas (I), (Ia), (Ib), (Ic), (I'), (I'a), (I'b), (I'c), (I''), (I''a), and (I''b), may be used in the therapeutic combinations of this disclosure.

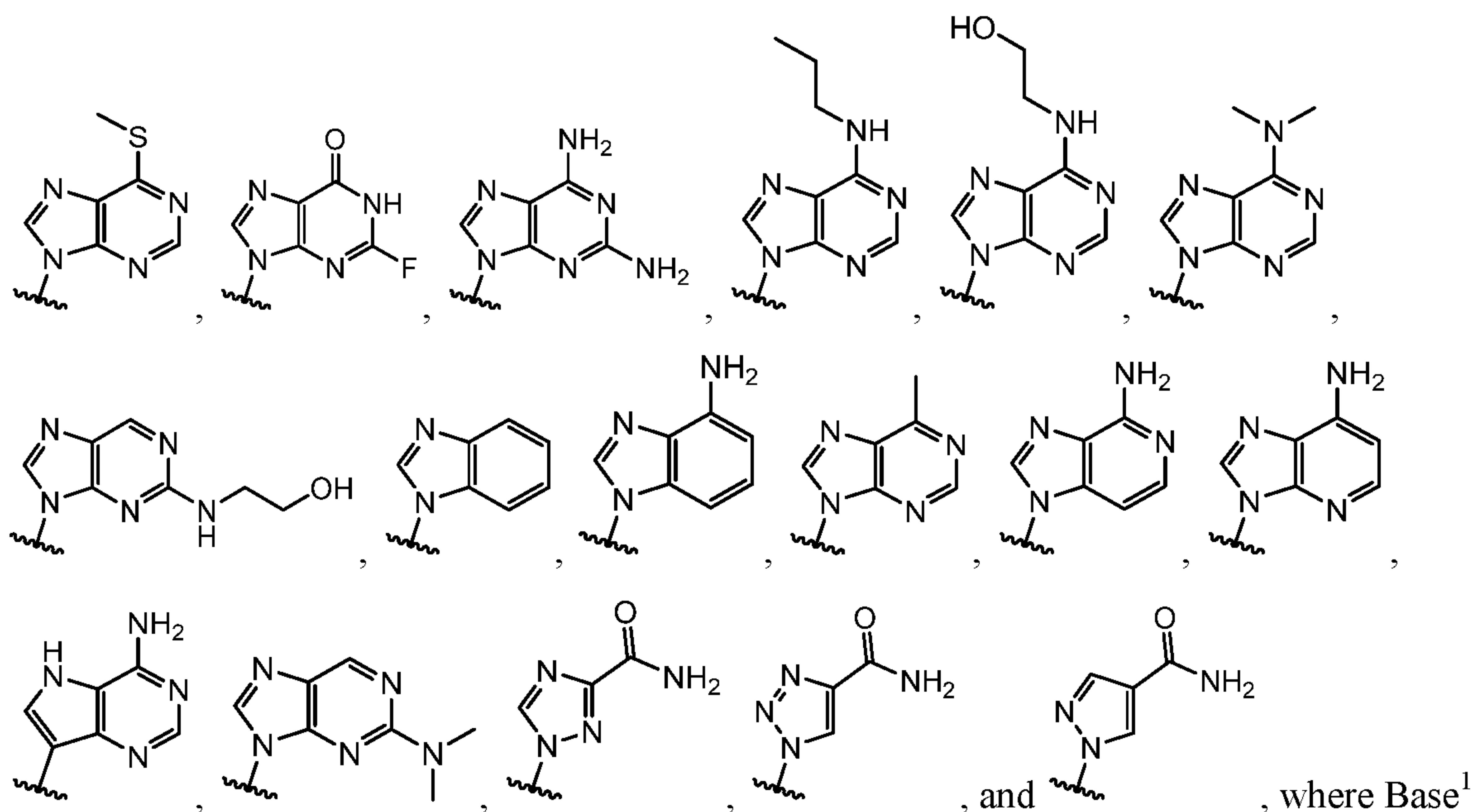
[0106] In embodiments, the CDN STING agonist is selected from cyclic di-nucleotide compounds of formula (I):



or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein Base¹ and

Base² are each independently selected from the group consisting of





, where Base¹ and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is

- 5 independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂; Y and Y^a are each independently selected from the group consisting of -O-, -S-, -SO₂-, -CH₂-, and -CF₂-; X^a and X^{a1} are each independently selected from the group consisting of O, C, and S; X^b and X^{b1} are each
- 10 independently selected from the group consisting of O, C, and S; X^c and X^{c1} are each independently selected from the group consisting of OR⁹, SR⁹, and NR⁹R⁹; X^d and X^{d1} are each independently selected from the group consisting of O and S; R¹ and R^{1a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆
- 15 alkenyl, and -O-C₂-C₆ alkynyl, where said R¹ and R^{1a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R² and R^{2a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆
- 20 haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R² and R^{2a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R³ is

selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R³ C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and

5 -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁴ and R^{4a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁴ and R^{4a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl,

10 C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁵ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁵ C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆

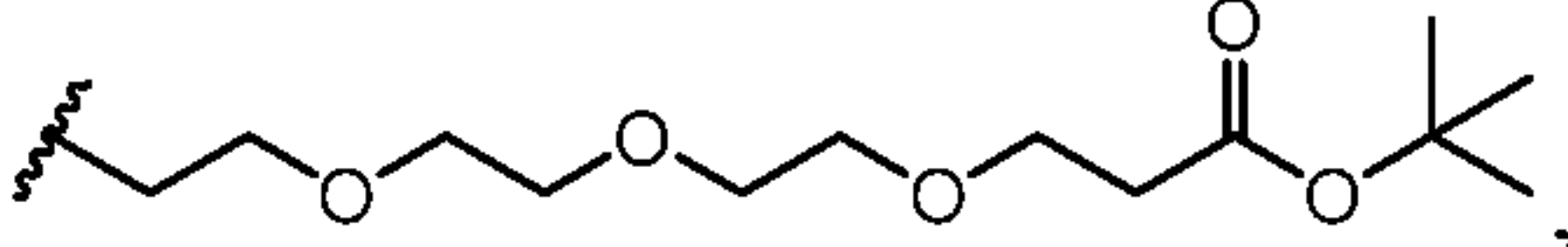
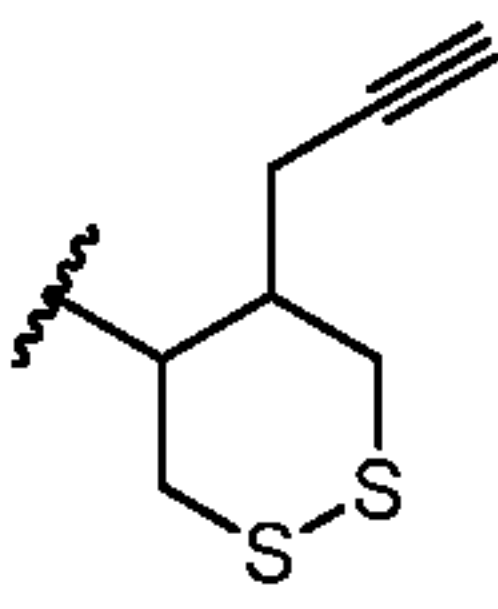
15 haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁶ and R^{6a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where

20 said R⁶ and R^{6a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁷ and R^{7a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl,

25 -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁷ and R^{7a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁸ and R^{8a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl,

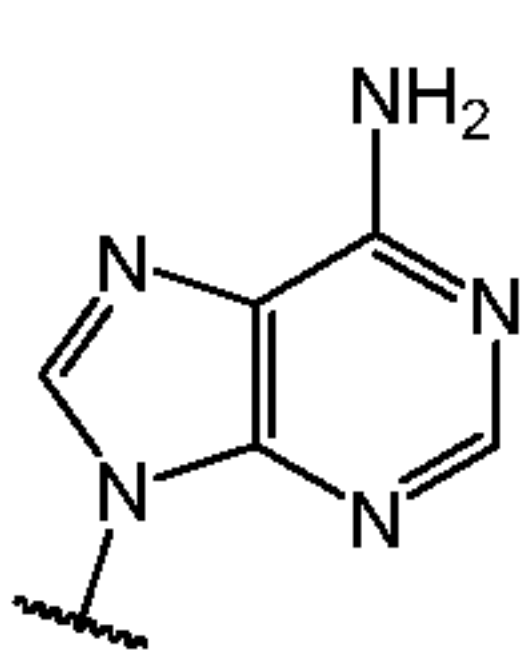
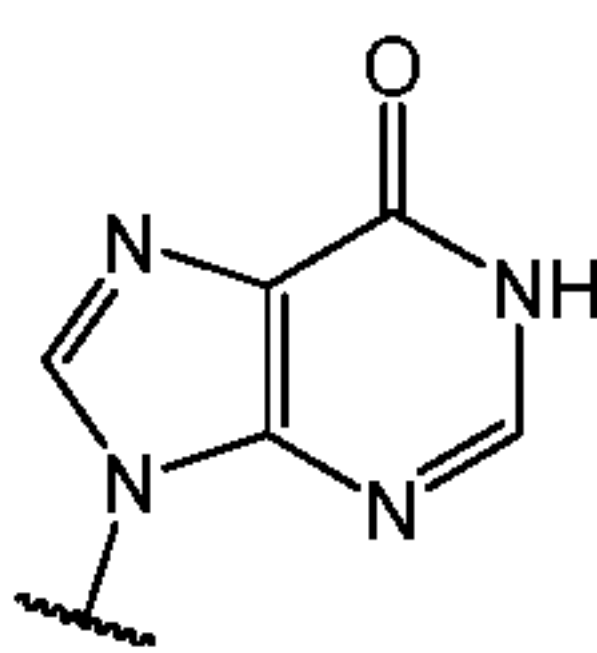
30 C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁸ and R^{8a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and

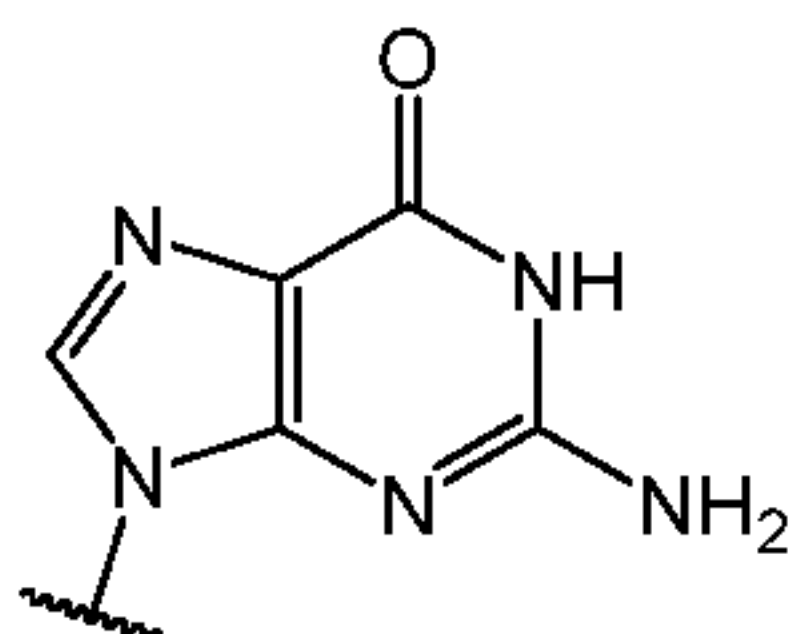
-O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; each R⁹ is independently selected from the group consisting of H,

C₁-C₂₀ alkyl, , and , where each R⁹ C₁-C₂₀ alkyl is

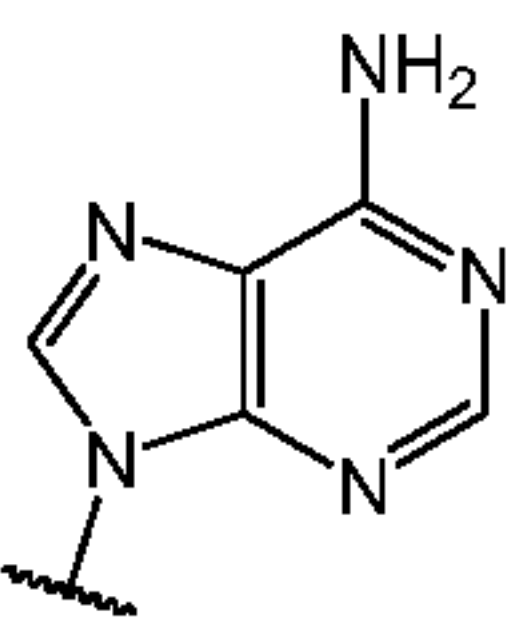
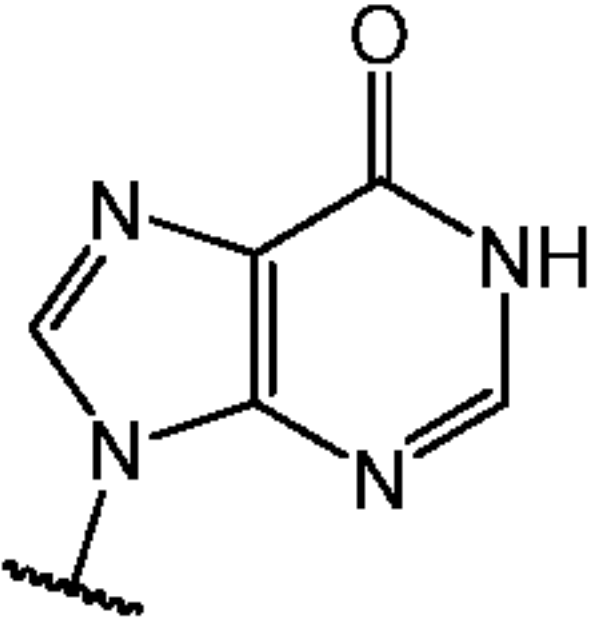
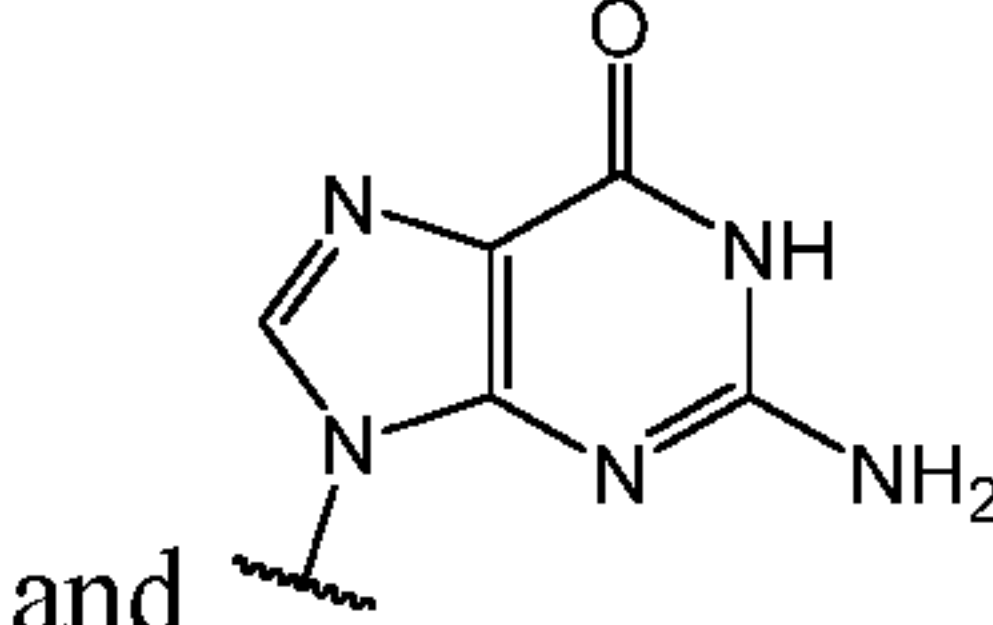
optionally substituted by 0 to 3 substituents independently selected from the group consisting of
 5 OH, -O-C₁-C₂₀ alkyl, -S-C(O)C₁-C₆ alkyl, and -C(O)OC₁-C₆ alkyl; optionally R^{1a} and R³ are
 connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene, -O-C₁-C₆ alkylene,
 -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R^{1a} and R³ are connected to form
 -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³
 position; optionally R^{2a} and R³ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆
 10 alkynylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where
 R^{2a} and R³ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆
 alkynylene, said O is bound at the R³ position; optionally R³ and R^{6a} are connected to form
 -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R³ and R^{6a} are
 connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is
 15 bound at the R³ position; optionally R⁴ and R⁵ are connected to form are connected to form
 C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or
 -O-C₂-C₆ alkynylene, such that where R⁴ and R⁵ are connected to form -O-C₁-C₆ alkylene,
 -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R⁵ position; optionally R⁵
 and R⁶ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene,
 20 such that where R⁵ and R⁶ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or
 -O-C₂-C₆ alkynylene, said O is bound at the R⁵ position; optionally R⁷ and R⁸ are connected to
 form C₁-C₆ alkylene, C₂-C₆ alkenylene, or C₂-C₆ alkynylene; and optionally R^{7a} and R^{8a} are
 connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, or C₂-C₆ alkynylene.

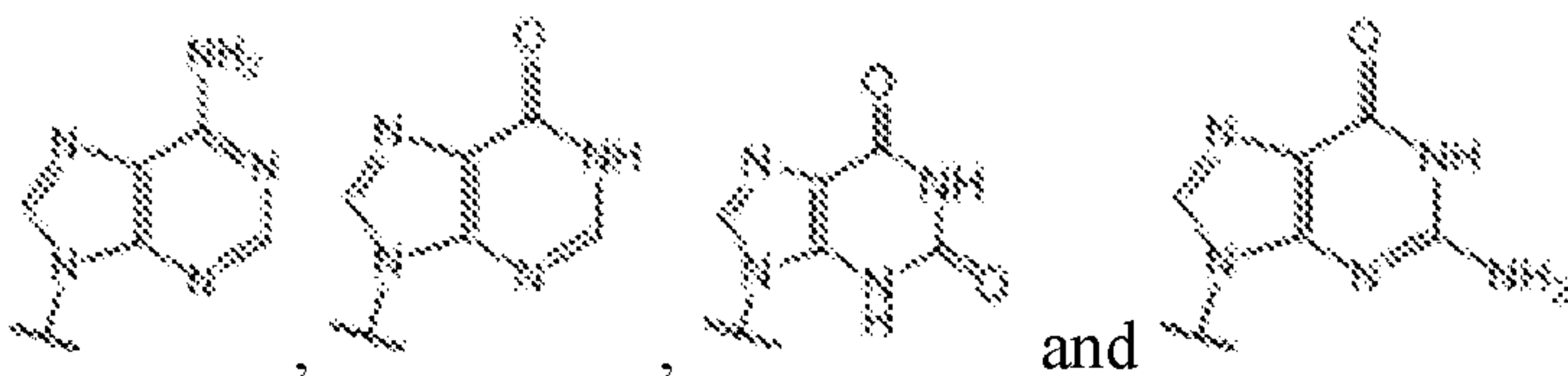
[0107] In specific aspects of this embodiment, when Y and Y^a are each O, X^a and X^{a1} are
 25 each O, X^b and X^{b1} are each O, and X^c and X^{c1} are each OH or SH, X^d and X^{d1} are each O, R¹
 and R^{1a} are each H, R² is H, R⁶ and R^{6a} are each H, R⁷ and R^{7a} are each H, R⁸ and R^{8a} are each H,

and Base¹ and Base² are each selected from the group consisting of , , and



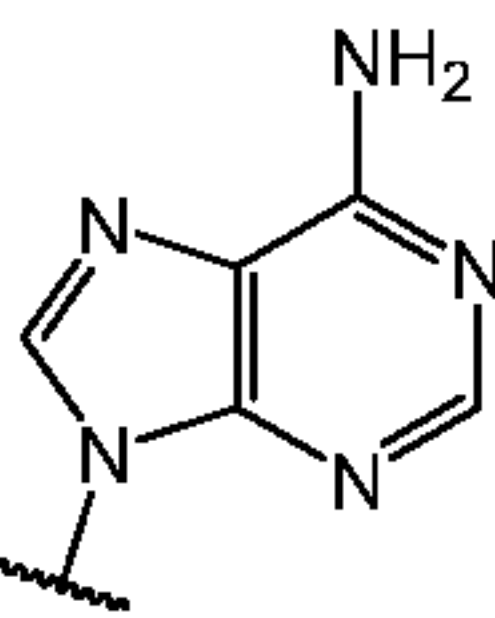
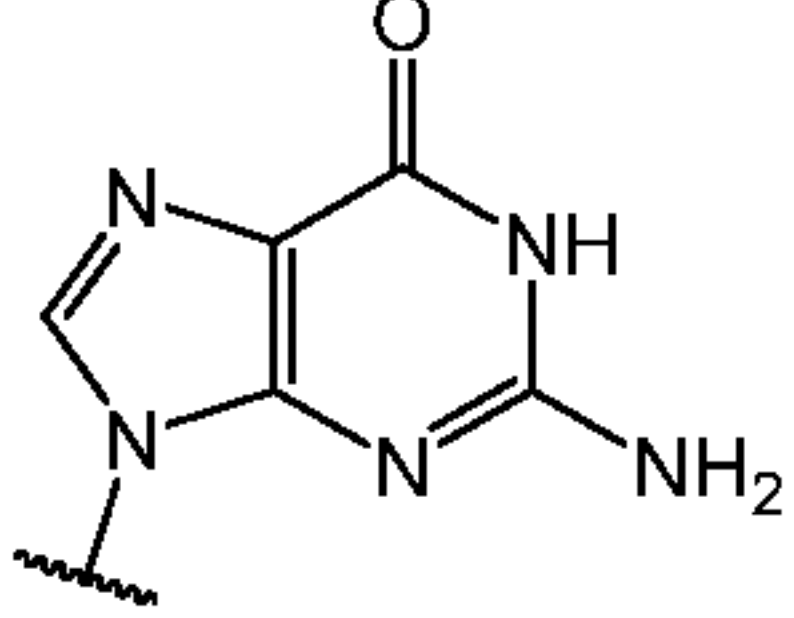
, R^5 and R^3 are not both selected from the group consisting of H, F and OH. That is, when Y and Y^a are each O, X^a and X^{a1} are each O, X^b and X^{b1} are each O, and X^c and X^{c1} are each OH or SH, X^d and X^{d1} are each O, R^1 and R^{1a} are each H, R^2 is H, R^6 and R^{6a} are each H, R^7 and R^{7a} are each H, R^8 and R^{8a} are each H, and $Base^1$ and $Base^2$ are each selected from the group

5 consisting of , , and , either only one of R^5 and R^3 is selected from the group consisting of H, F, and OH, or neither R^5 and R^3 is selected from the group consisting of H, F, and OH. In specific instances of this aspect, when Y and Y^a are each O, X^a and X^{a1} are each O, X^b and X^{b1} are each O, and X^c and X^{c1} are each OH, X^d and X^{d1} are each O or S, R^1 and R^{1a} are each H, R^2 is H, R^6 and R^{6a} are each H, R^7 and R^{7a} are each H, R^8 and R^{8a} are each H, and $Base^1$ and $Base^2$ are each selected from the group consisting of

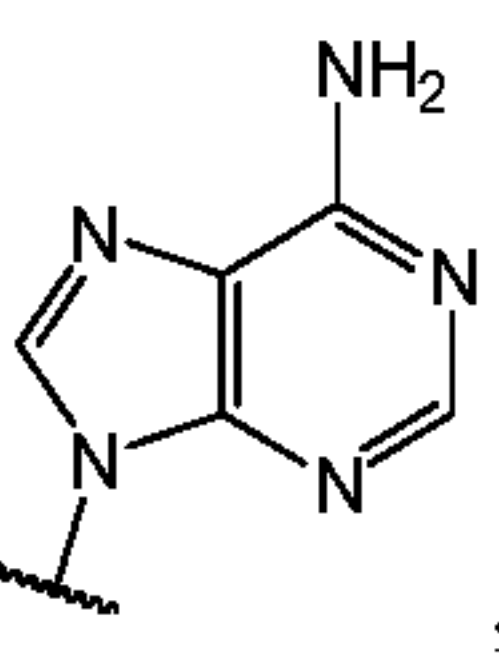
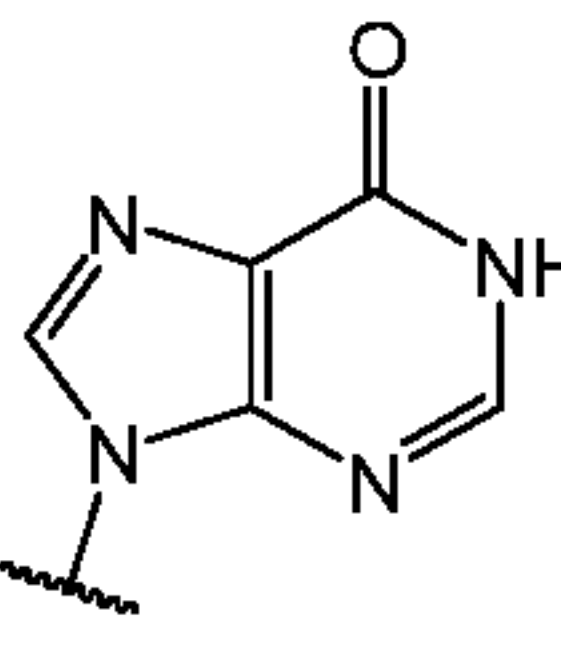
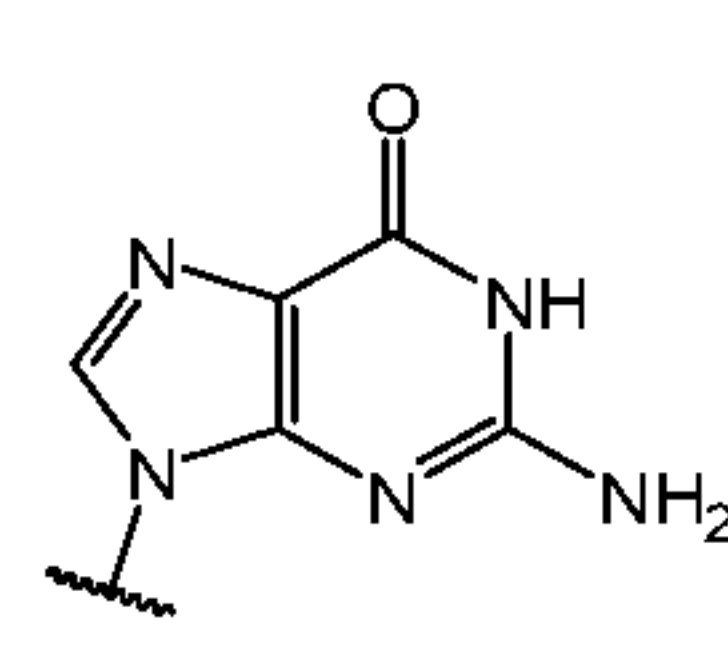
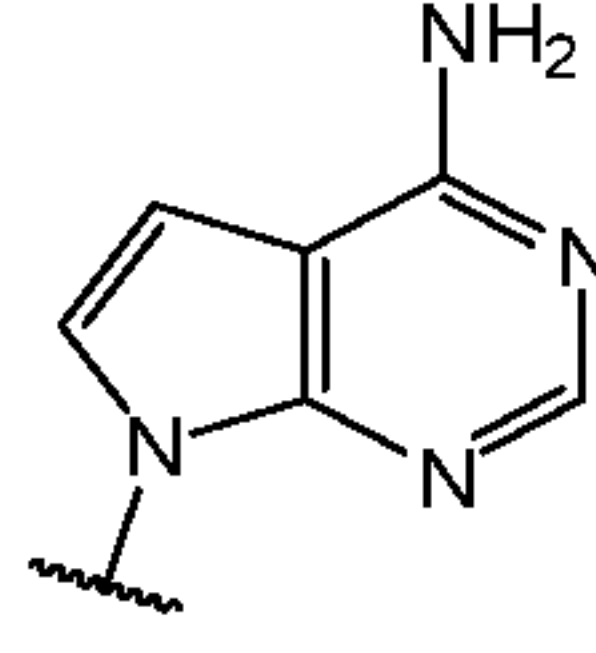
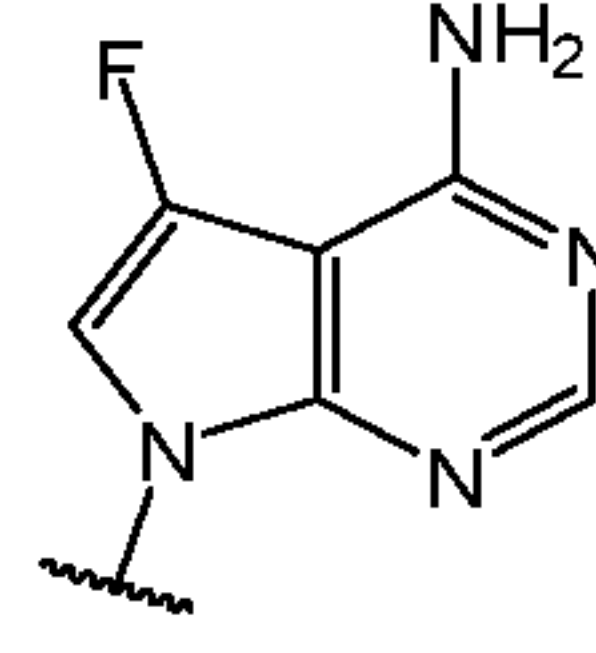


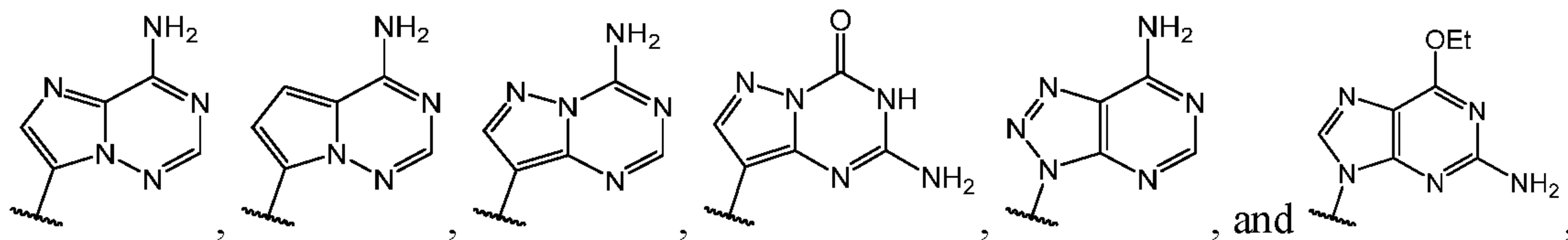
and R^5 and R^3 are not both selected from the group consisting of H, C_1 - C_6 alkyl, C_2 - C_6 alkenyl, C_2 - C_6 alkynyl, where said C_1 - C_6 alkyl, C_2 - C_6 alkenyl and C_2 - C_6 alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I and OH.

15 [0108] In further specific aspects of this embodiment, when $Base^1$ and $Base^2$ are each

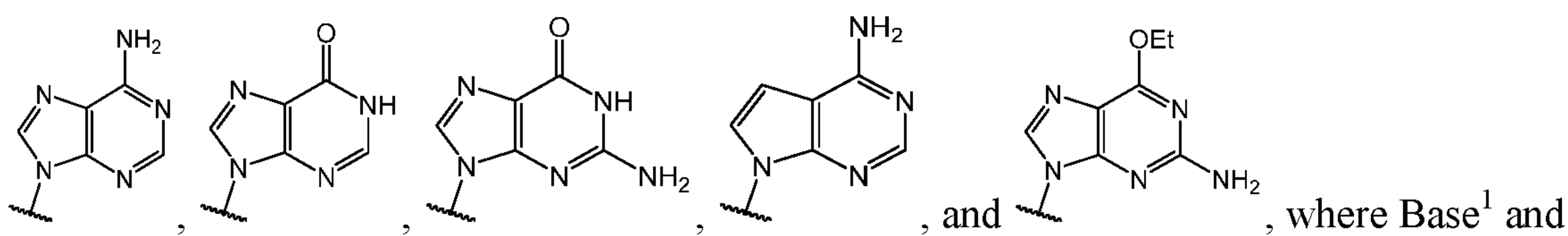
selected from the group consisting of  and , and R^{2a} is F and R^5 is F, at least one of X^c and X^{c1} is SR^9 .

[0109] In aspects of this embodiment, $Base^1$ and $Base^2$ are each independently selected

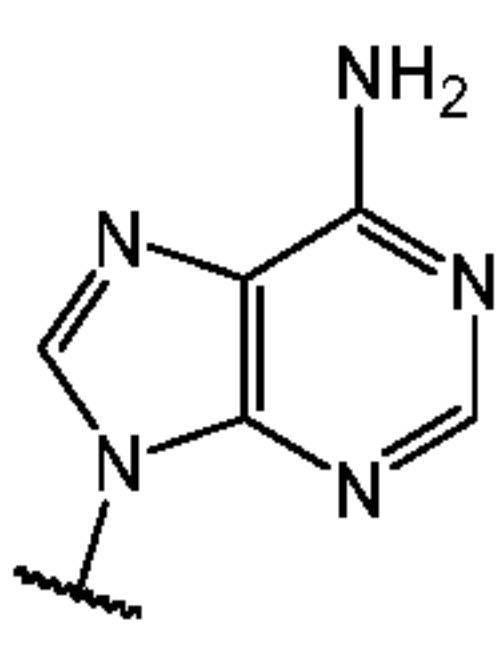
from the group consisting of , , , , ,

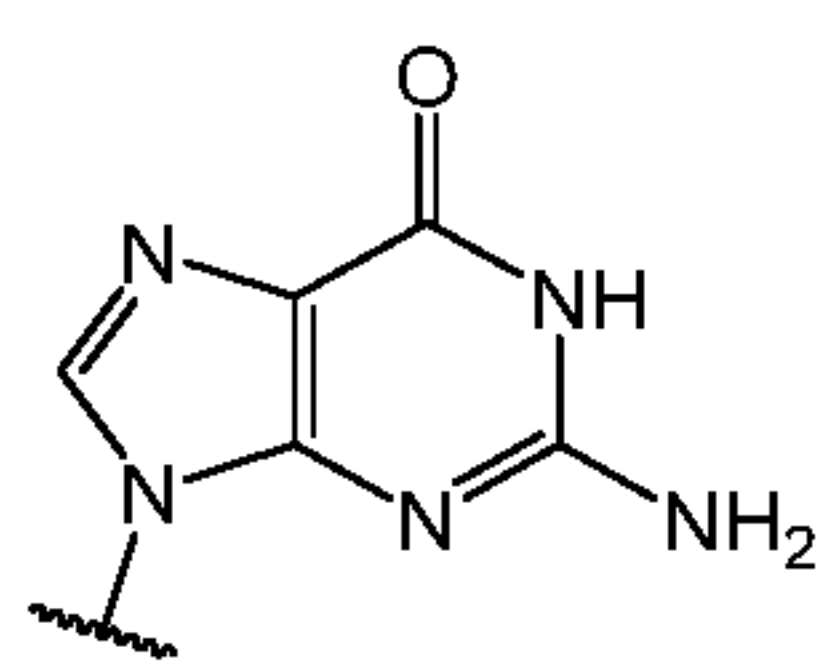


where Base¹ and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂. In particular instances, Base¹ and Base² are each independently selected from the group consisting of



where Base¹ and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂. In even more particular instances,

Base¹ and Base² are each independently selected from the group consisting of  and



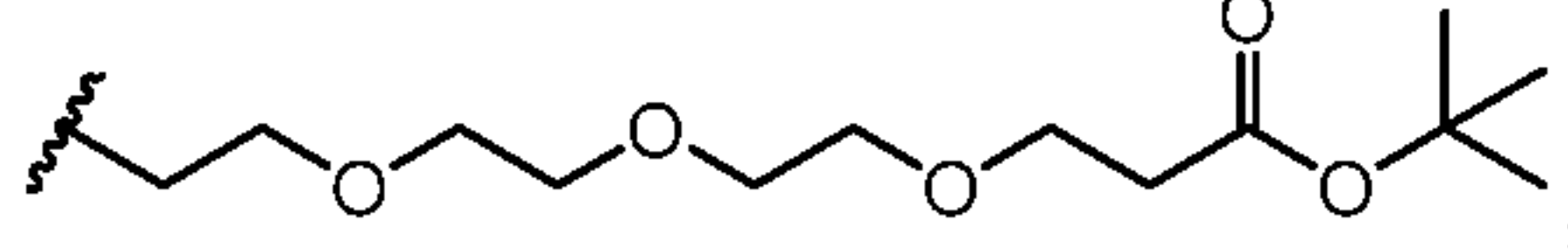
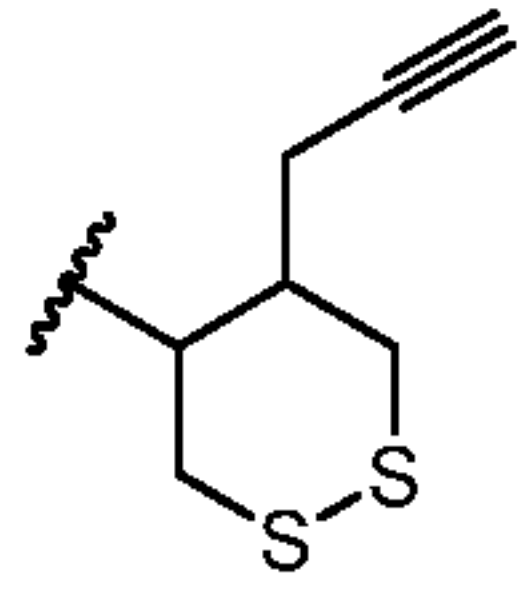
, where Base¹ and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂. In this aspect, all other groups are as provided in the formula (I) above.

[0110] In aspects of this embodiment, Y and Y^a are each independently selected from the group consisting of -O- and -S-. In this aspect, all other groups are as provided in the formula (I) above or in the aspect described above.

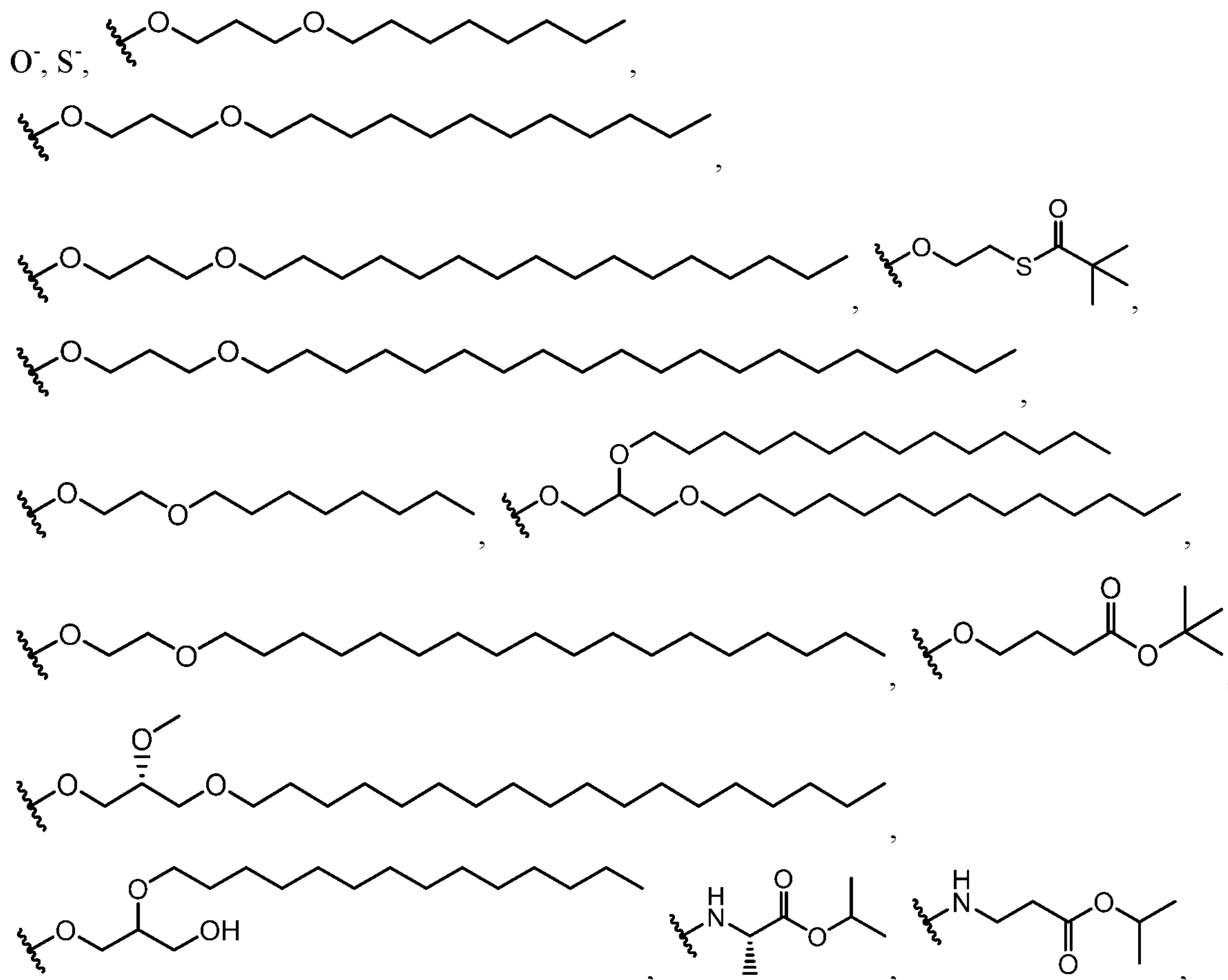
[0111] In aspects of this embodiment, X^a and X^{a1} are each independently selected from the group consisting of O and S. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

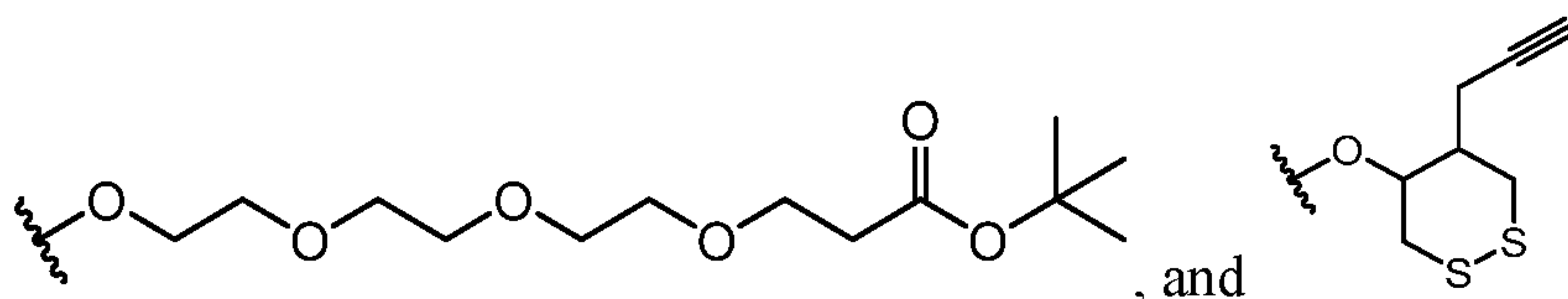
[0112] In aspects of this embodiment, X^b and X^{b1} are each independently selected from the group consisting of O and S. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0113] In aspects of this embodiment, X^c and X^{c1} are each independently selected from the group consisting of O^- , S^- , OR^9 , and NR^9R^9 , where each R^9 is independently selected from

the group consisting of H, C_1 - C_{20} alkyl, , and ,

where each R^9 C_1 - C_{20} alkyl is optionally substituted by 0 to 3 substituents independently selected from the group consisting of OH, $-O$ - C_1 - C_{20} alkyl, $-S$ - $C(O)C_1$ - C_6 alkyl, and $-C(O)OC_1$ - C_6 alkyl. In particular instances, X^c and X^{c1} are each independently selected from the group consisting of





, and . In all instances of this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0114] In aspects of this embodiment, X^d and X^{d1} are each independently selected from the group consisting of O and S. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0115] In aspects of this embodiment, R^1 and R^{1a} are each H. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0116] In aspects of this embodiment, R^2 and R^{2a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^2 and R^{2a} C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 . In particular instances, R^2 and R^{2a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, CN, N_3 , CF_3 , CH_3 , CH_2OH , and CH_2CH_3 . In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0117] In aspects of this embodiment, R^3 is selected from the group consisting H, F, Cl, I, Br, OH, N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^3 C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 . In particular instances, R^3 are each independently selected from the group consisting of H, F, Cl, I, Br, OH, CN, N_3 , CF_3 , CH_3 , CH_2OH , and CH_2CH_3 . In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0118] In aspects of this embodiment, R^4 and R^{4a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^4 and R^{4a} C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 . In particular instances, R^4 and R^{4a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, CN, N_3 , CF_3 , CH_3 , CH_2OH , and CH_2CH_3 . In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0119] In aspects of this embodiment, R^5 is selected from the group consisting of H, F, Cl, I, Br, OH, N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^5 C_1 - C_6 alkyl or C_1 - C_6 haloalkyl

are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃. In particular instances, R⁵ are each independently selected from the group consisting of H, F, Cl, I, Br, OH, CN, N₃, CF₃, CH₃, CH₂OH, and CH₂CH₃. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

5 [0120] In aspects of this embodiment, R⁶ and R^{6a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, C₁-C₆ alkyl, C₂-C₆ alkenyl, and C₂-C₆ alkynyl. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0121] In aspects of this embodiment, R⁷ and R^{7a} are each H. In this aspect, all other
10 groups are as provided in the formula (I) above or in the aspects described above.

[0122] In aspects of this embodiment, R⁸ and R^{8a} are each H. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0123] In aspects of this embodiment, R^{1a} and R³ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆
15 alkynylene, such that where R^{1a} and R³ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0124] In aspects of this embodiment, R^{2a} and R³ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆
20 alkynylene, such that where R^{2a} and R³ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0125] In aspects of this embodiment, R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, and -O-C₂-C₆ alkynylene, such that where R³ and R^{6a} are
25 connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

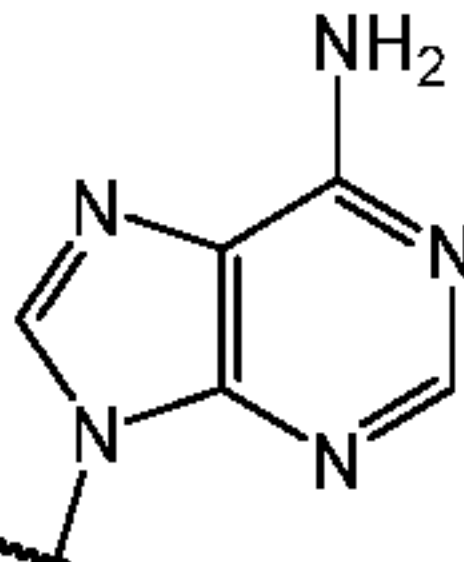
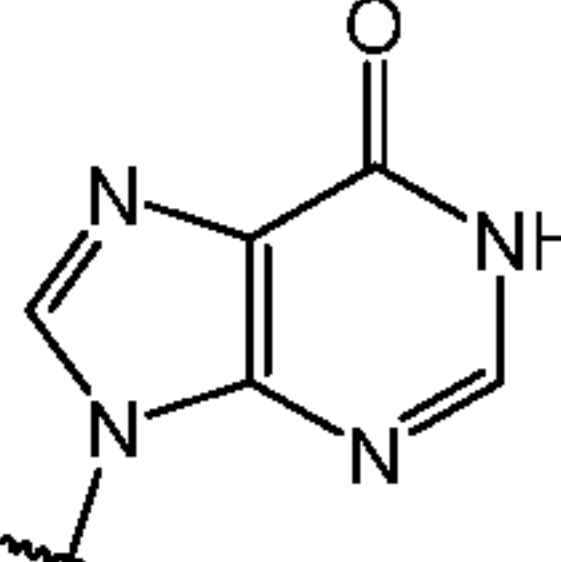
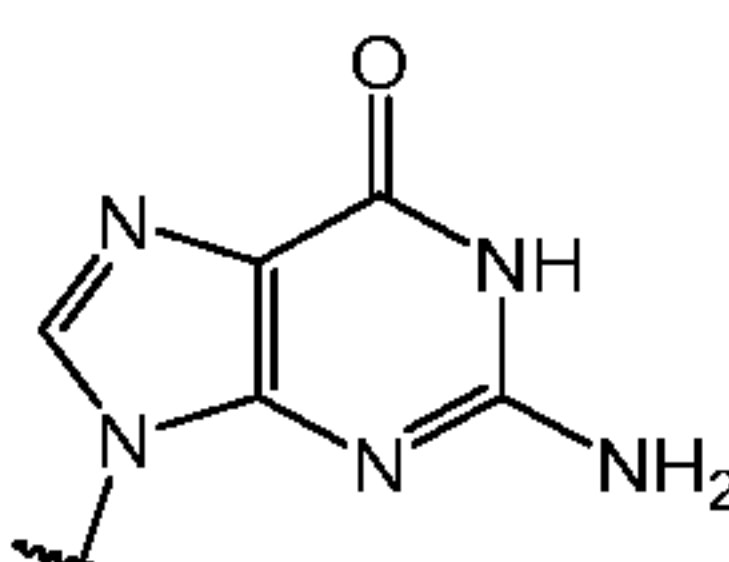
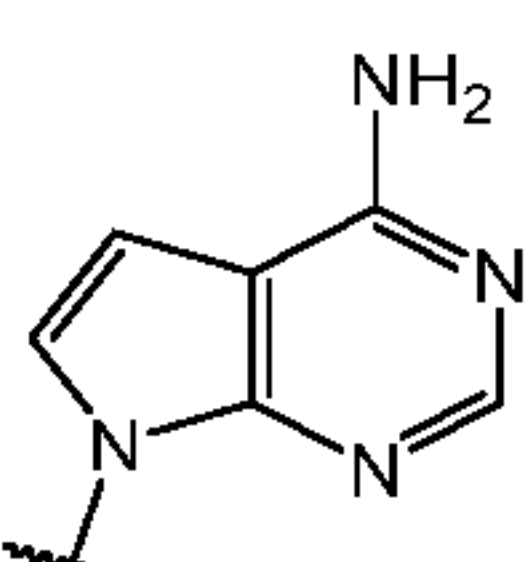
[0126] In aspects of this embodiment, R⁴ and R⁵ are connected by C₁-C₆ alkylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R⁴ and R⁵ are
30 connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R⁵ position. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

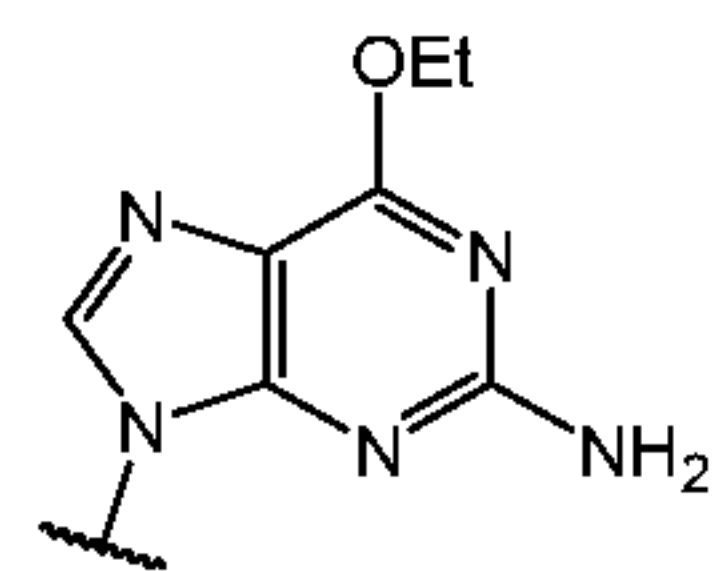
[0127] In aspects of this embodiment, R^5 and R^6 are connected to form $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, such that where R^5 and R^6 are connected to form $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, said O is bound at the R^5 position. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0128] In aspects of this embodiment, R^7 and R^8 are connected to form C_1-C_6 alkylene, C_2-C_6 alkenylene, or C_2-C_6 alkynylene. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0129] In aspects of this embodiment, R^{7a} and R^{8a} are connected to form C_1-C_6 alkylene, C_2-C_6 alkenylene, or C_2-C_6 alkynylene. In this aspect, all other groups are as provided in the formula (I) above or in the aspects described above.

[0130] In aspects of this embodiment, $Base^1$ and $Base^2$ are each independently selected

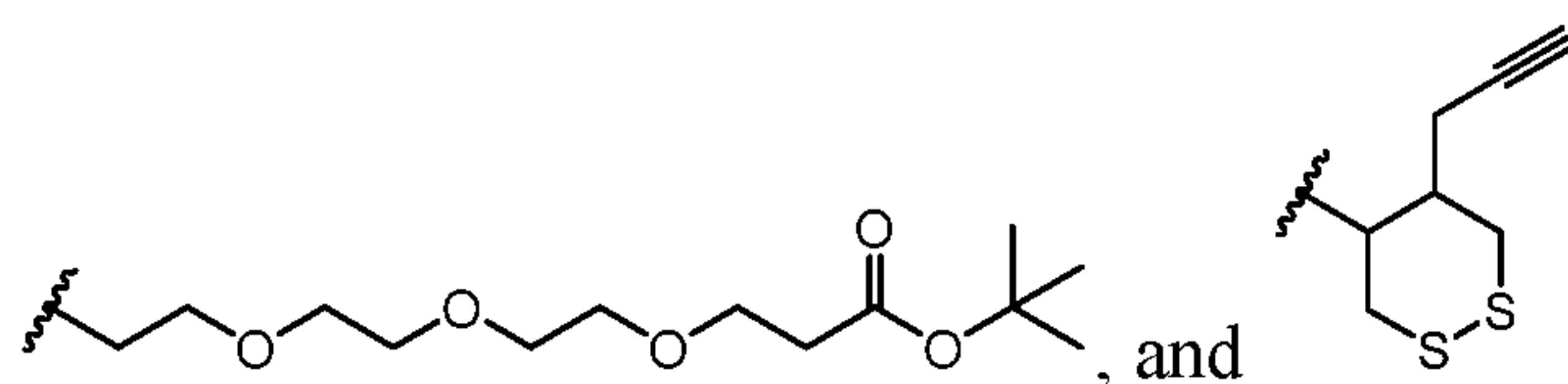
from the group consisting of , , , , and



, where $Base^1$ and $Base^2$ each may be independently substituted by 0-3 substituents

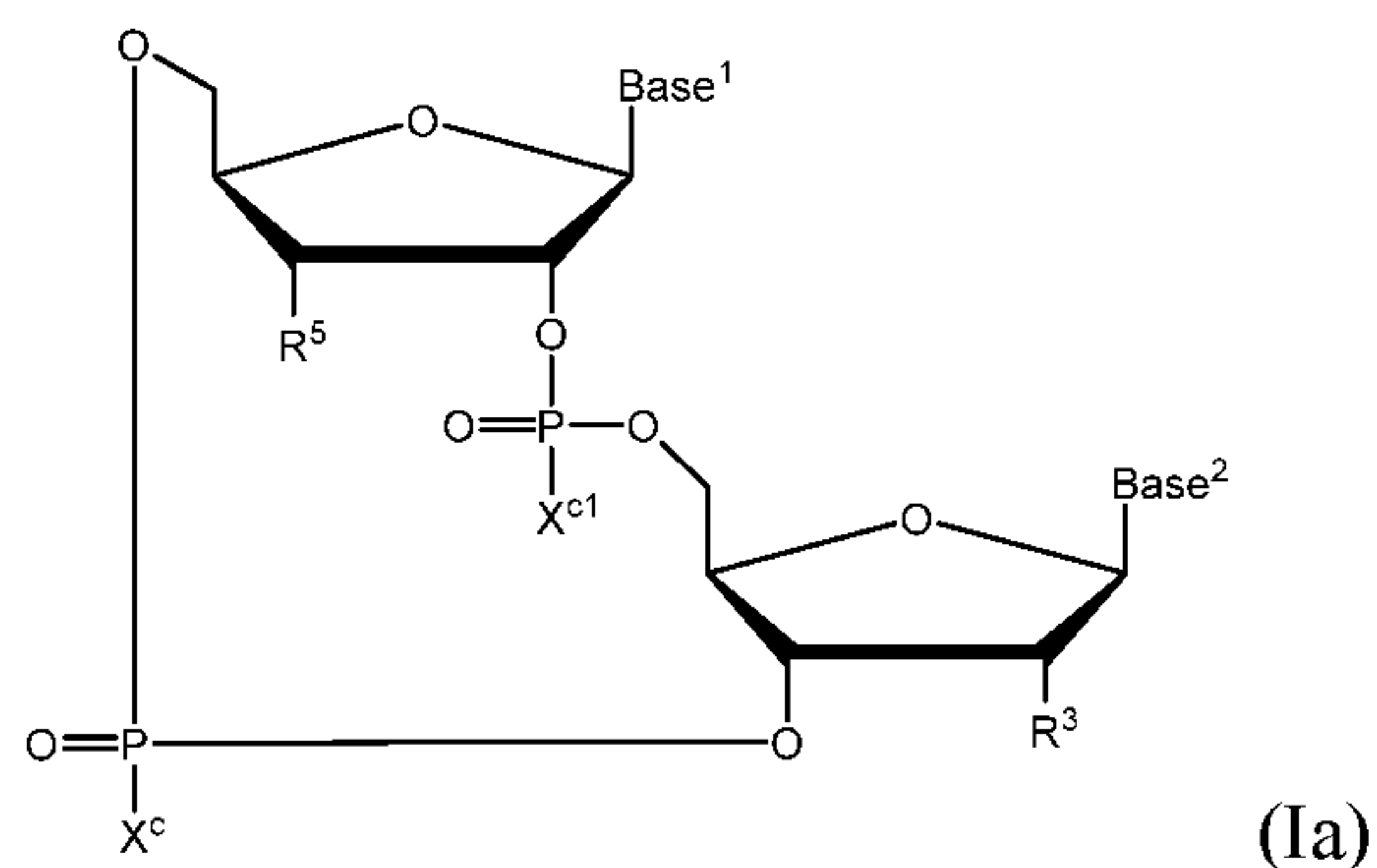
R^{10} , where each R^{10} is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH_2 , C_{1-3} alkyl, C_{3-6} cycloalkyl, $O(C_{1-3}$ alkyl), $O(C_{3-6}$ cycloalkyl), $S(C_{1-3}$ alkyl), $S(C_{3-6}$ cycloalkyl), $NH(C_{1-3}$ alkyl), $NH(C_{3-6}$ cycloalkyl), $N(C_{1-3}$ alkyl) $_2$, and $N(C_{3-6}$ cycloalkyl) $_2$; Y and Y^a are each independently selected from the group consisting of $-O-$, $-S-$, $-SO_2-$, $-CH_2-$, and $-CF_2-$; X^a and X^{a1} are each independently selected from the group consisting of O and S; X^b and X^{b1} are each independently selected from the group consisting of O and S; X^c and X^{c1} are each independently selected from the group consisting of O^- , S^- , OR^9 , and NR^9R^9 ; X^d and X^{d1} are each independently selected from the group consisting of O and S; R^1 and R^{1a} are each H; R^2 and R^{2a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, N_3 , C_1-C_6 alkyl, and C_1-C_6 haloalkyl, where said R^2 and R^{2a} C_1-C_6 alkyl or C_1-C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 ; R^3 is selected from the group consisting of H, F, Cl, I, Br, OH, CN, N_3 , C_1-C_6 alkyl, and C_1-C_6 haloalkyl, where said R^3 C_1-C_6 alkyl or C_1-C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 ; R^4 and R^{4a} are each independently selected

from the group consisting of H, C₁-C₆ alkyl, and C₁-C₆ haloalkyl, where said R⁴ and R^{4a} C₁-C₆ alkyl or C₁-C₆ haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁵ is selected from the group consisting of H, F, Cl, I, Br, OH, N₃, C₁-C₆ alkyl, and C₁-C₆ haloalkyl, where said R⁵ C₁-C₆ alkyl or C₁-C₆ haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁶ and R^{6a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, and C₁-C₆ haloalkyl, where said R⁶ and R^{6a} C₁-C₆ alkyl or C₁-C₆ haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁷ and R^{7a} are each H; R⁸ and R^{8a} are each H; each R⁹ is independently selected from the group consisting of H, C₂-C₃ alkyl,

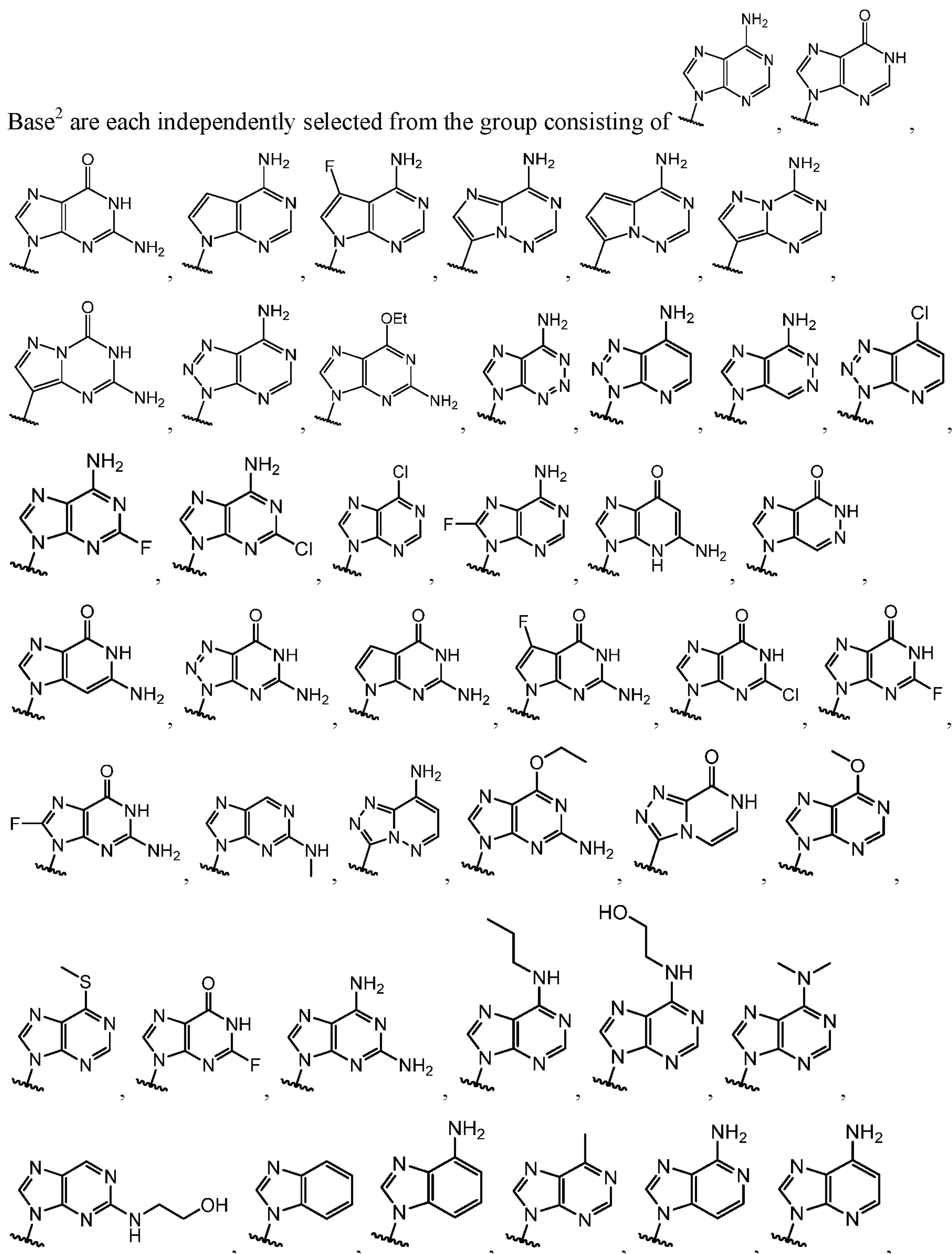


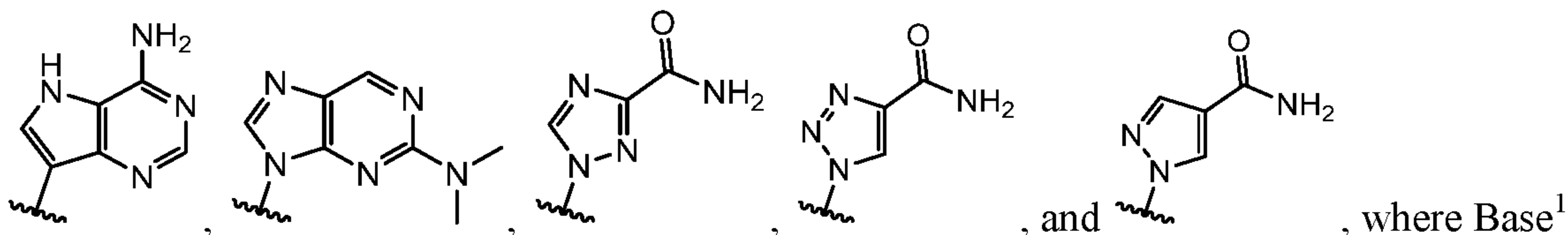
, where each R⁹ C₂-C₃ alkyl is optionally substituted by 1 to 2 substituents independently selected from the group consisting of OH, -O-C₁-C₂₀ alkyl, -S-C(O)C₁-C₆ alkyl, and -C(O)OC₁-C₆ alkyl; optionally R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, and -O-C₂-C₆ alkynylene, such that where R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position or optionally R⁴ and R⁵ are connected by C₁-C₆ alkylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R⁴ and R⁵ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R⁵ position. In this aspect, all other groups are as provided in the formula (I) above.

[0131] In aspects of this embodiment, the compound of formula (I) is a compound of formula (Ia):



or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein Base¹ and

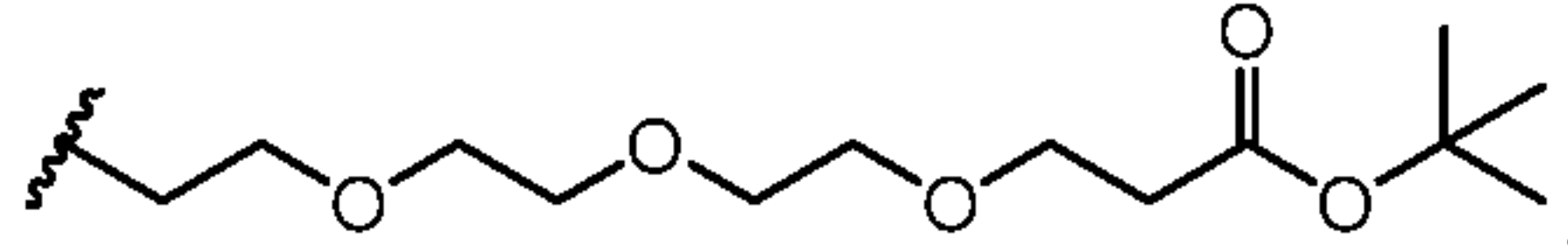
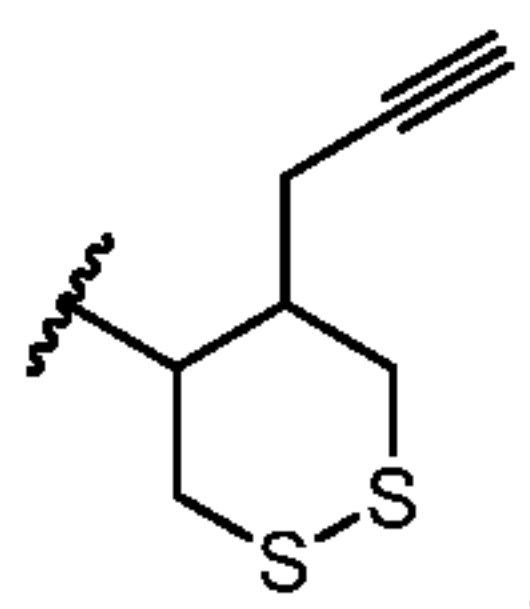




, where Base¹ and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is

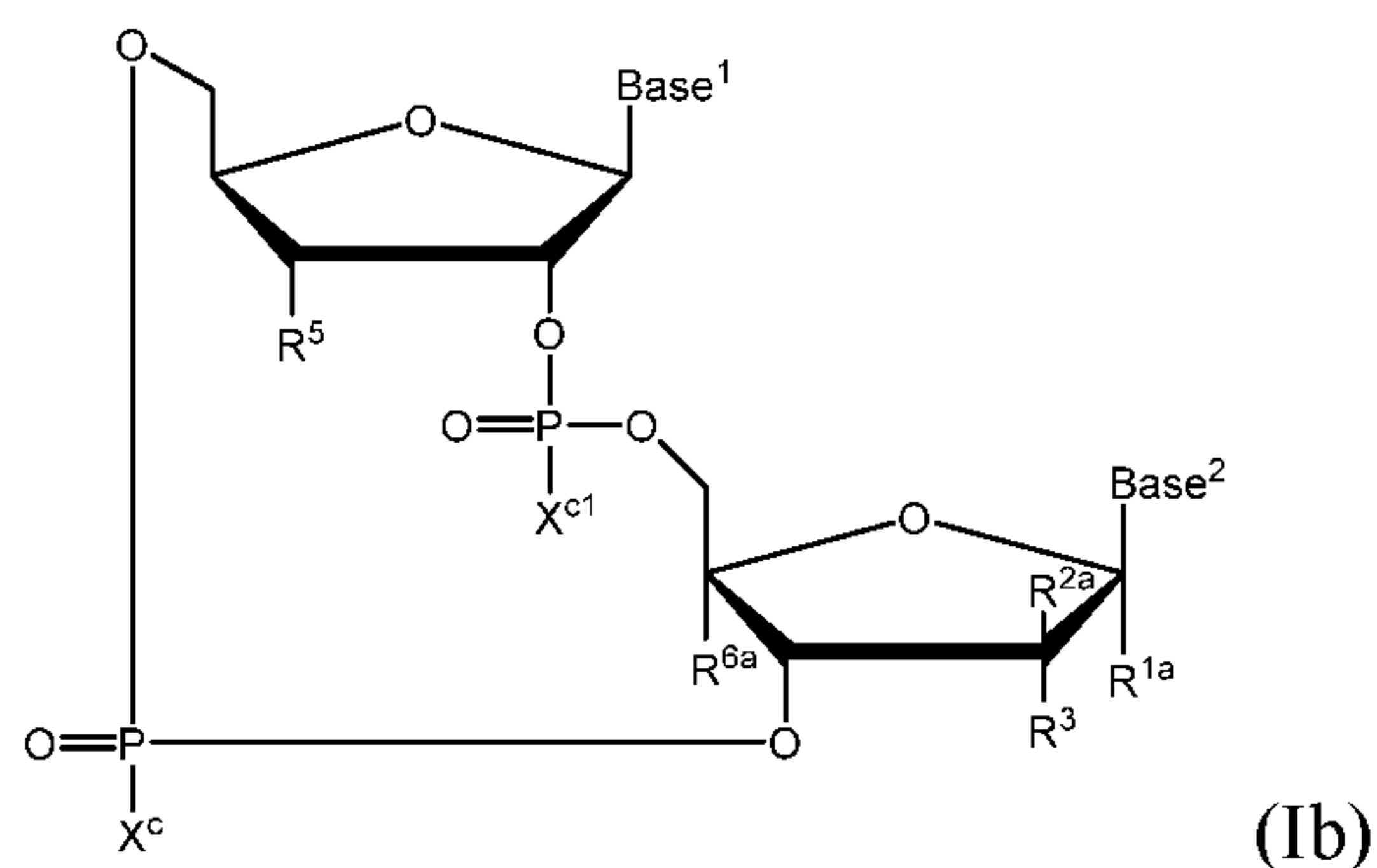
independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl),

5 NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂; X^c and X^{c1} are each independently selected from the group consisting of O⁻, S⁻, OR⁹, and NR⁹R⁹; R³ is selected from the group consisting of H, F, Cl, I, Br, OH, CN, N₃, C₁₋₆ alkyl, and C₁₋₆ haloalkyl, where said R³ C₁₋₆ alkyl or C₁₋₆ haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, I, Br, and OH; R⁵ is selected from the group consisting of H, F, Cl, I, Br, OH, CN, N₃,
10 C₁₋₆ alkyl, and C₁₋₆ haloalkyl, where said R⁵ C₁₋₆ alkyl or C₁₋₆ haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, I, Br, and OH; R³ and R⁵ are not both selected from the group consisting of OH, C₁₋₆ alkyl substituted with OH, and C₁₋₆ haloalkyl substituted with OH; and each R⁹ is independently selected from the group consisting

of H, C₂₋₃ alkyl, , and , where each R⁹ C₂₋₃ alkyl

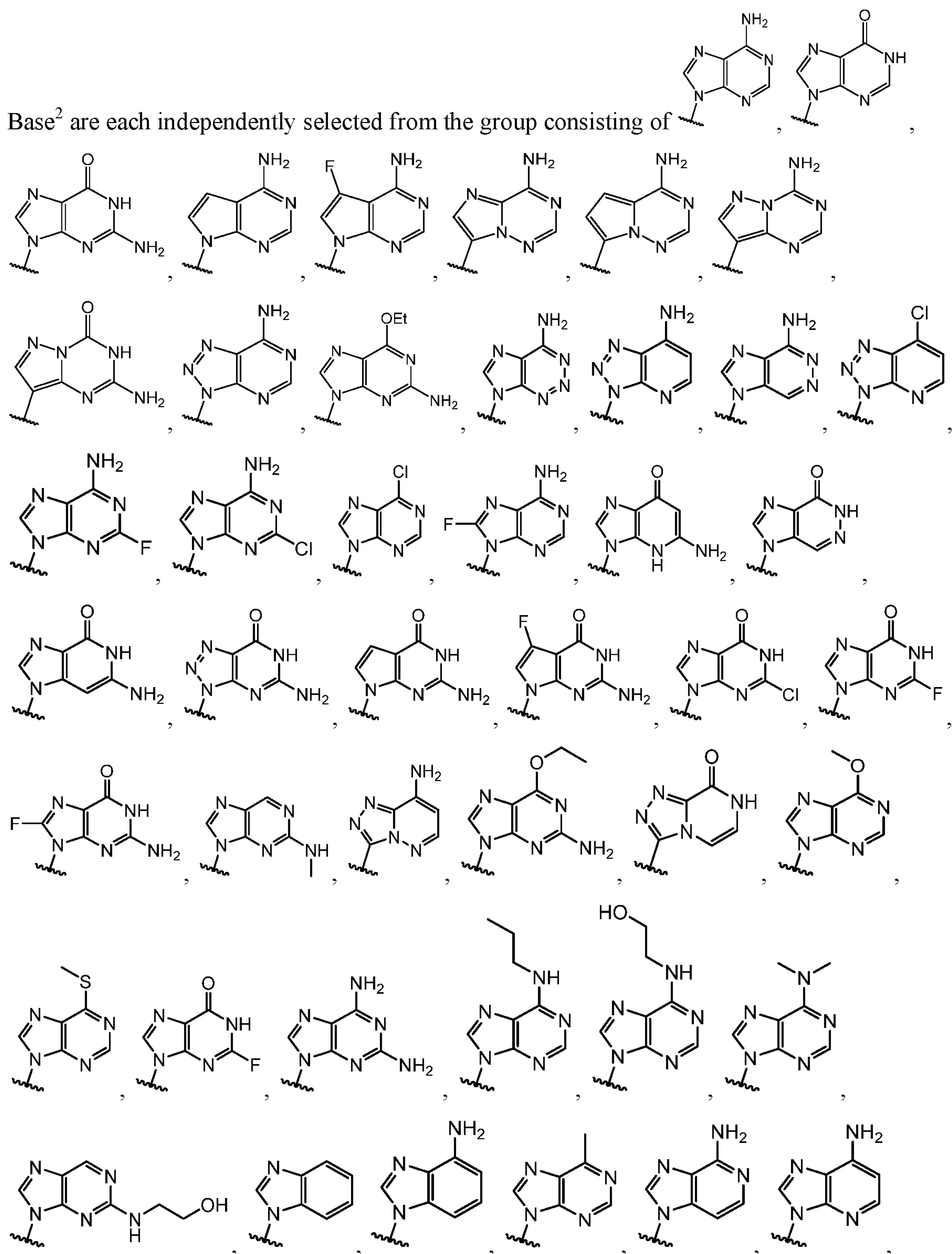
15 is optionally substituted by 1 to 2 substituents independently selected from the group consisting of OH, -O-C₁₋₂₀ alkyl, -S-C(O)C₁₋₆ alkyl, and -C(O)OC₁₋₆ alkyl. In this aspect, all other groups are as provided in the formula (I) above.

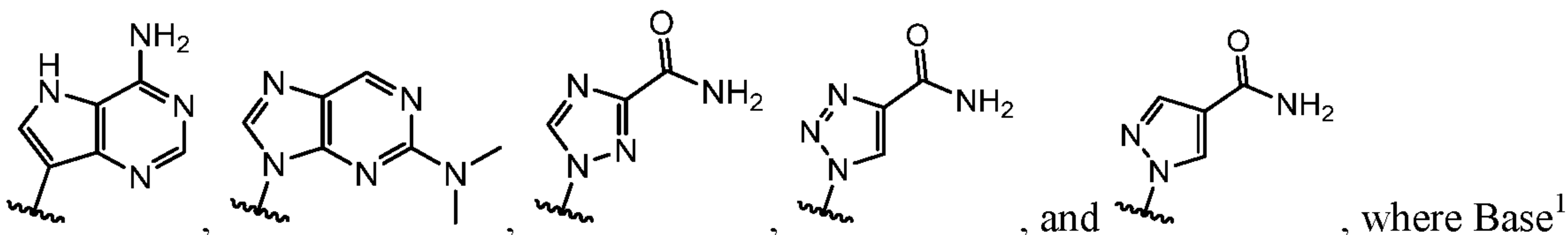
[0132] In aspects of this embodiment, the compound of formula (I) is a compound of formula (Ib):



(Ib)

or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein Base¹ and





, where Base¹ and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is

independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl),

5 NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂; X^c and X^{cl} are each independently selected from the group consisting of OR⁹, SR⁹, and NR⁹R⁹; R^{1a} is selected from the group

consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and

-O-C₂₋₆ alkynyl, where said R^{1a} C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl,

10 C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN,

and N₃; R^{2a} is selected from the group consisting of H, F, Cl, I, Br, OH, CN, N₃, C₁₋₆ alkyl, and C₁₋₆ haloalkyl, where said R^{2a} C₁₋₆ alkyl or C₁₋₆ haloalkyl are substituted by 0 to 3

substituents selected from the group consisting of F, Cl, I, Br, and OH; R³ is selected from the

15 group consisting of H, F, Cl, I, Br, OH, CN, N₃, C₁₋₆ alkyl, and C₁₋₆ haloalkyl, where said R³ C₁₋₆ alkyl or C₁₋₆ haloalkyl are substituted by 0 to 3 substituents selected from the group

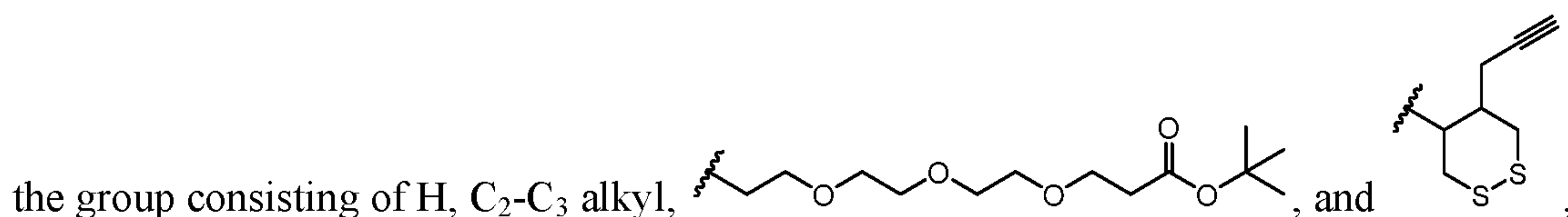
consisting of F, Cl, I, Br, and OH; R⁵ is selected from the group consisting of H, F, Cl, I, Br, OH, CN, N₃, C₁₋₆ alkyl, and C₁₋₆ haloalkyl, where said R⁵ C₁₋₆ alkyl or C₁₋₆ haloalkyl are

substituted by 0 to 3 substituents selected from the group consisting of F, Cl, I, Br, and OH; R³

20 and R⁵ are not both selected from the group consisting of OH, C₁₋₆ alkyl substituted with OH,

and C₁₋₆ haloalkyl substituted with OH; R^{6a} is selected from the group consisting of H, F, Cl, I,

Br, OH, C₁₋₆ alkyl, C₂₋₆ alkenyl, and C₂₋₆ alkynyl; each R⁹ is independently selected from



the group consisting of H, C₂₋₃ alkyl, where each R⁹ C₂₋₃ alkyl is optionally substituted by 1 to 2 substituents independently selected

25 from the group consisting of OH, -O-C₁₋₂₀ alkyl, -S-C(O)C₁₋₆ alkyl, and -C(O)OC₁₋₆ alkyl;

and optionally R³ and R^{6a} are connected to form -O-C₁₋₆ alkylene, -O-C₂₋₆ alkenylene, and

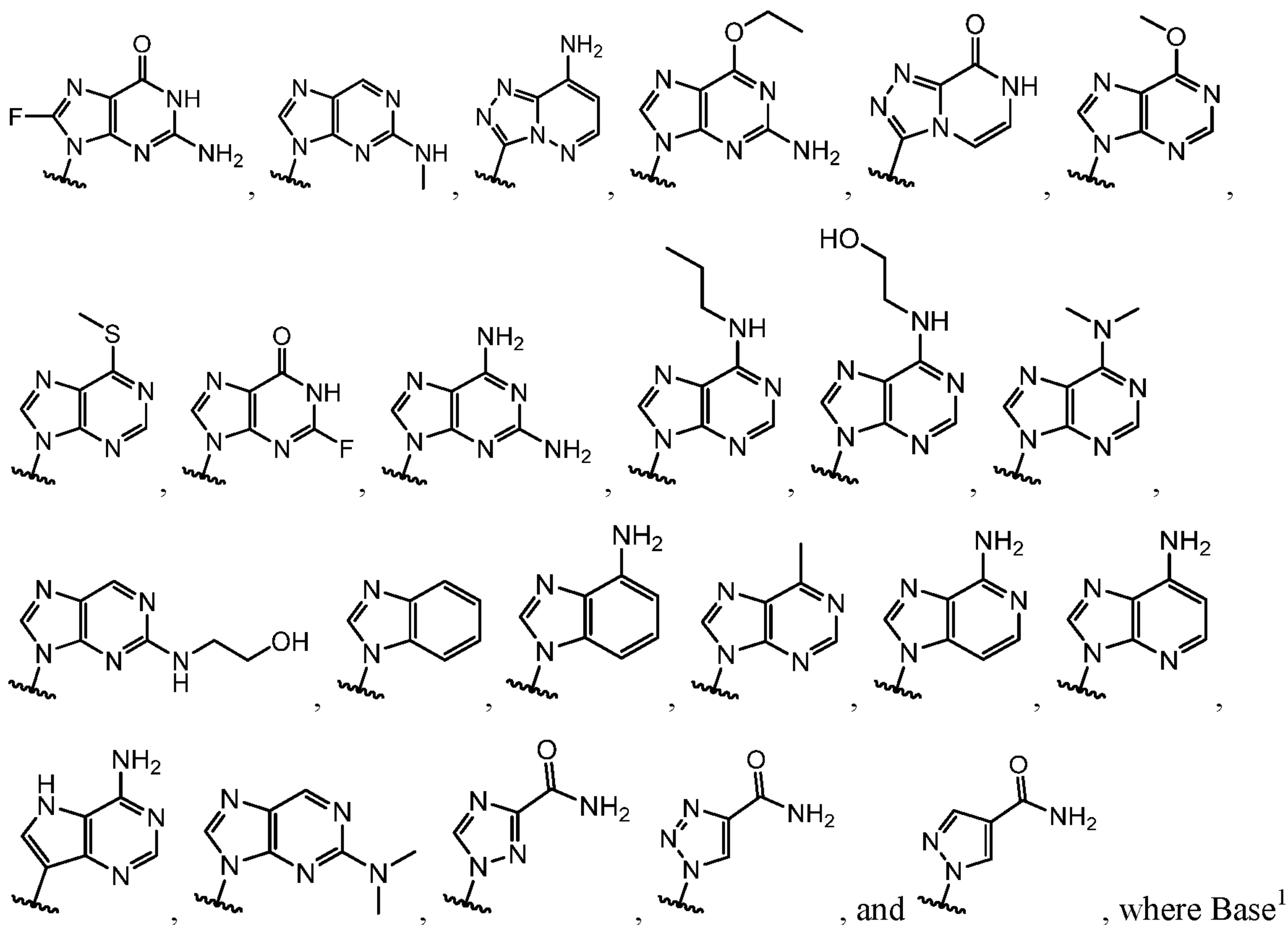
-O-C₂₋₆ alkynylene, such that where R³ and R^{6a} are connected to form -O-C₁₋₆ alkylene,

[0133] In aspects of this embodiment, the compound of formula (I) is a compound of formula (Ic):



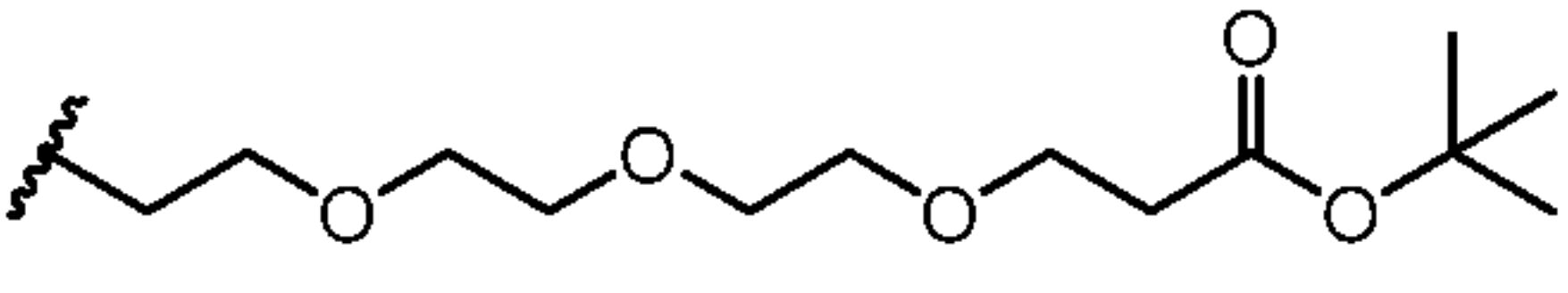
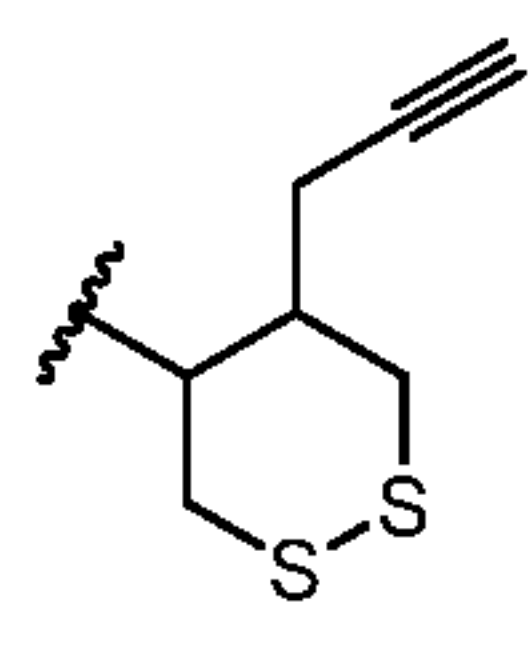
Base² are each independently selected from the group consisting of  ,  ,



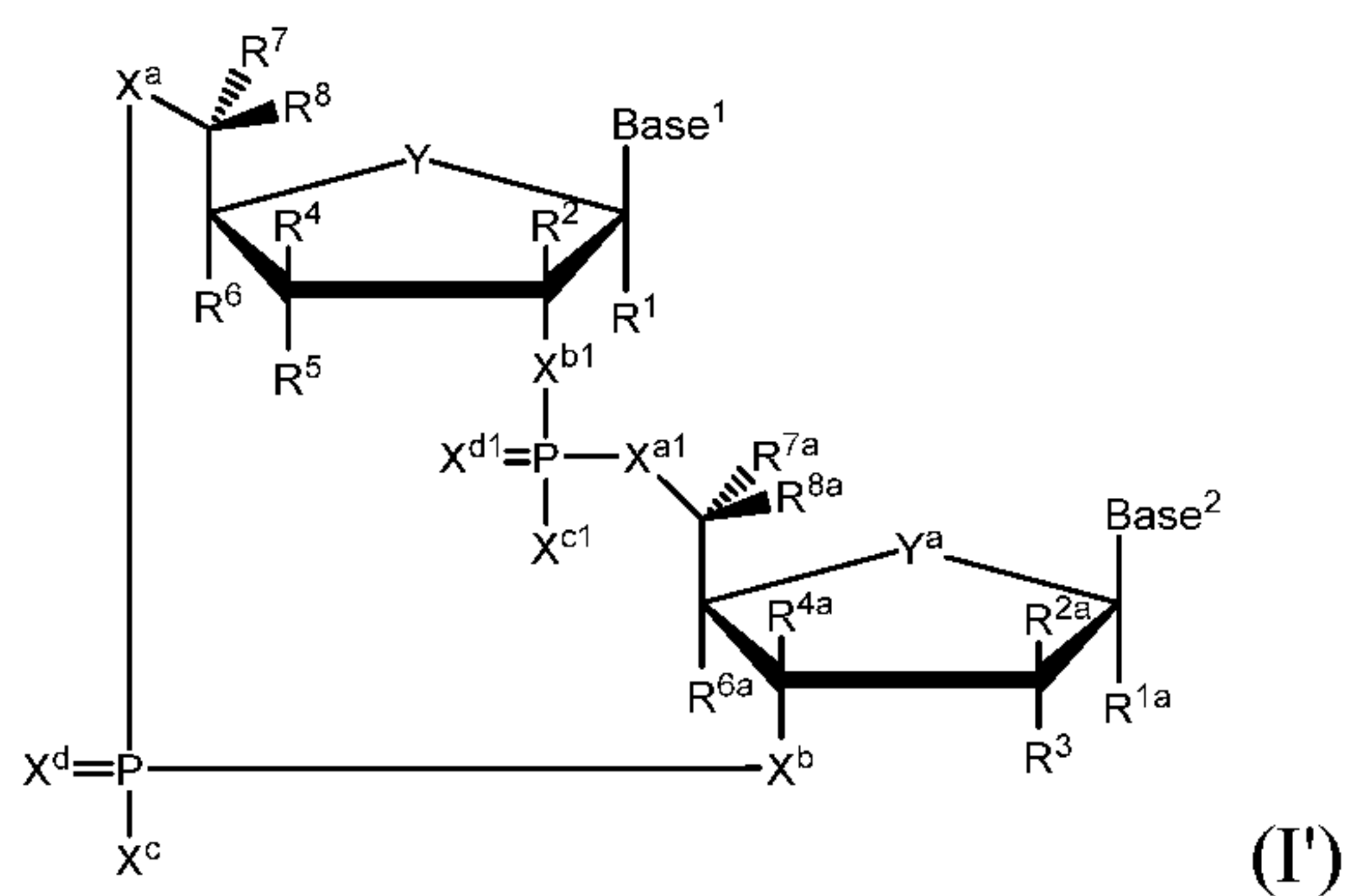


- 5 and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂; X^c and X^{cl} are each independently selected from the group consisting of OR⁹, SR⁹, and NR⁹R⁹; R³ is selected from the group
- 10 consisting of H, F, Cl, I, Br, OH, CN, N₃, C_{1-C6} alkyl, and C_{1-C6} haloalkyl, where said R³ C_{1-C6} alkyl or C_{1-C6} haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, I, Br, and OH; R⁴ is selected from the group consisting of H, F, OH, CN, N₃, C_{1-C6} alkyl, and C_{1-C6} haloalkyl, where said R⁴ C_{1-C6} alkyl or C_{1-C6} haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, I, Br, and OH; R⁵ is selected from the
- 15 group consisting of H, F, Cl, I, Br, OH, CN, N₃, C_{1-C6} alkyl, and C_{1-C6} haloalkyl, where said R⁵ C_{1-C6} alkyl or C_{1-C6} haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, I, Br, and OH; R³ and R⁵ are not both selected from the group consisting of OH, C_{1-C6} alkyl substituted with OH, and C_{1-C6} haloalkyl substituted with OH; R^{6a} is selected from the group consisting of H, F, Cl, I, Br, OH, CN, N₃, C_{1-C6} alkyl, and C_{1-C6} haloalkyl,
- 20 where said R^{6a} C_{1-C6} alkyl or C_{1-C6} haloalkyl are substituted by 0 to 3 substituents selected from

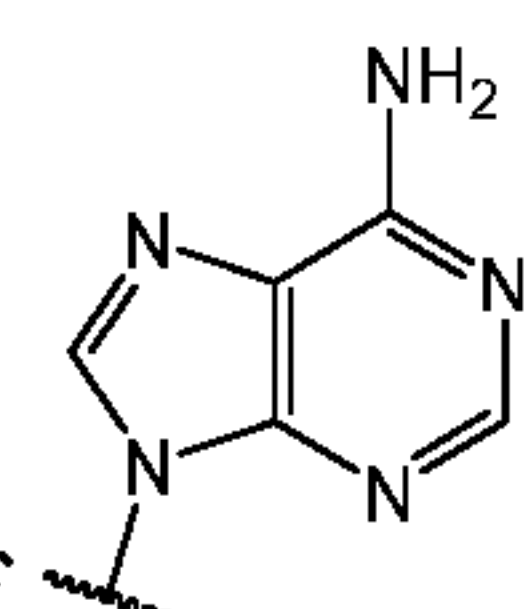
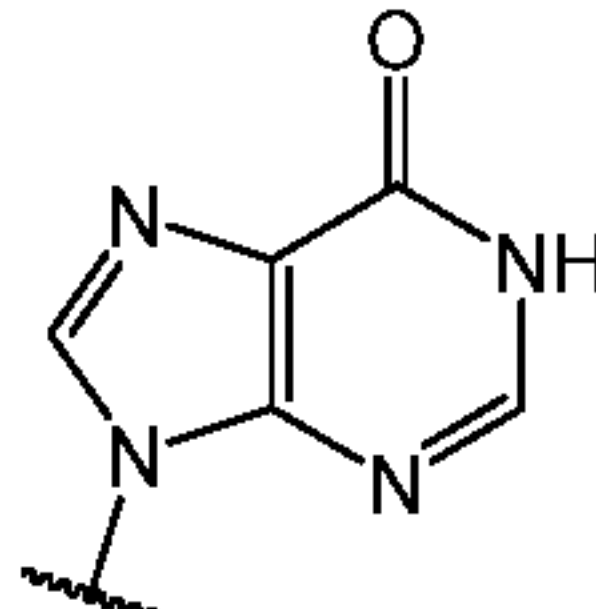
the group consisting of F, Cl, I, Br, and OH; each R^9 is independently selected from the group

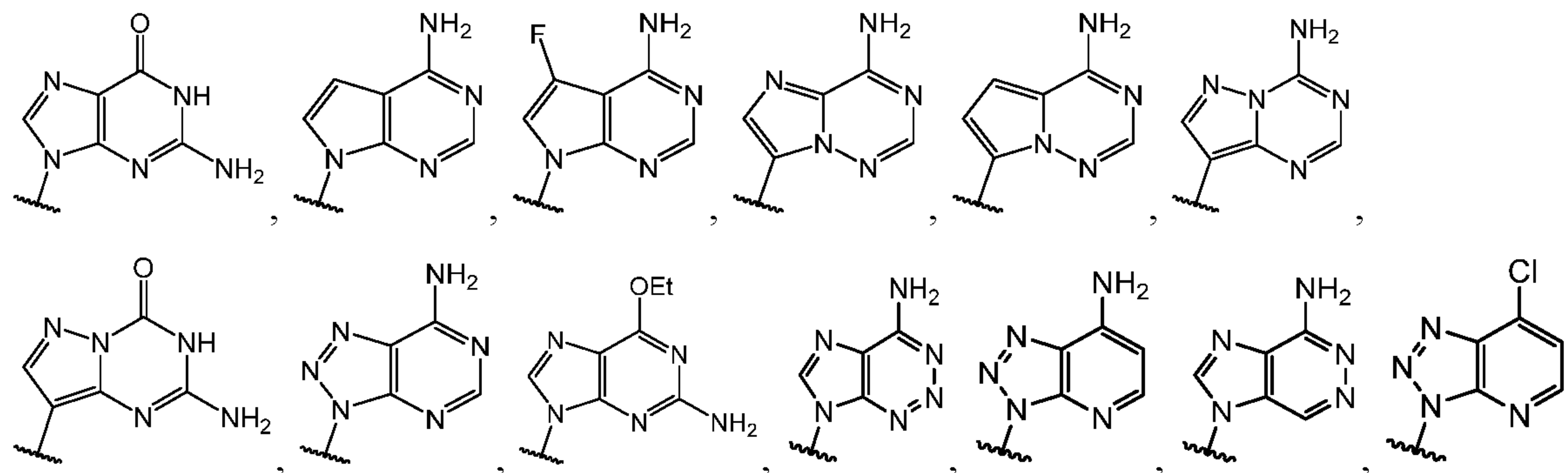
consisting of H, C₂-C₃ alkyl, , and , where each R^9 C₂-C₃ alkyl is optionally substituted by 1 to 2 substituents independently selected from the group consisting of OH, -O-C₁-C₂₀ alkyl, -S-C(O)C₁-C₆ alkyl, and -C(O)OC₁-C₆ alkyl; and optionally
 5 R^4 and R^5 are connected by C₁-C₆ alkylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R^4 and R^5 are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R^5 position. In this aspect, all other groups are as provided in the formula (I) above.

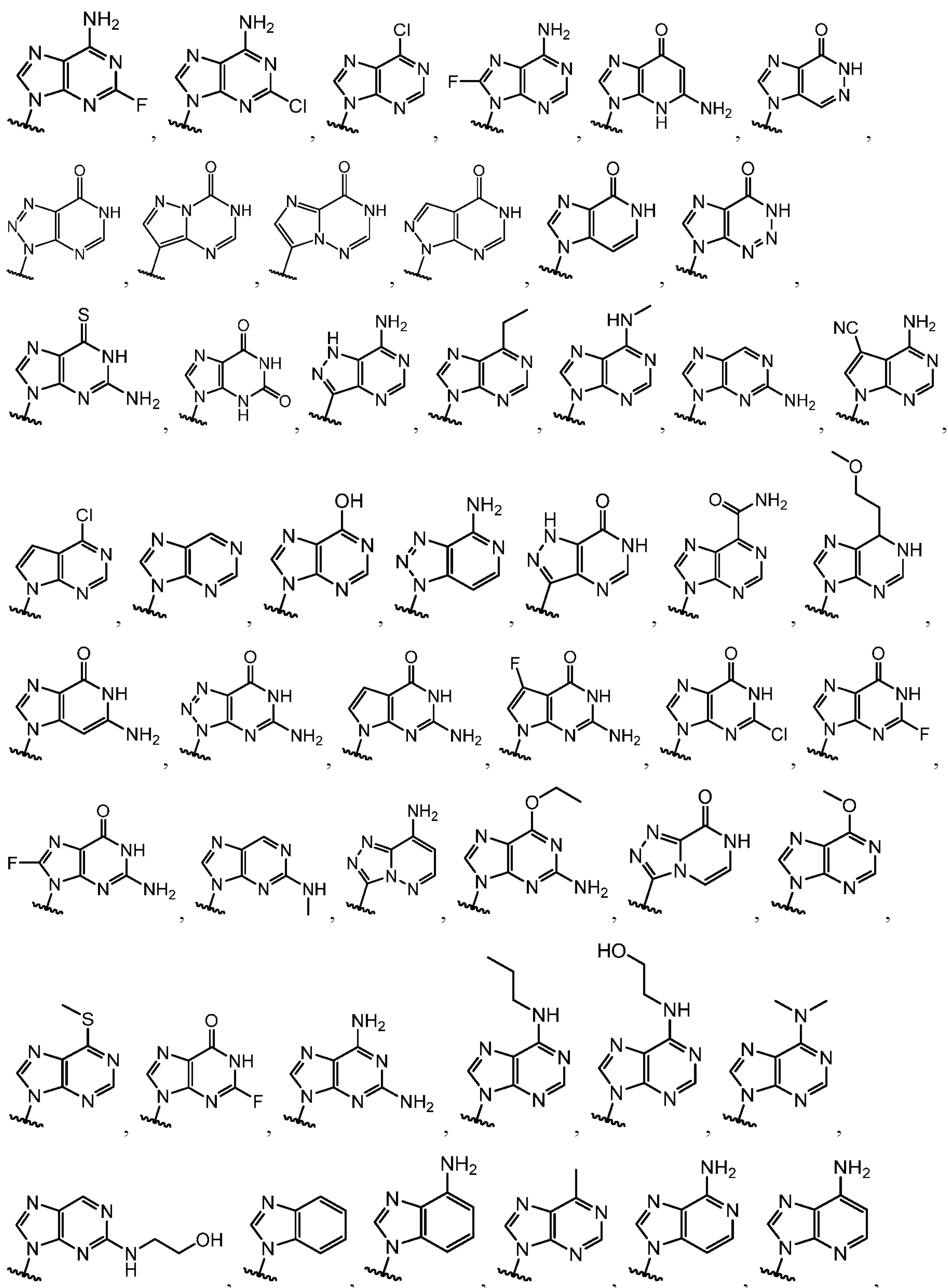
[0134] In an additional embodiment, the CDN STING agonist is selected from cyclic di-
 10 nucleotide compounds of formula (I'):

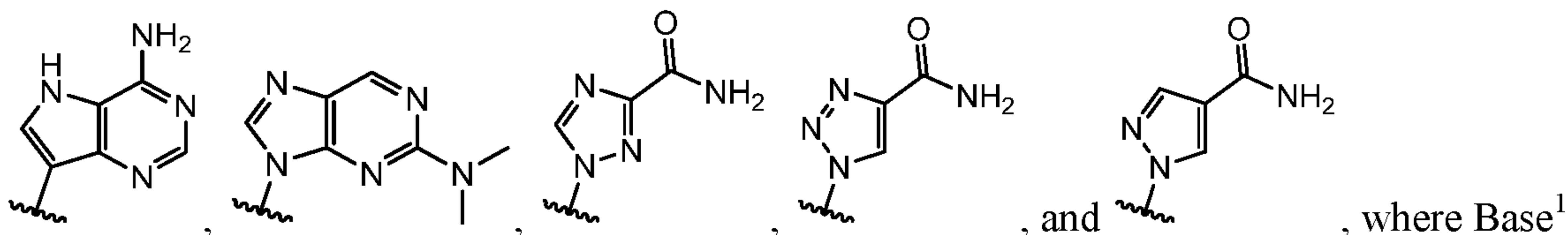


or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein Base¹ and

Base² are each independently selected from the group consisting of , ,

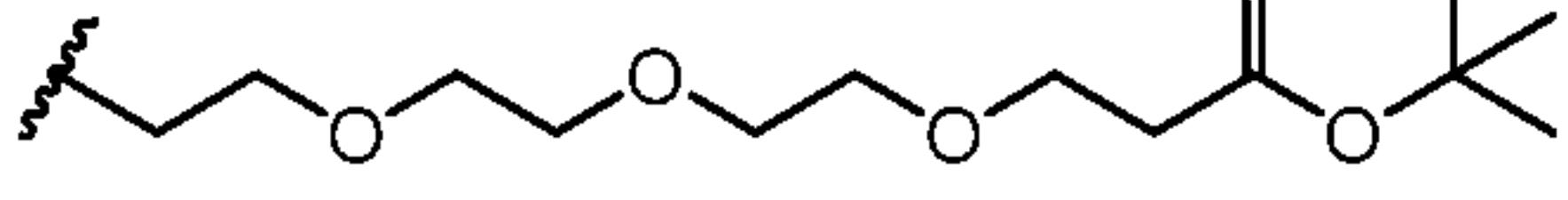
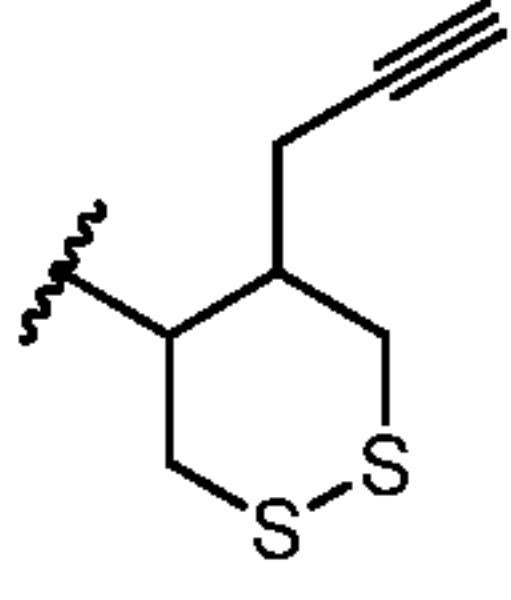






and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂; Y and Y^a are each independently selected from the group consisting of -O- and -S-; X^a and X^{a1} are each independently selected from the group consisting of O, and S; X^b and X^{b1} are each independently selected from the group consisting of O, and S; X^c and X^{c1} are each independently selected from the group consisting of OR⁹, SR⁹, and NR⁹R⁹; X^d and X^{d1} are each independently selected from the group consisting of O and S; R¹ and R^{1a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl, where said R¹ and R^{1a} C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R² and R^{2a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl, where said R² and R^{2a} C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R³ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl, where said R³ C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁴ and R^{4a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl, where said R⁴ and

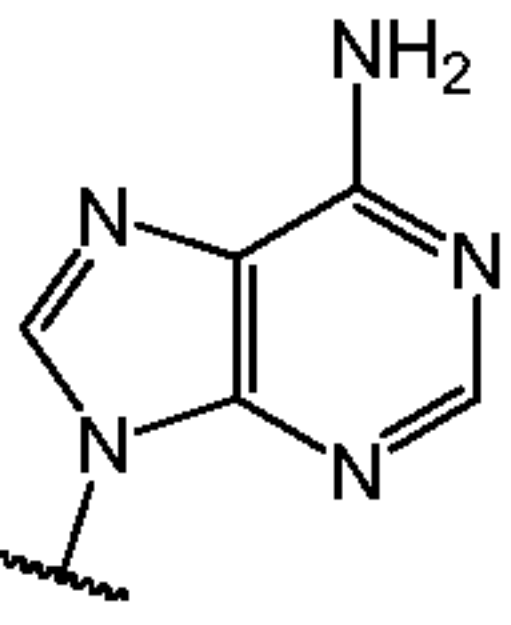
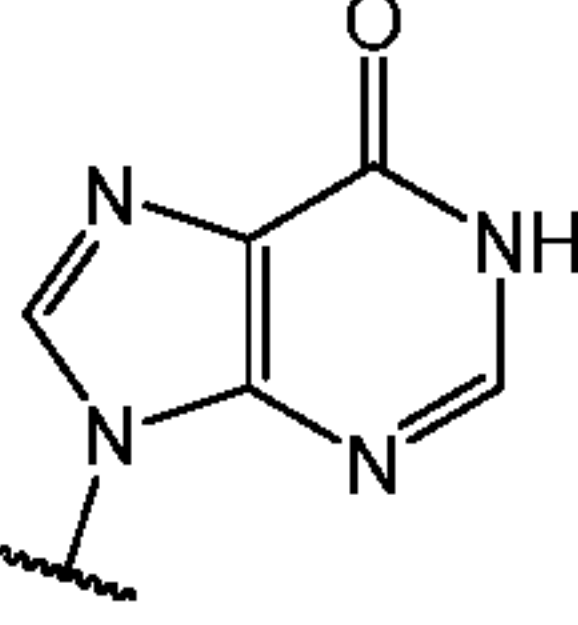
R^{4a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁵ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, NH₂, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁵ C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, NR⁹R⁹, and N₃; R⁶ and R^{6a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁶ and R^{6a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁷ and R^{7a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁷ and R^{7a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁸ and R^{8a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁸ and R^{8a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; each R⁹ is independently selected

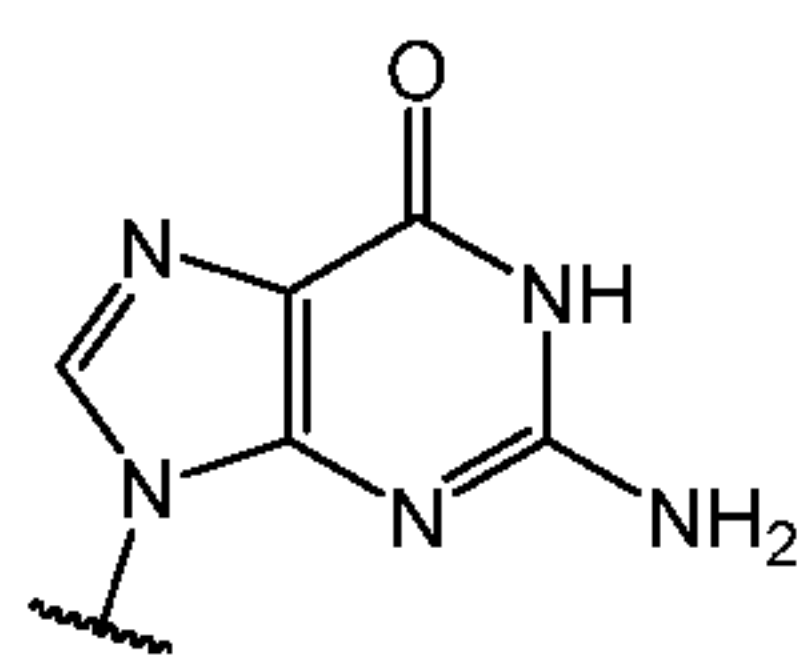
from the group consisting of H, C₁-C₂₀ alkyl, , and ,

where each R⁹ C₁-C₂₀ alkyl is optionally substituted by 0 to 3 substituents independently selected from the group consisting of OH, -O-C₁-C₂₀ alkyl, -S-C(O)C₁-C₆ alkyl, and -C(O)OC₁-C₆ alkyl; optionally R^{1a} and R³ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene,

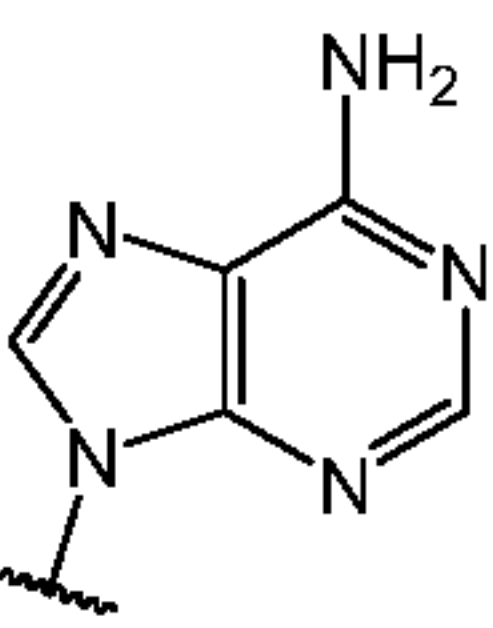
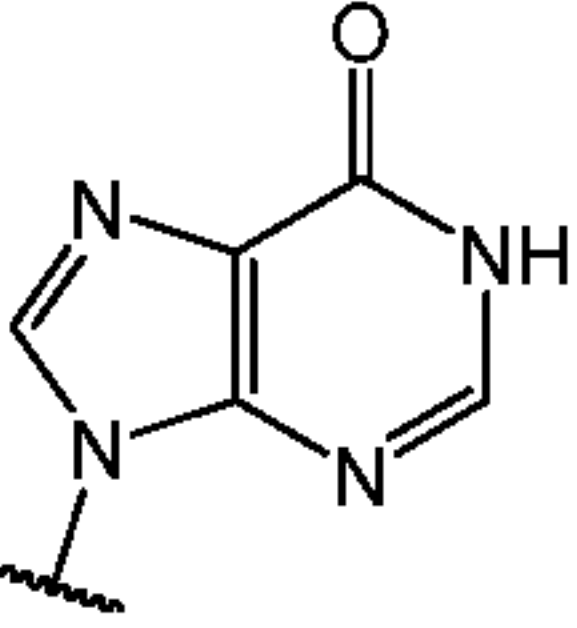
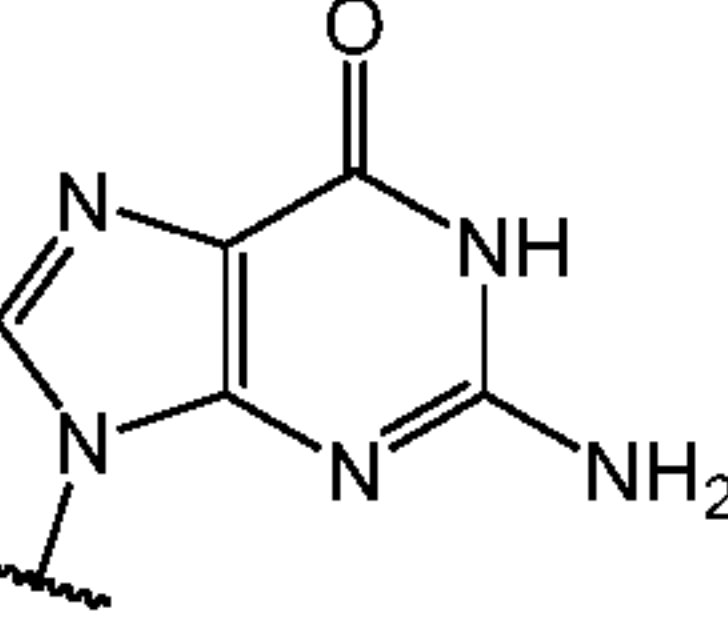
-O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R^{1a} and R³ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position; optionally R^{2a} and R³ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R^{2a} and R³ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position; optionally R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position; optionally R⁴ and R⁵ are connected to form are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R⁴ and R⁵ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R⁵ position; optionally R⁵ and R⁶ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R⁵ and R⁶ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R⁵ position; optionally R⁷ and R⁸ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, or C₂-C₆ alkynylene; and optionally R^{7a} and R^{8a} are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, or C₂-C₆ alkynylene.

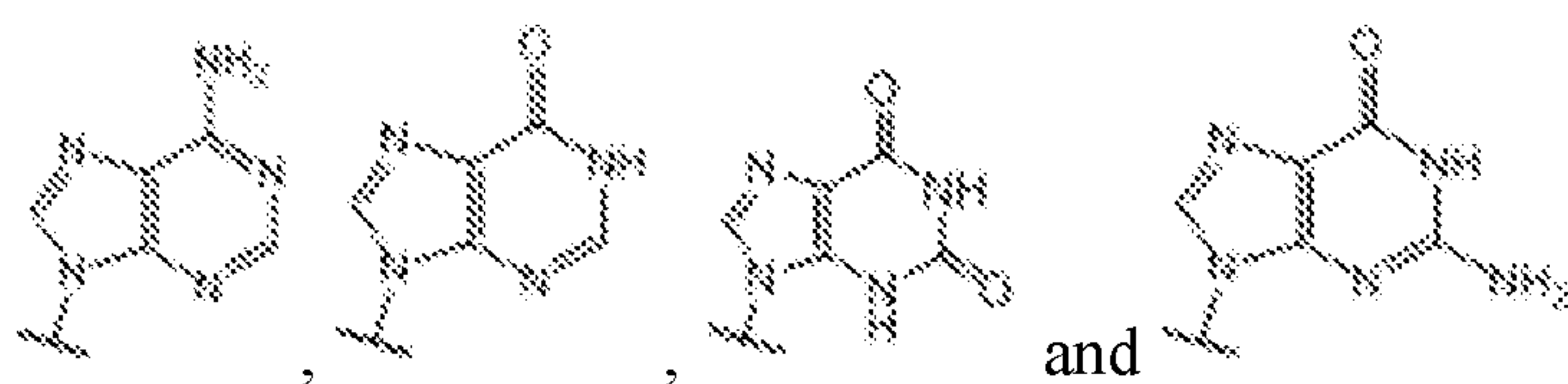
[0135] In specific aspects of this embodiment, when Y and Y^a are each O, X^a and X^{a1} are each O, X^b and X^{b1} are each O, and X^c and X^{c1} are each OH or SH, X^d and X^{d1} are each O, R¹ and R^{1a} are each H, R² is H, R⁶ and R^{6a} are each H, R⁷ and R^{7a} are each H, R⁸ and R^{8a} are each H,

and Base¹ and Base² are each selected from the group consisting of , , and



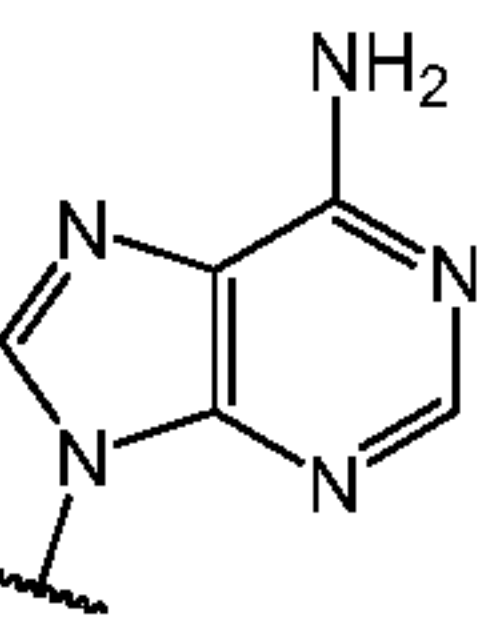
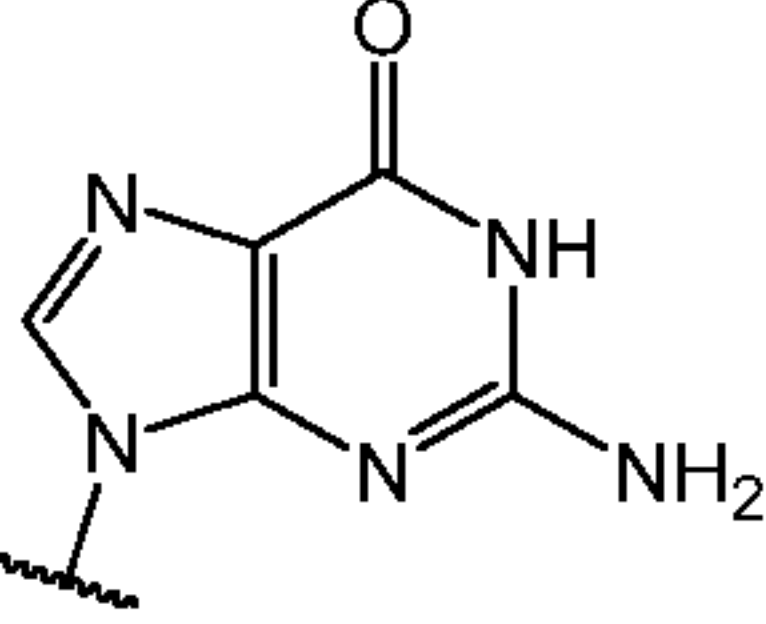
, R⁵ and R³ are not both selected from the group consisting of H, F and OH. That is, when Y and Y^a are each O, X^a and X^{a1} are each O, X^b and X^{b1} are each O, and X^c and X^{c1} are each OH or SH, X^d and X^{d1} are each O, R¹ and R^{1a} are each H, R² is H, R⁶ and R^{6a} are each H, R⁷ and R^{7a} are each H, R⁸ and R^{8a} are each H, and Base¹ and Base² are each selected from the group

consisting of , , and , either only one of R⁵ and R³ is selected from the group consisting of H, F, and OH, or neither R⁵ and R³ is selected from the group consisting of H, F, and OH. In specific instances of this aspect, when Y and Y^a are each O, X^a and X^{a1} are each O, X^b and X^{b1} are each O, and X^c and X^{c1} are each OH, X^d and X^{d1} are each O or S, R¹ and R^{1a} are each H, R² is H, R⁶ and R^{6a} are each H, R⁷ and R^{7a} are each H, R⁸ and R^{8a} are each H, and Base¹ and Base² are each selected from the group consisting of

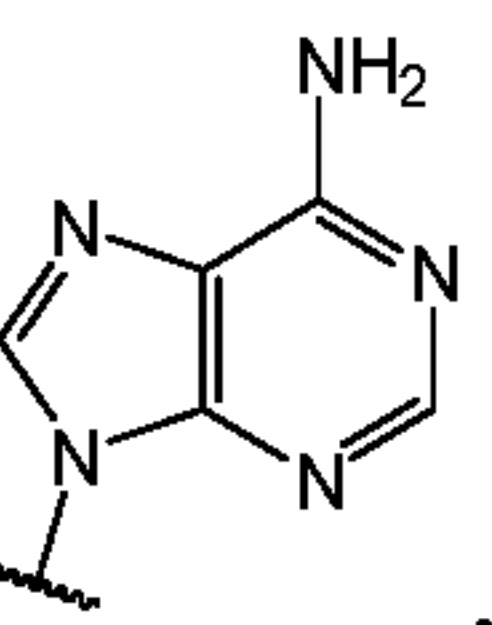
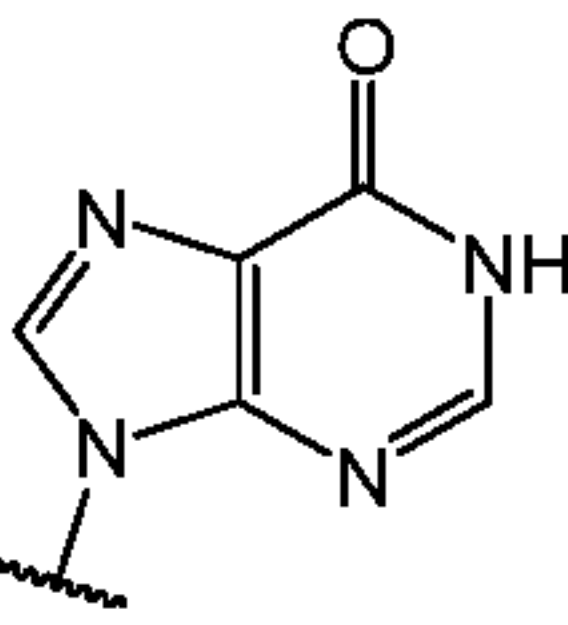
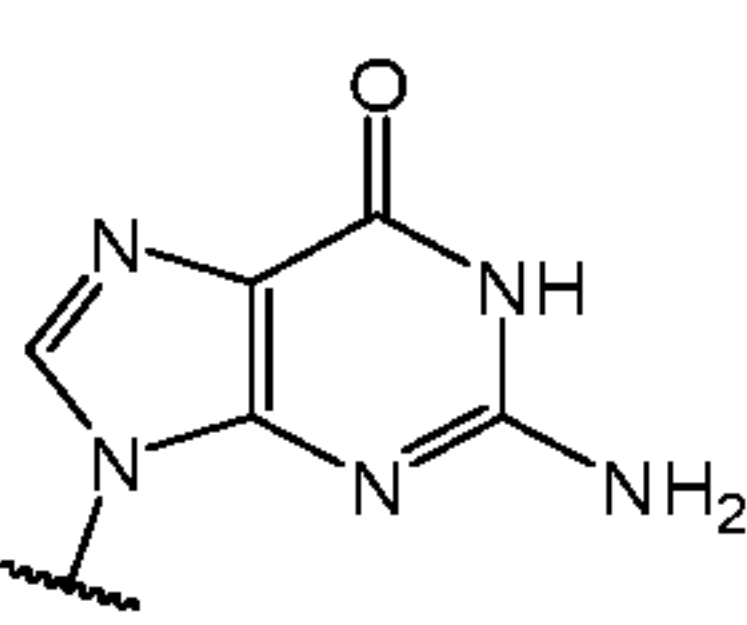
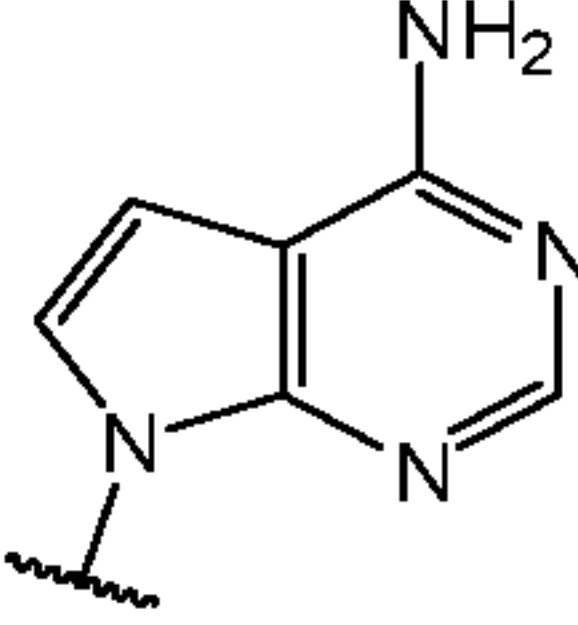
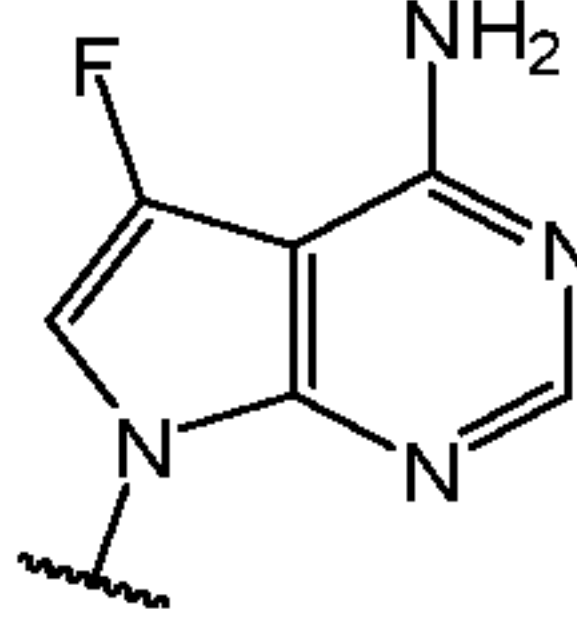
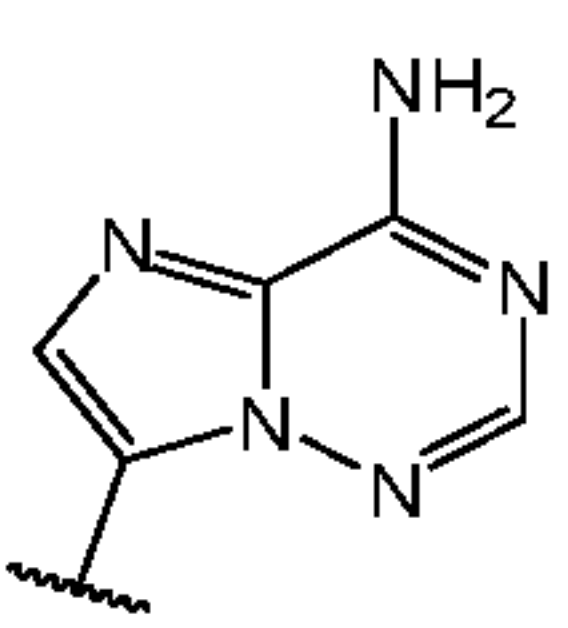
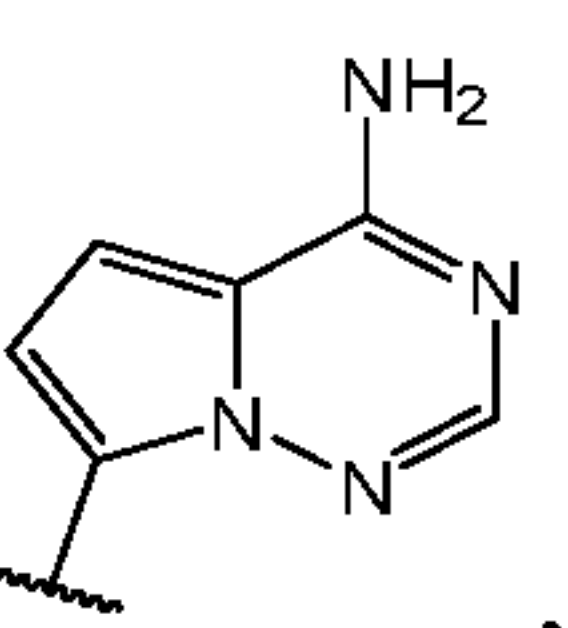
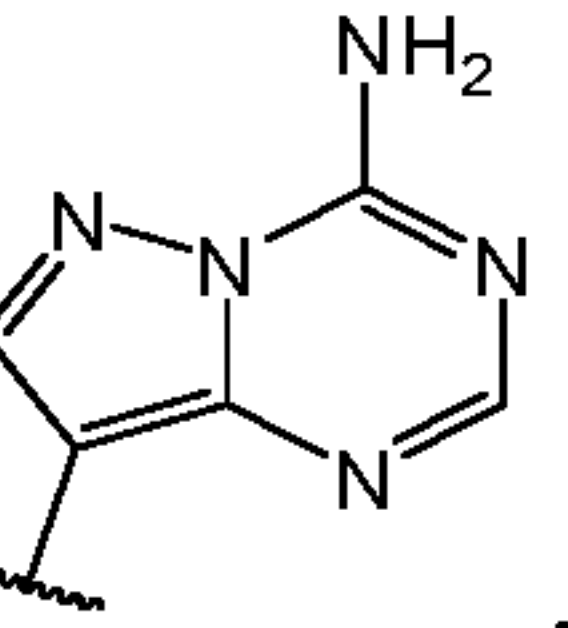
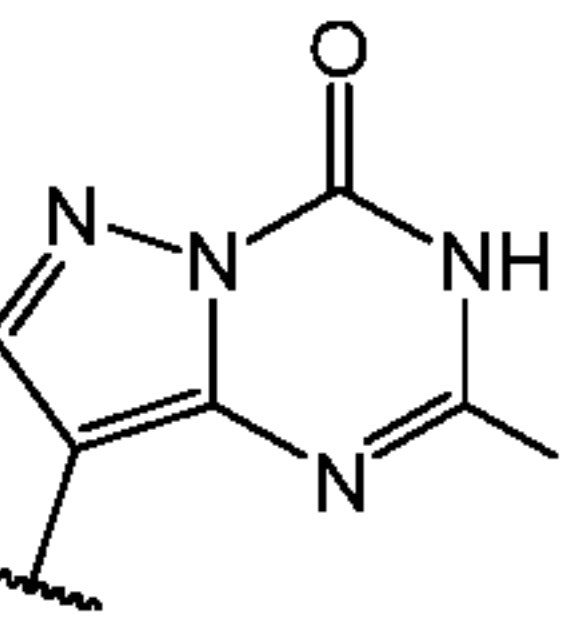
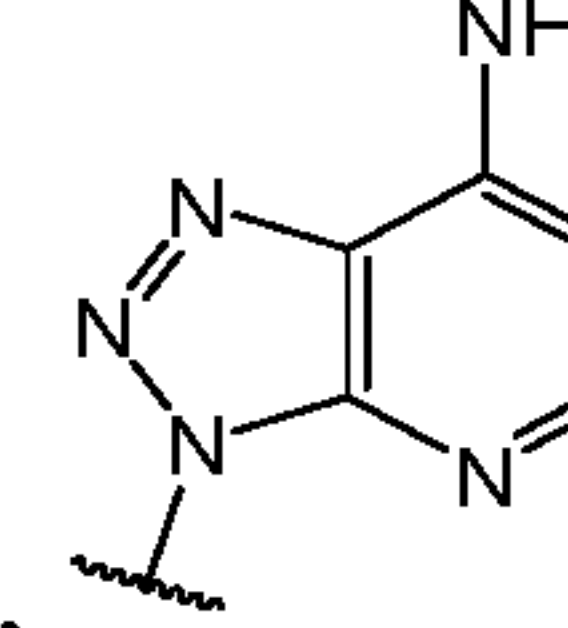
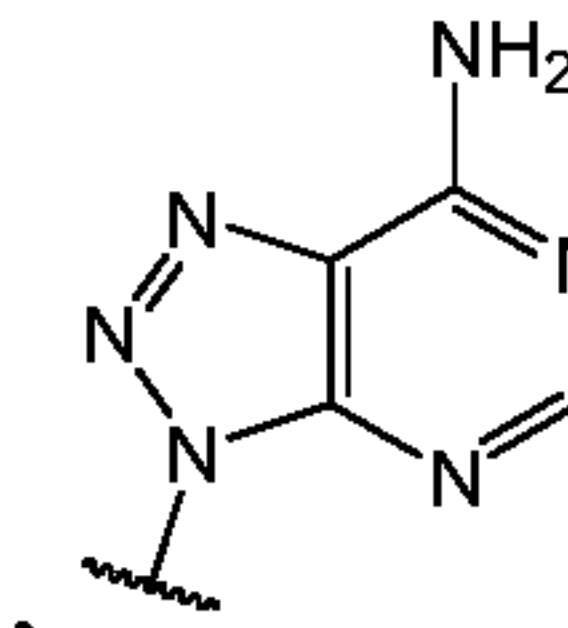
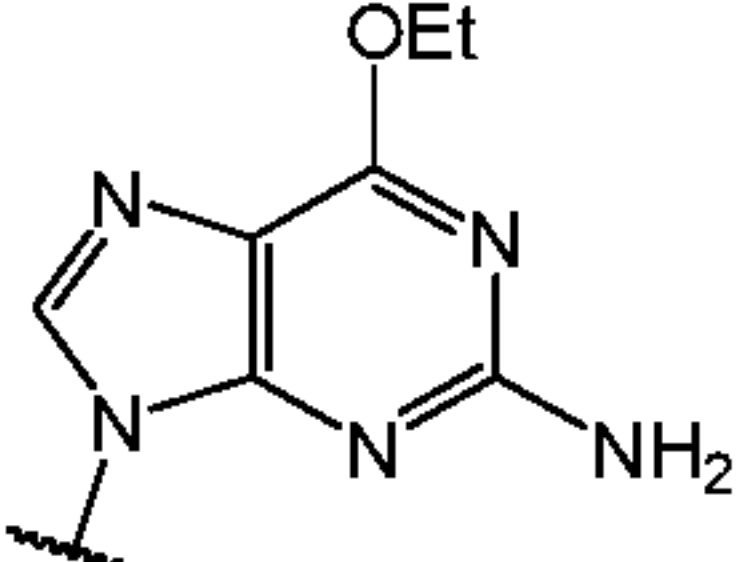


, R⁵ and R³ are not both selected from the group consisting of H, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, where said C₁-C₆ alkyl, C₂-C₆ alkenyl and C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I and OH.

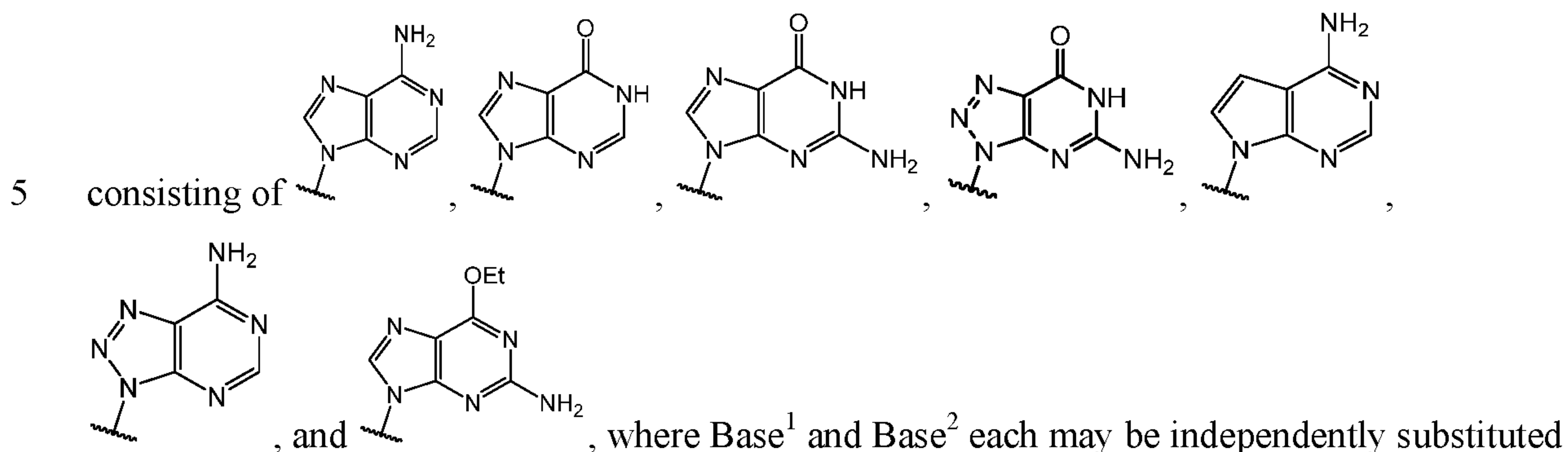
[0136] In further aspects of this embodiment, when Base¹ and Base² are each selected

from the group consisting of  and , and R^{2a} is F and R⁵ is F, at least one of X^c and X^{c1} is SR⁹.

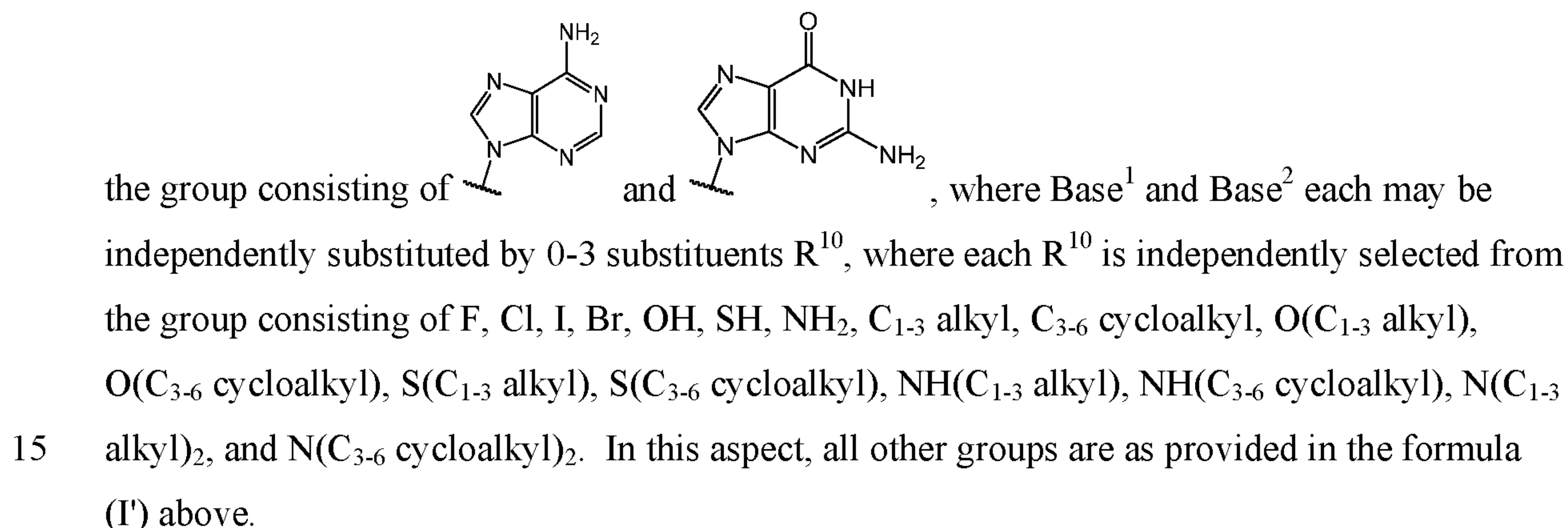
[0137] In aspects of this embodiment, Base¹ and Base² are each independently selected

from the group consisting of , , , , , , , , , , , and , where Base¹ and Base² each may be independently substituted by 0-3

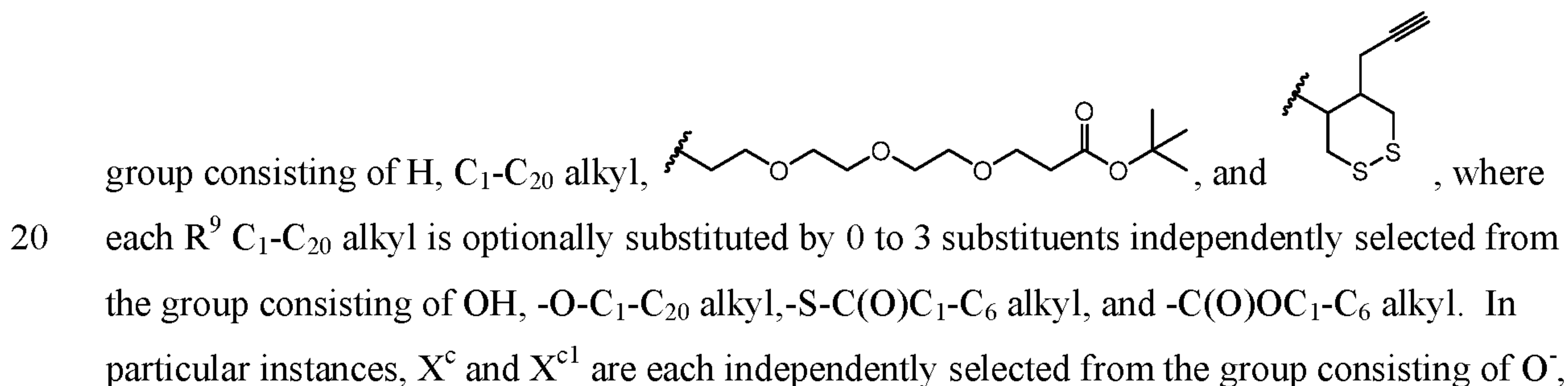
substituents R^{10} , where each R^{10} is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH_2 , C_{1-3} alkyl, C_{3-6} cycloalkyl, $O(C_{1-3}$ alkyl), $O(C_{3-6}$ cycloalkyl), $S(C_{1-3}$ alkyl), $S(C_{3-6}$ cycloalkyl), $NH(C_{1-3}$ alkyl), $NH(C_{3-6}$ cycloalkyl), $N(C_{1-3}$ alkyl) $_2$, and $N(C_{3-6}$ cycloalkyl) $_2$. In particular instances, $Base^1$ and $Base^2$ are each independently selected from the group

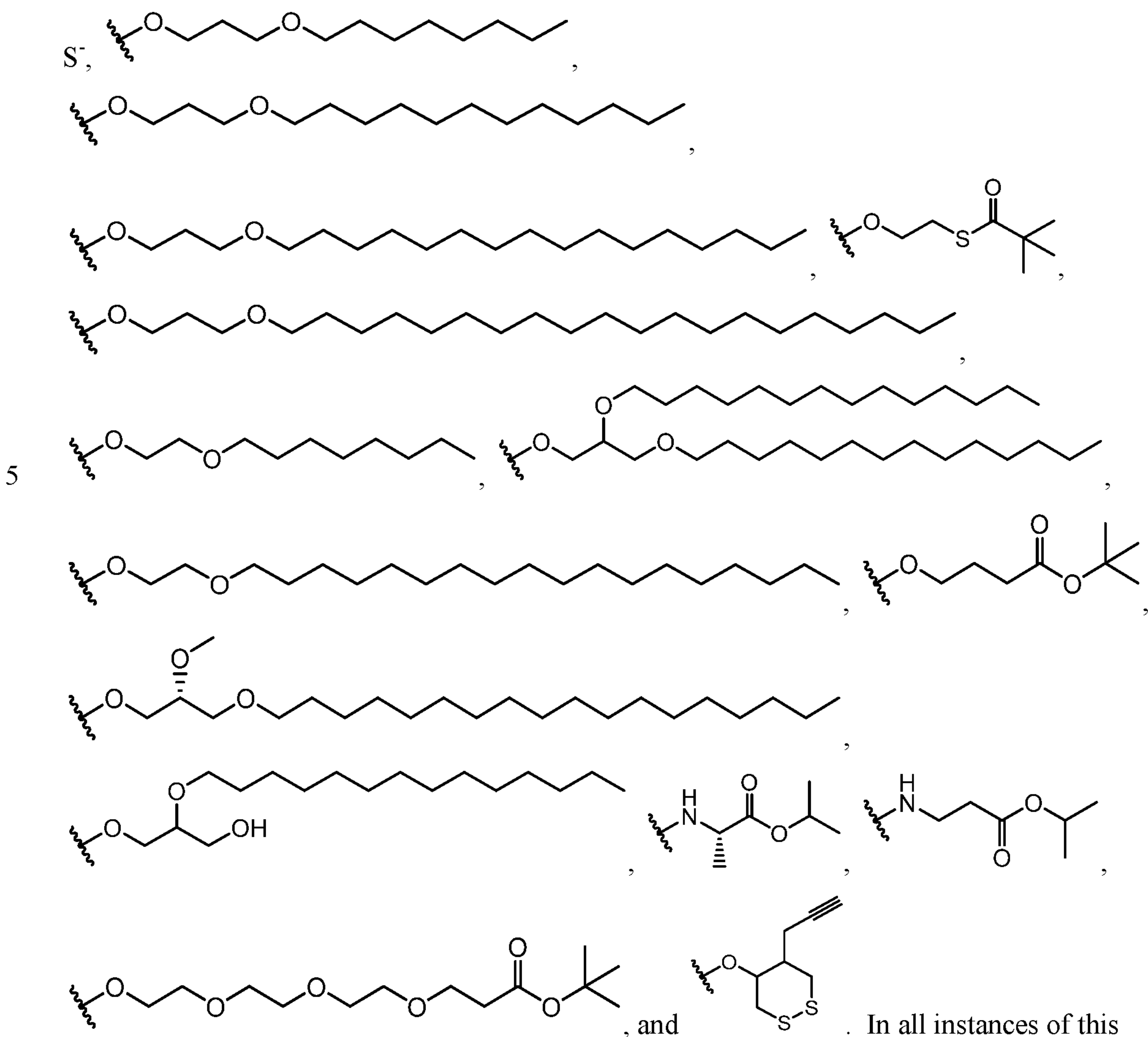


by 0-3 substituents R^{10} , where each R^{10} is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH_2 , C_{1-3} alkyl, C_{3-6} cycloalkyl, $O(C_{1-3}$ alkyl), $O(C_{3-6}$ cycloalkyl), $S(C_{1-3}$ alkyl), $S(C_{3-6}$ cycloalkyl), $NH(C_{1-3}$ alkyl), $NH(C_{3-6}$ cycloalkyl), $N(C_{1-3}$ alkyl) $_2$, and $N(C_{3-6}$ cycloalkyl) $_2$. In more particular instances, $Base^1$ and $Base^2$ are each independently selected from



[0138] In aspects of this embodiment, X^c and X^{c1} are each independently selected from the group consisting of OR^9 , SR^9 , and NR^9R^9 , where each R^9 is independently selected from the





aspect, all other groups are as provided in the formula (I') above or in the aspect described above.

[0139] In aspects of this embodiment, R^1 and R^{1a} are each H. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0140] In aspects of this embodiment, R^2 and R^{2a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^2 and R^{2a} C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 . In particular instances, R^2 and R^{2a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, CN, N_3 , CF_3 , CH_3 , CH_2OH , and CH_2CH_3 . In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0141] In aspects of this embodiment, R^3 is selected from the group consisting of H, F, Cl, I, Br, OH, N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^3 C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 . In particular instances, R^3 are each independently selected from the group consisting of H, F, Cl, I, Br, OH, CN, N_3 , CF_3 , CH_3 , CH_2OH , and CH_2CH_3 . In more particular instances, R^3 is selected from NH_2 and N_3 . In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0142] In aspects of this embodiment, R^4 and R^{4a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^4 and R^{4a} C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 . In particular instances, R^4 and R^{4a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, CN, N_3 , CF_3 , CH_3 , CH_2OH , and CH_2CH_3 . In more particular instances, R^4 and R^{4a} are each F. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0143] In aspects of this embodiment, R^5 is selected from the group consisting of H, F, Cl, Br, I, OH, NH_2 , N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^5 C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, NR^9R^9 , and N_3 . In particular instances, R^5 are each independently selected from the group consisting of H, F, Cl, I, Br, OH, CN, N_3 , CF_3 , CH_3 , CH_2OH , and CH_2CH_3 . In even more particular instances, R^5 is selected from NH_2 and N_3 . In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0144] In aspects of this embodiment, R^6 and R^{6a} are each independently selected from the group consisting of H, F, Cl, I, Br, OH, C_1 - C_6 alkyl, C_2 - C_6 alkenyl, and C_2 - C_6 alkynyl. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0145] In aspects of this embodiment, R^7 and R^{7a} are each independently selected from the group consisting of H and C_1 - C_6 alkyl. In particular instances, R^7 and R^{7a} are each independently selected from the group consisting of H and CH_3 . In more particular instances, R^{7a} is CH_3 . In additional instances, R^7 and R^{7a} are each H. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0146] In aspects of this embodiment, R^8 and R^{8a} are each independently selected from the group consisting of H and C_1 - C_6 alkyl. In particular instances, R^8 and R^{8a} are each

independently selected from the group consisting of H and CH₃. In more particular instances, R^{8a} is CH₃. In additional instances, R⁸ and R^{8a} are each H. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0147] In aspects of this embodiment, R^{1a} and R³ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R^{1a} and R³ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0148] In aspects of this embodiment, R^{2a} and R³ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R^{2a} and R³ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0149] In aspects of this embodiment, R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, and -O-C₂-C₆ alkynylene, such that where R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

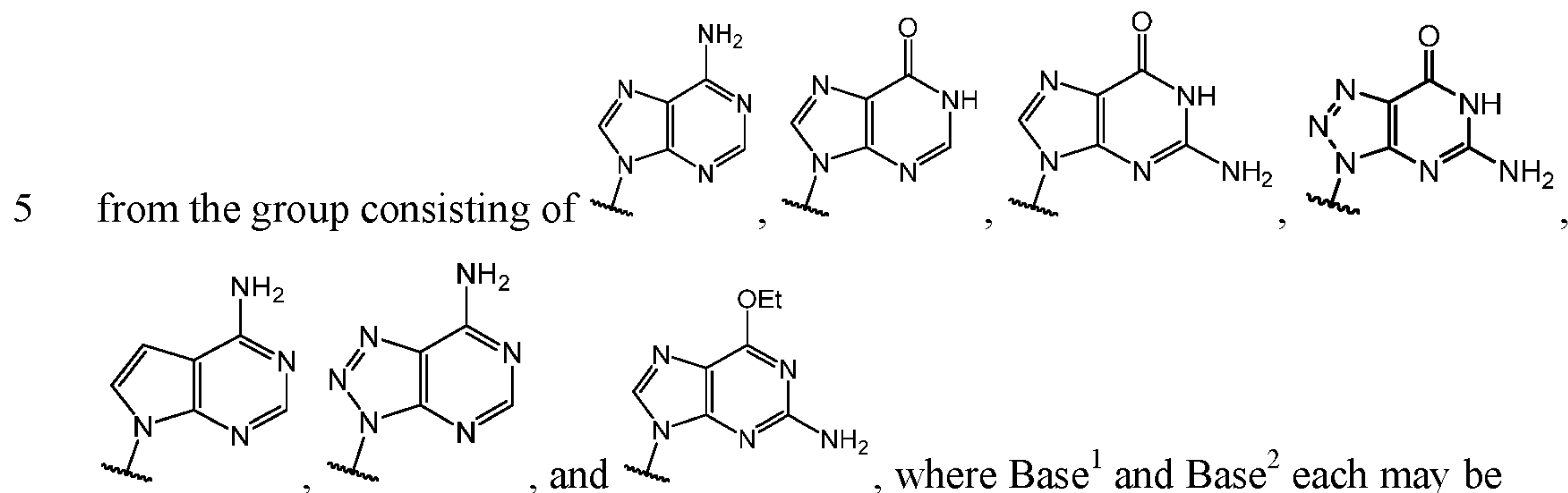
[0150] In aspects of this embodiment, R⁴ and R⁵ are connected by C₁-C₆ alkylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R⁴ and R⁵ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R⁵ position. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0151] In aspects of this embodiment, R⁵ and R⁶ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R⁵ and R⁶ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R⁵ position. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0152] In aspects of this embodiment, R⁷ and R⁸ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, or C₂-C₆ alkynylene. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

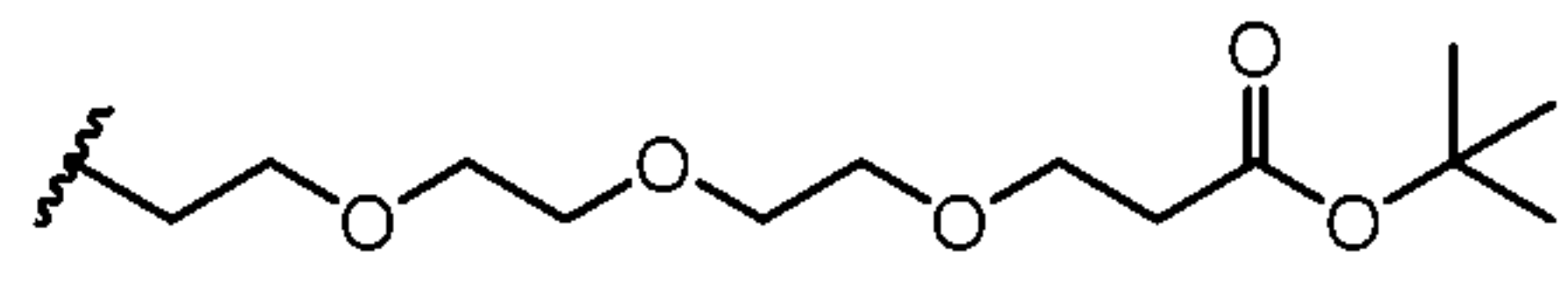
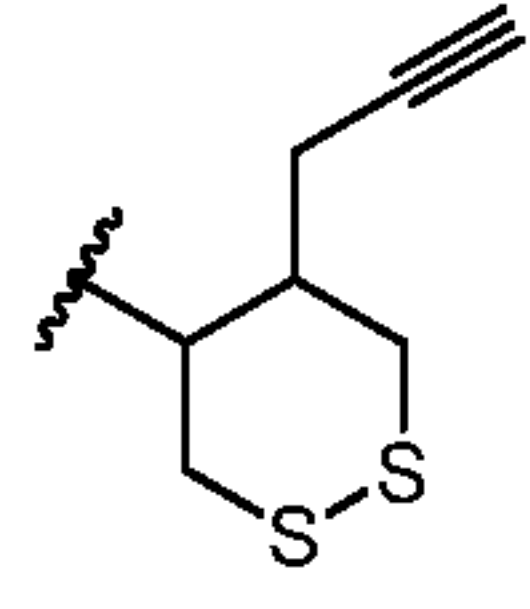
[0153] In aspects of this embodiment, R^{7a} and R^{8a} are connected to form C_1 - C_6 alkylene, C_2 - C_6 alkenylene, or C_2 - C_6 alkynylene. In this aspect, all other groups are as provided in the formula (I') above or in the aspects described above.

[0154] In aspects of this embodiment, $Base^1$ and $Base^2$ are each independently selected



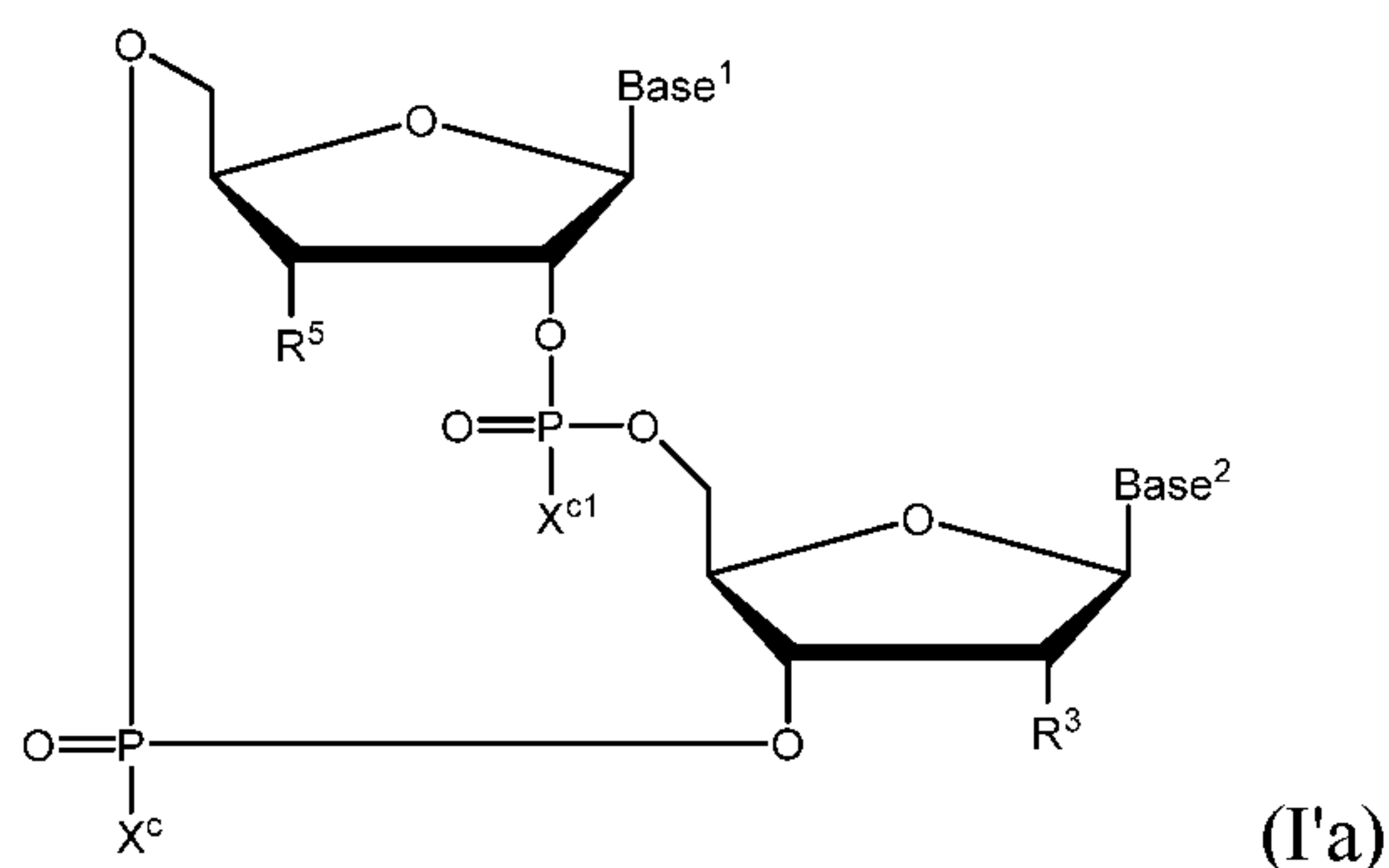
where $Base^1$ and $Base^2$ each may be independently substituted by 0-3 substituents R^{10} , where each R^{10} is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH_2 , C_{1-3} alkyl, C_{3-6} cycloalkyl, $O(C_{1-3}$ alkyl), $O(C_{3-6}$ cycloalkyl), $S(C_{1-3}$ alkyl), $S(C_{3-6}$ cycloalkyl), $NH(C_{1-3}$ alkyl), $NH(C_{3-6}$ cycloalkyl), $N(C_{1-3}$ alkyl)₂, and $N(C_{3-6}$ cycloalkyl)₂; Y and Y^a are each independently selected from the group consisting of -O- and -S-; X^a and X^{a1} are each independently selected from the group consisting of O and S; X^b and X^{b1} are each independently selected from the group consisting of O and S; X^c and X^{c1} are each independently selected from the group consisting of OR^9 , SR^9 , and NR^9R^9 ; X^d and X^{d1} are each independently selected from the group consisting of O and S; R^1 and R^{1a} are each H; R^2 and R^{2a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^2 and R^{2a} C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 ; R^3 is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^3 C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 ; R^4 and R^{4a} are each independently selected from the group consisting of H, C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^4 and R^{4a} C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 ; R^5 is selected from the group consisting of H, F, Cl, Br, I, OH, NH_2 , N_3 , C_1 - C_6 alkyl, and C_1 - C_6 haloalkyl, where said R^5 C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, NR^9R^9 , and N_3 ; R^6 and R^{6a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, C_1 - C_6 alkyl, C_2 - C_6 alkenyl, C_2 - C_6 alkynyl, and C_1 - C_6

haloalkyl, where said R^6 and R^{6a} C_1 - C_6 alkyl or C_1 - C_6 haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N_3 ; R^7 and R^{7a} are each H; R^8 and R^{8a} are each H; each R^9 is independently selected from the group consisting of H,

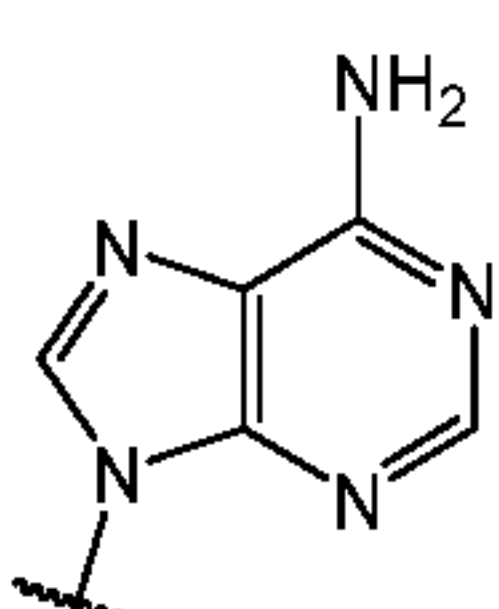
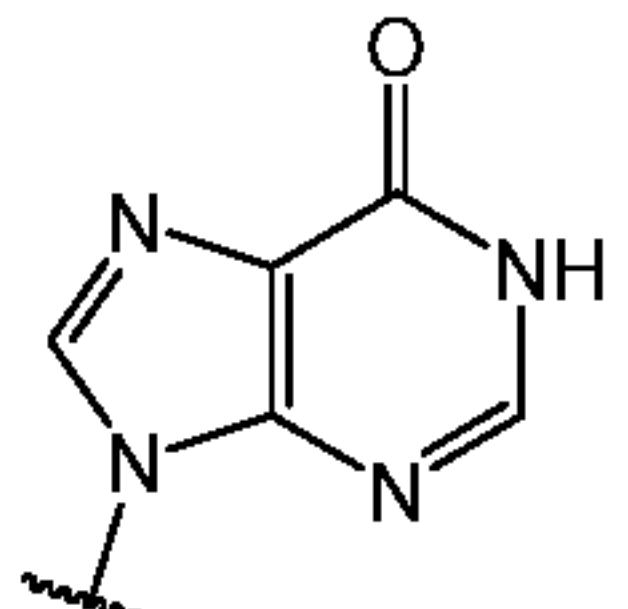
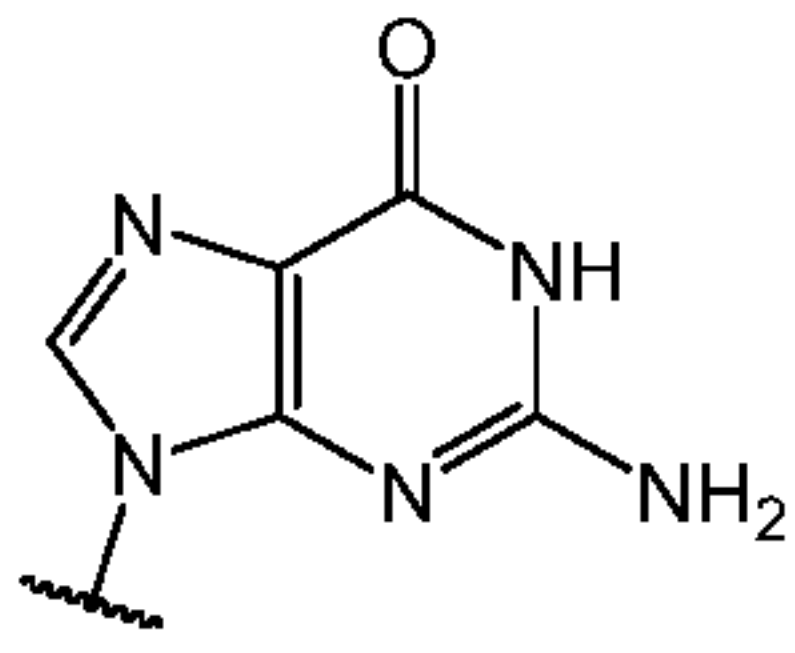
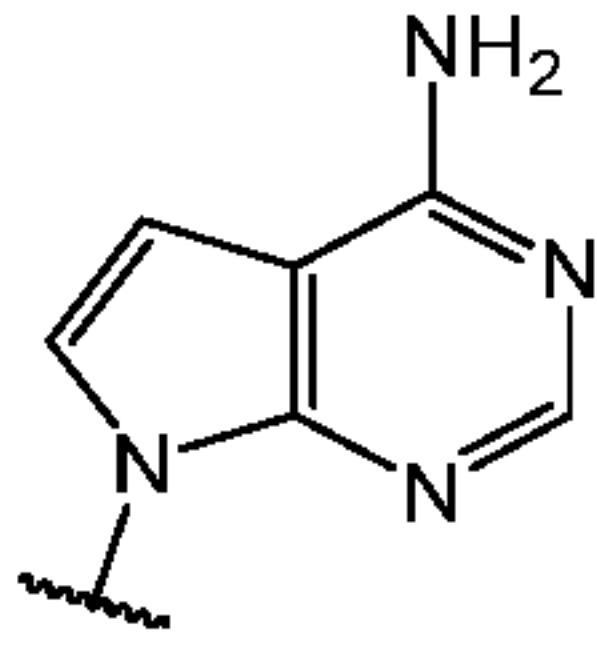
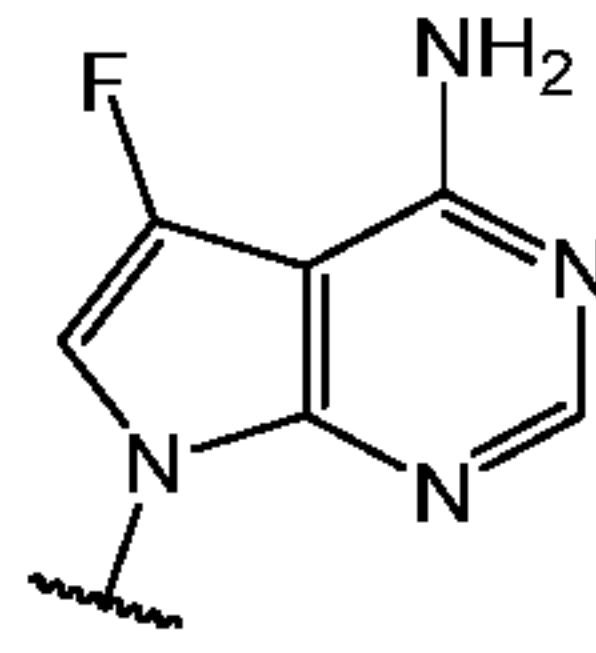
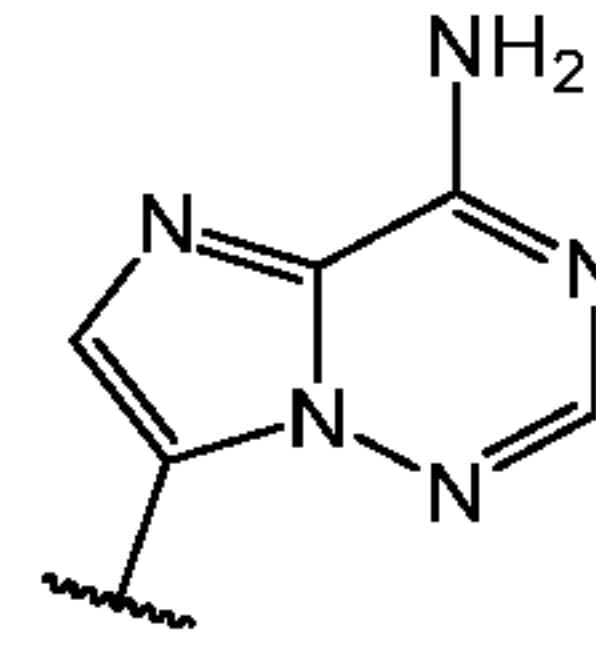
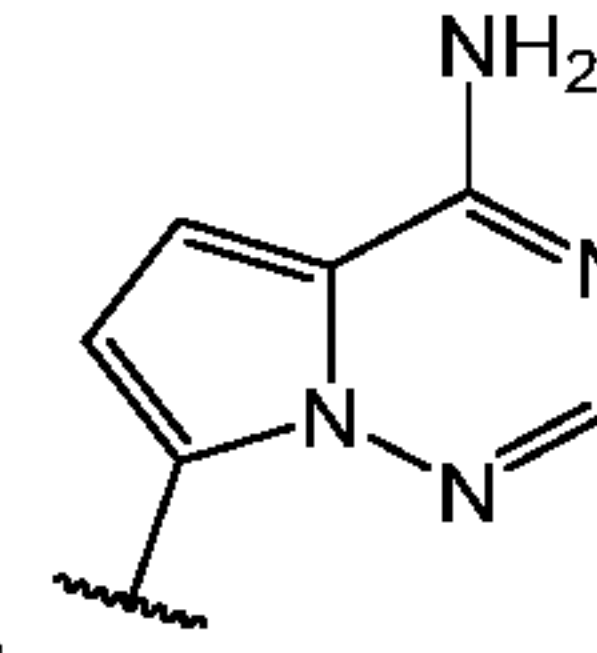
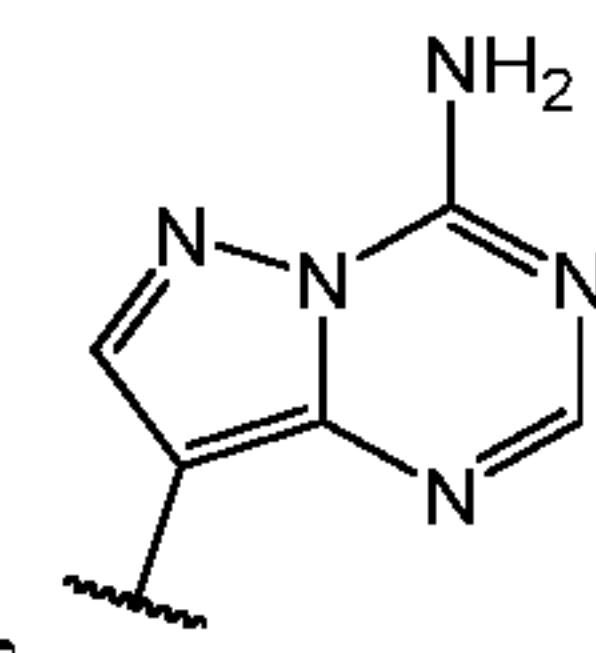
C_2 - C_3 alkyl, , and , where each R^9 C_2 - C_3 alkyl is

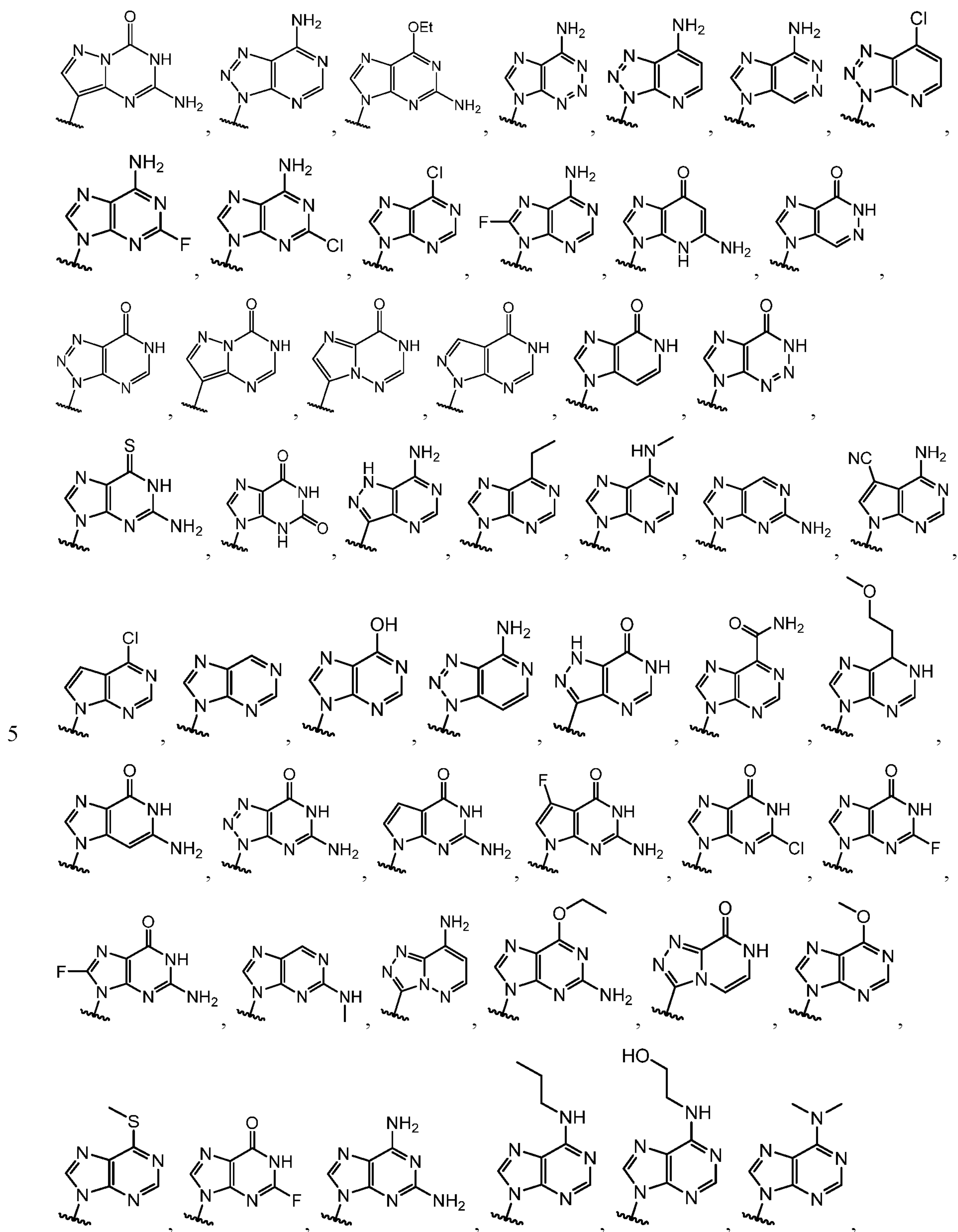
optionally substituted by 1 to 2 substituents independently selected from the group consisting of OH, -O- C_1 - C_{20} alkyl, -S-C(O) C_1 - C_6 alkyl, and -C(O)OC $_1$ - C_6 alkyl; optionally R^3 and R^{6a} are connected to form -O- C_1 - C_6 alkylene, -O- C_2 - C_6 alkenylene, and -O- C_2 - C_6 alkynylene, such that where R^3 and R^{6a} are connected to form -O- C_1 - C_6 alkylene, -O- C_2 - C_6 alkenylene, or -O- C_2 - C_6 alkynylene, said O is bound at the R^3 position or optionally R^4 and R^5 are connected by C_1 - C_6 alkylene, -O- C_1 - C_6 alkylene, -O- C_2 - C_6 alkenylene, or -O- C_2 - C_6 alkynylene, such that where R^4 and R^5 are connected to form -O- C_1 - C_6 alkylene, -O- C_2 - C_6 alkenylene, or -O- C_2 - C_6 alkynylene, said O is bound at the R^5 position. In all instances of this aspect, all other groups are as provided in the formula (I') above.

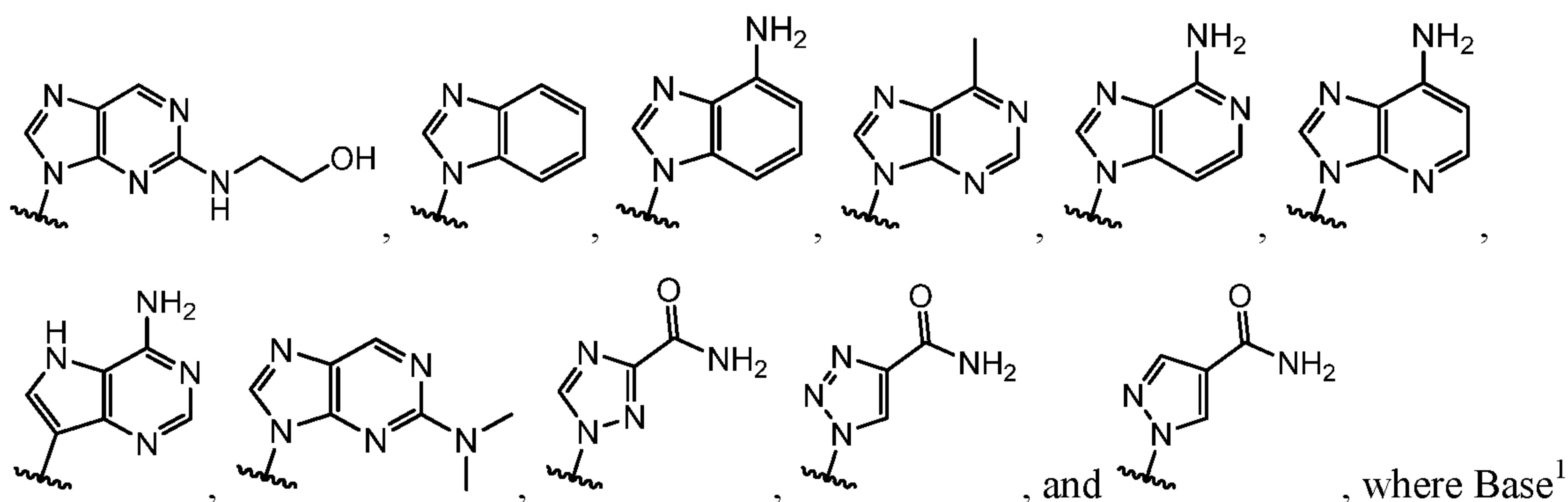
[0155] In aspects of this embodiment, the compound of formula (I') is a compound of formula (I'a):



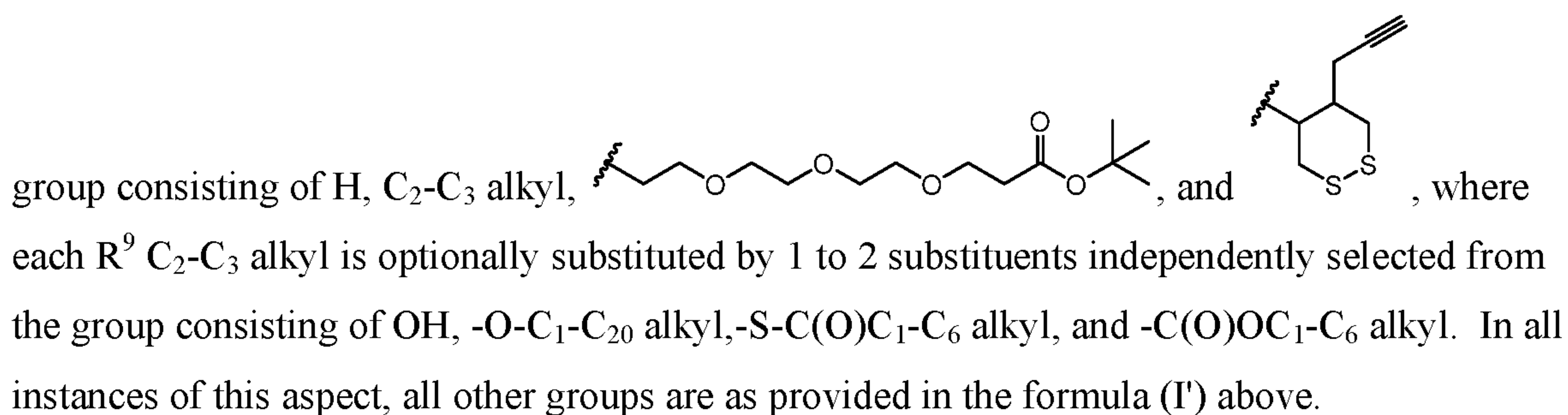
or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein $Base^1$ and

$Base^2$ are each independently selected from the group consisting of , , , , , , , , , , , ,

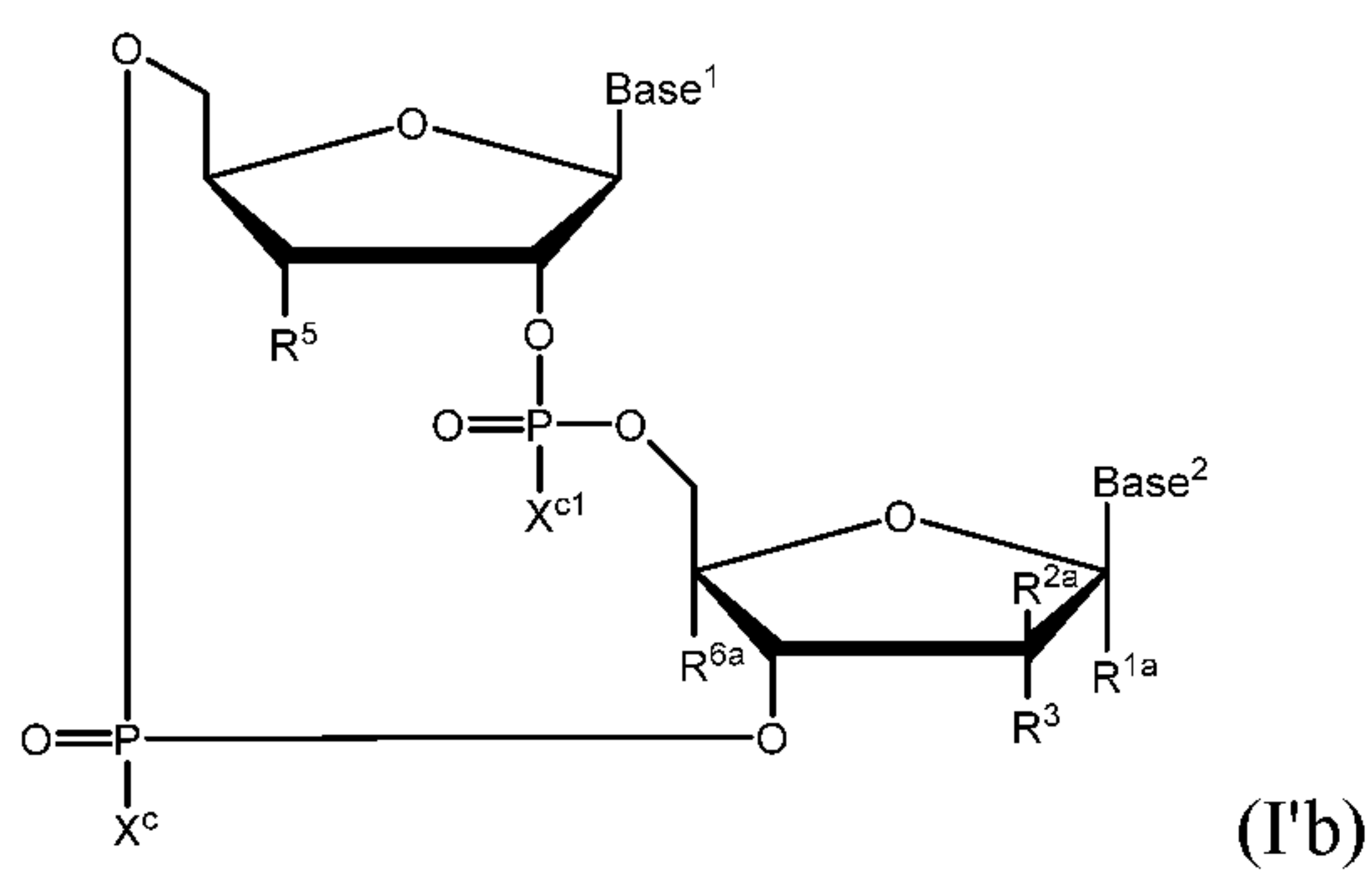




and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂; X^c and X^{c1} are each independently selected from the group consisting of OR⁹, SR⁹, and NR⁹R⁹; R³ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C_{1-C6} alkyl, and C_{1-C6} haloalkyl, where said R³ C_{1-C6} alkyl or C_{1-C6} haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, and OH; R⁵ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, NH₂, N₃, C_{1-C6} alkyl, and C_{1-C6} haloalkyl, where said R⁵ C_{1-C6} alkyl or C_{1-C6} haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, and OH; R³ and R⁵ are not both selected from the group consisting of: OH, R⁵ C_{1-C6} alkyl substituted with OH, or C_{1-C6} haloalkyl substituted with OH; and each R⁹ is independently selected from the

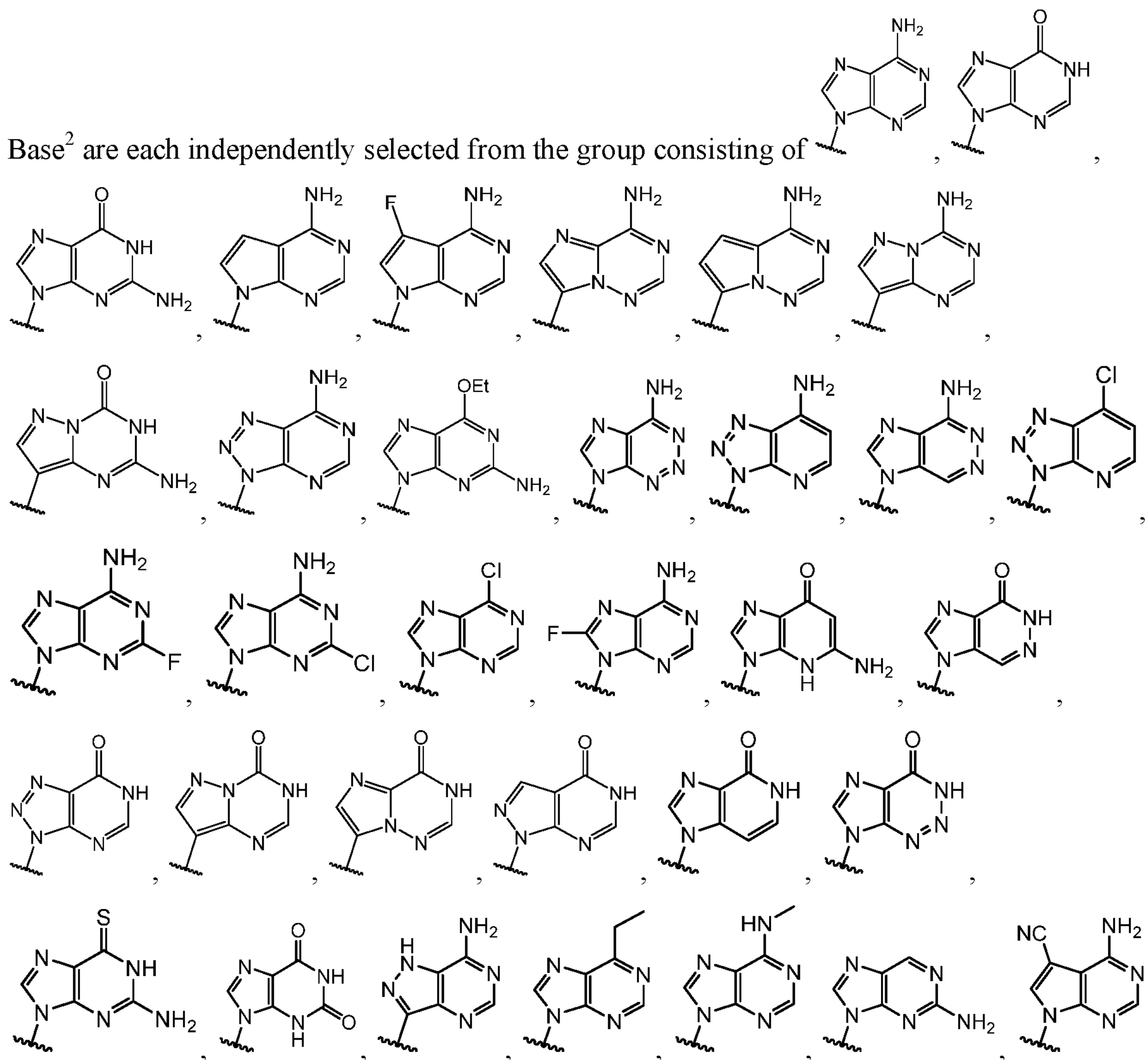


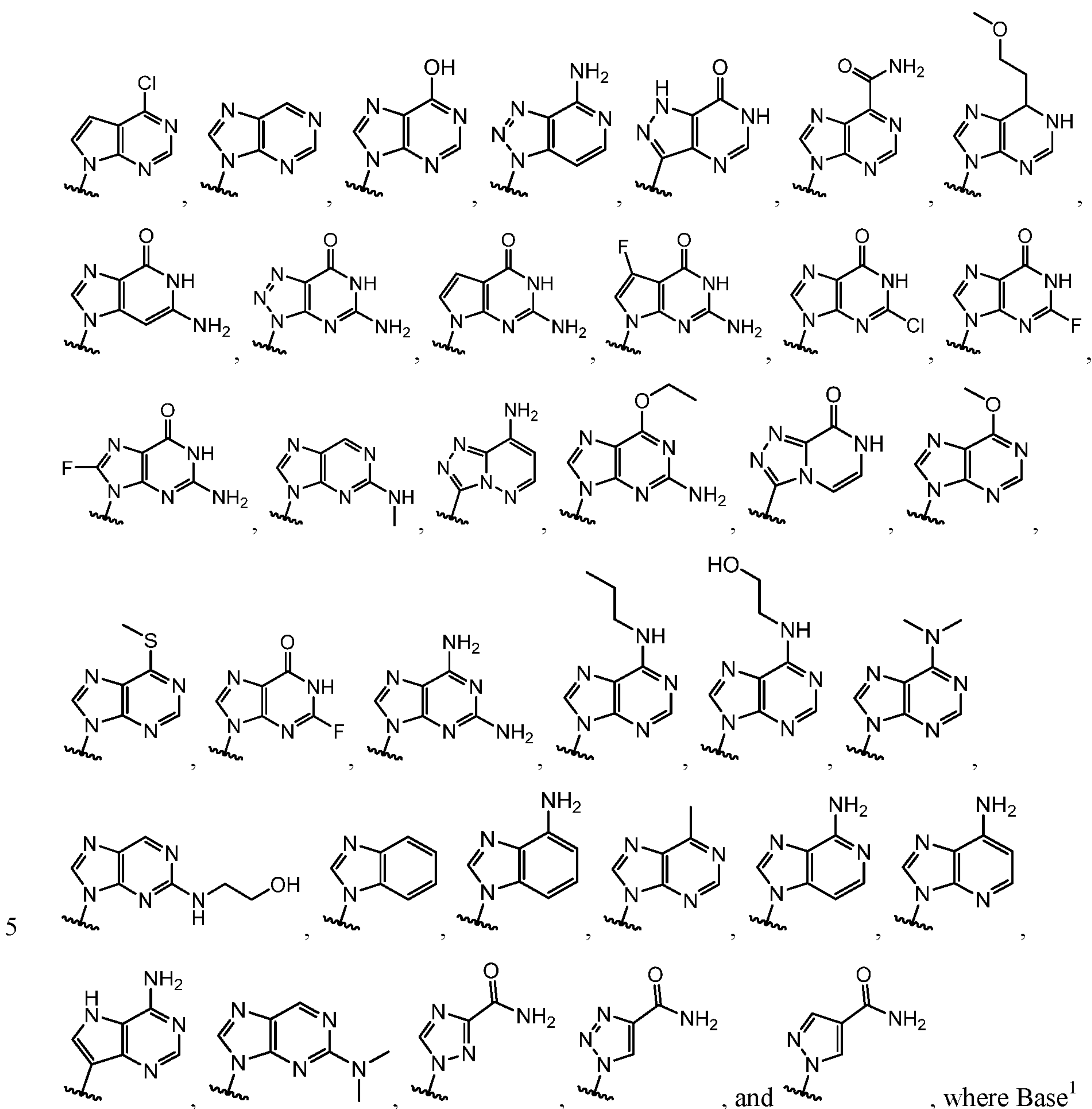
[0156] In aspects of this embodiment, the compound of formula (I') is a compound of formula (I'b):



or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein Base¹ and

Base² are each independently selected from the group consisting of

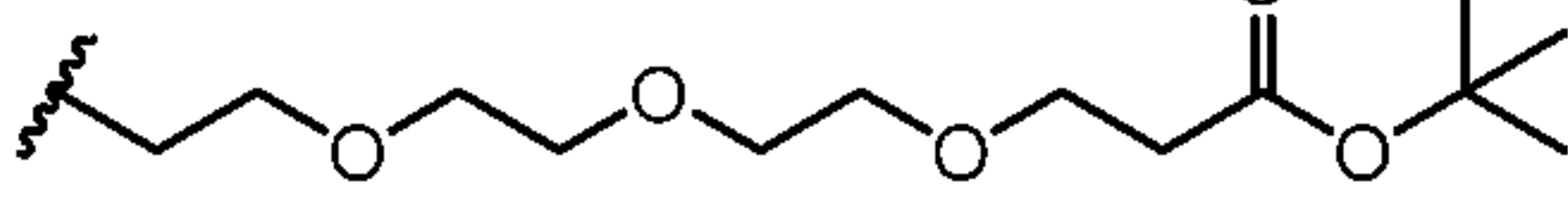
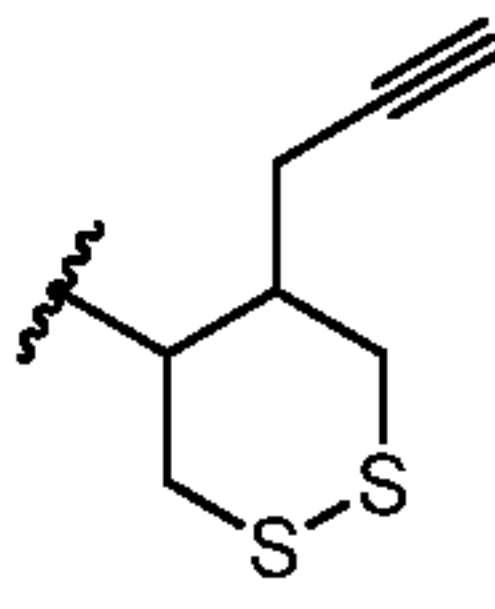




and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂; X^c and X^{cl} are each independently selected from the group consisting of OR⁹, SR⁹, and NR⁹R⁹; R^{1a} is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl, where said R^{1a} C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl,

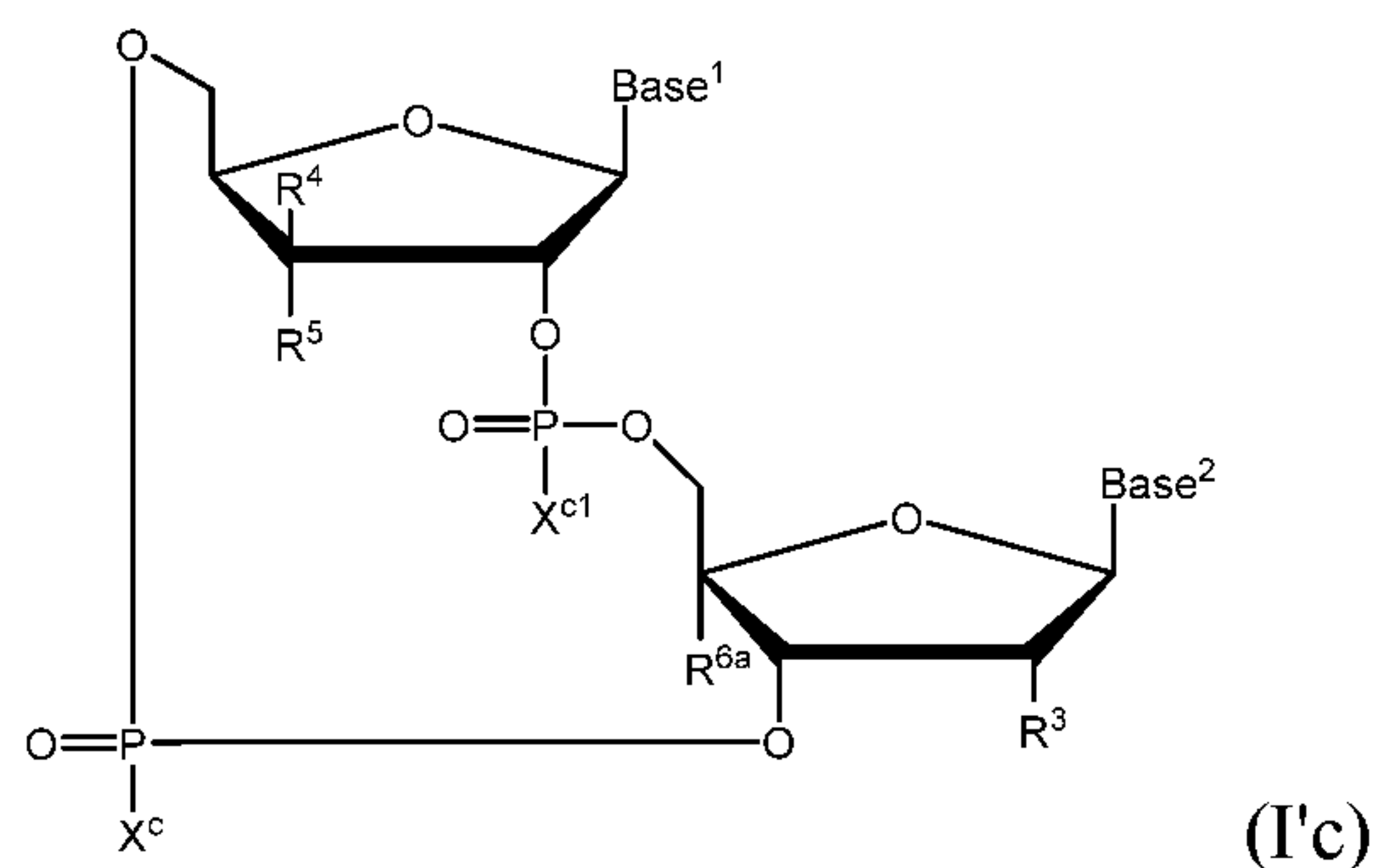
C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R^{2a} is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, and C₁-C₆ haloalkyl, where said R^{2a} C₁-C₆ alkyl or C₁-C₆ haloalkyl are substituted by 0 to 3

- 5 substituents selected from the group consisting of F, Cl, Br, I, and OH; R³ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, and C₁-C₆ haloalkyl, where said R³ C₁-C₆ alkyl or C₁-C₆ haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, and OH; R⁵ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, NH₂, N₃, C₁-C₆ alkyl, and C₁-C₆ haloalkyl, where said R⁵ C₁-C₆ alkyl or C₁-C₆ haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, and OH; R³ and R⁵ are not both selected from the group consisting of OH, C₁-C₆ alkyl substituted with OH, and C₁-C₆ haloalkyl substituted with OH; R^{6a} is selected from the group consisting of H, F, Cl, Br, I, OH, C₁-C₆ alkyl, C₂-C₆ alkenyl, and C₂-C₆ alkynyl; each R⁹ is independently selected from

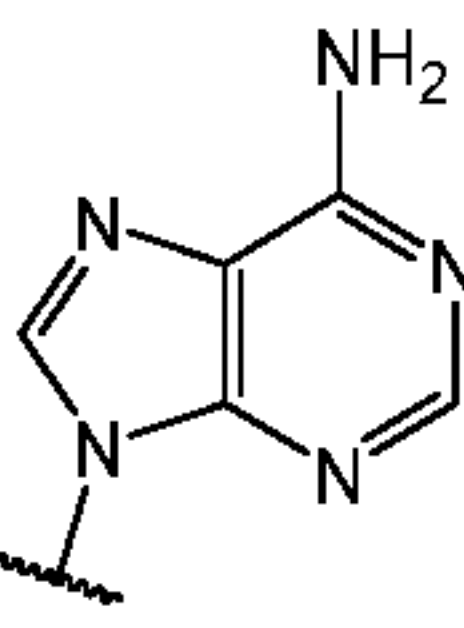
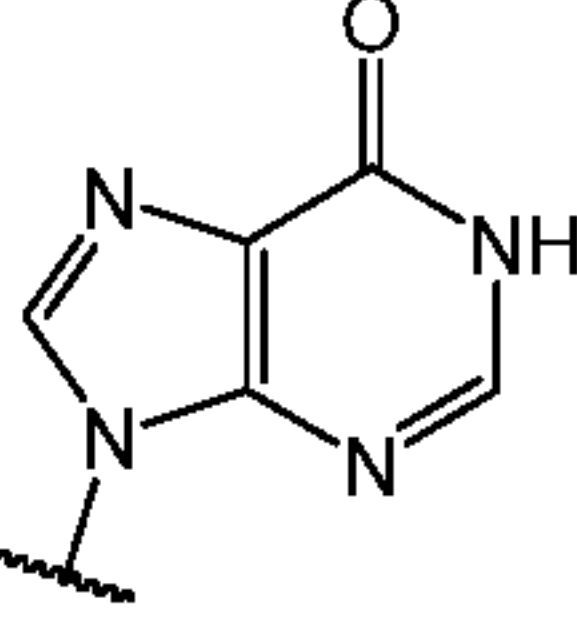
the group consisting of H, C₂-C₃ alkyl, , and ,

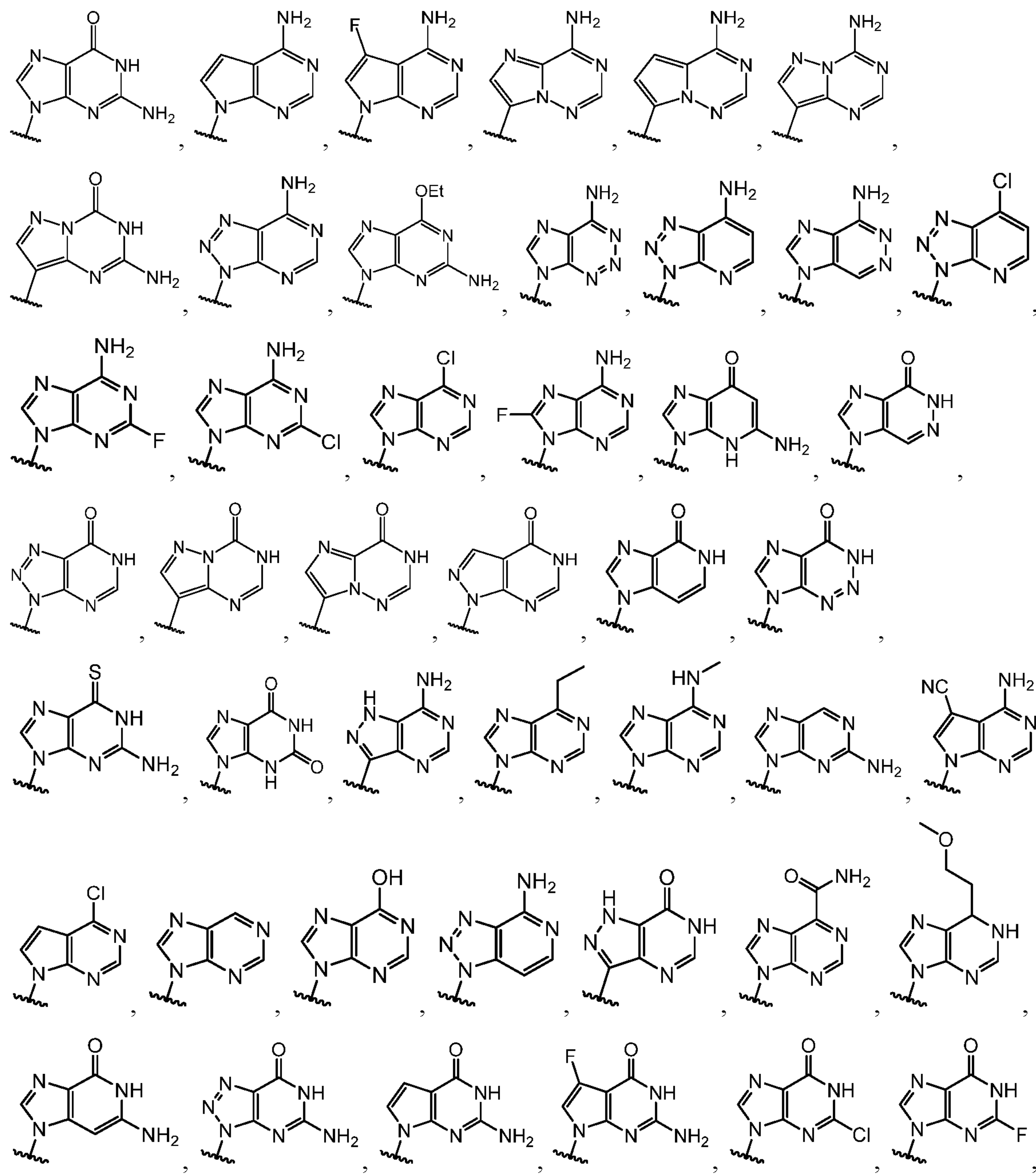
- 15 where each R⁹ C₂-C₃ alkyl is optionally substituted by 1 to 2 substituents independently selected from the group consisting of OH, -O-C₁-C₂₀ alkyl, -S-C(O)C₁-C₆ alkyl, and -C(O)OC₁-C₆ alkyl; and optionally R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, and -O-C₂-C₆ alkynylene, such that where R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position. In this aspect,
- 20 all other groups are as provided in the formula (I') above.

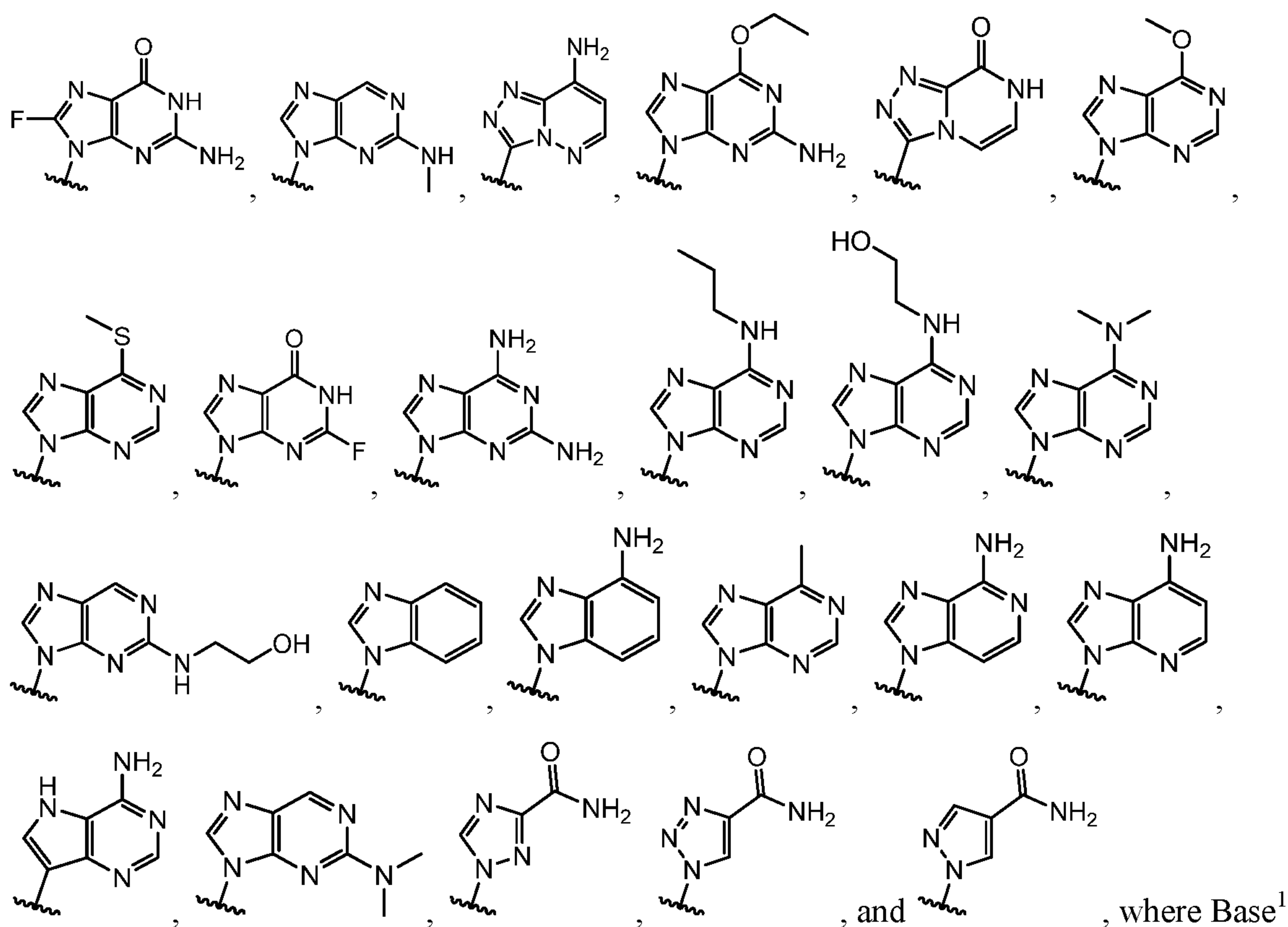
[0157] In aspects of this embodiment, the compound of formula (I') is a compound of formula (I'c):



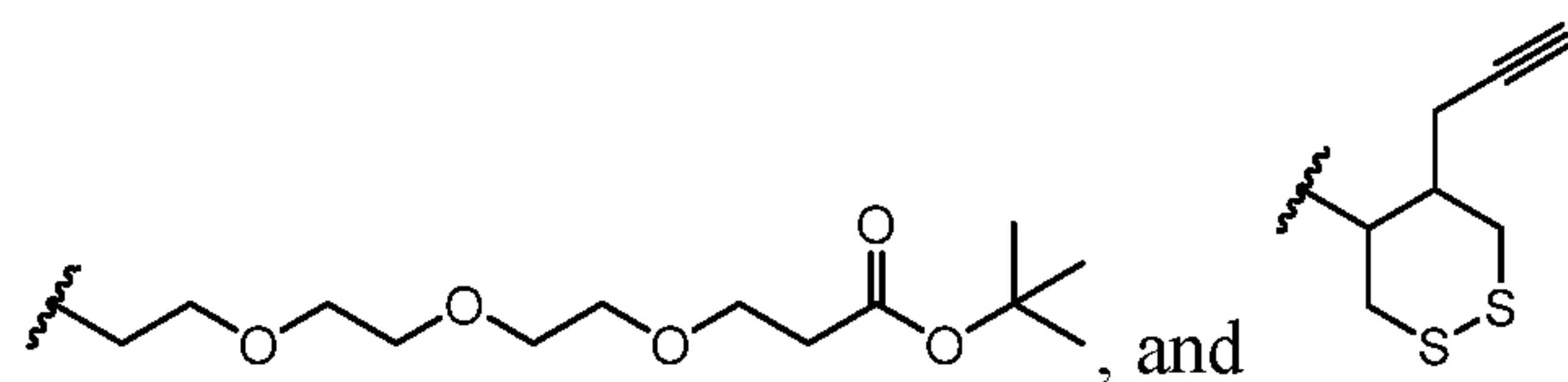
or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein Base¹ and

Base² are each independently selected from the group consisting of  ,  ,



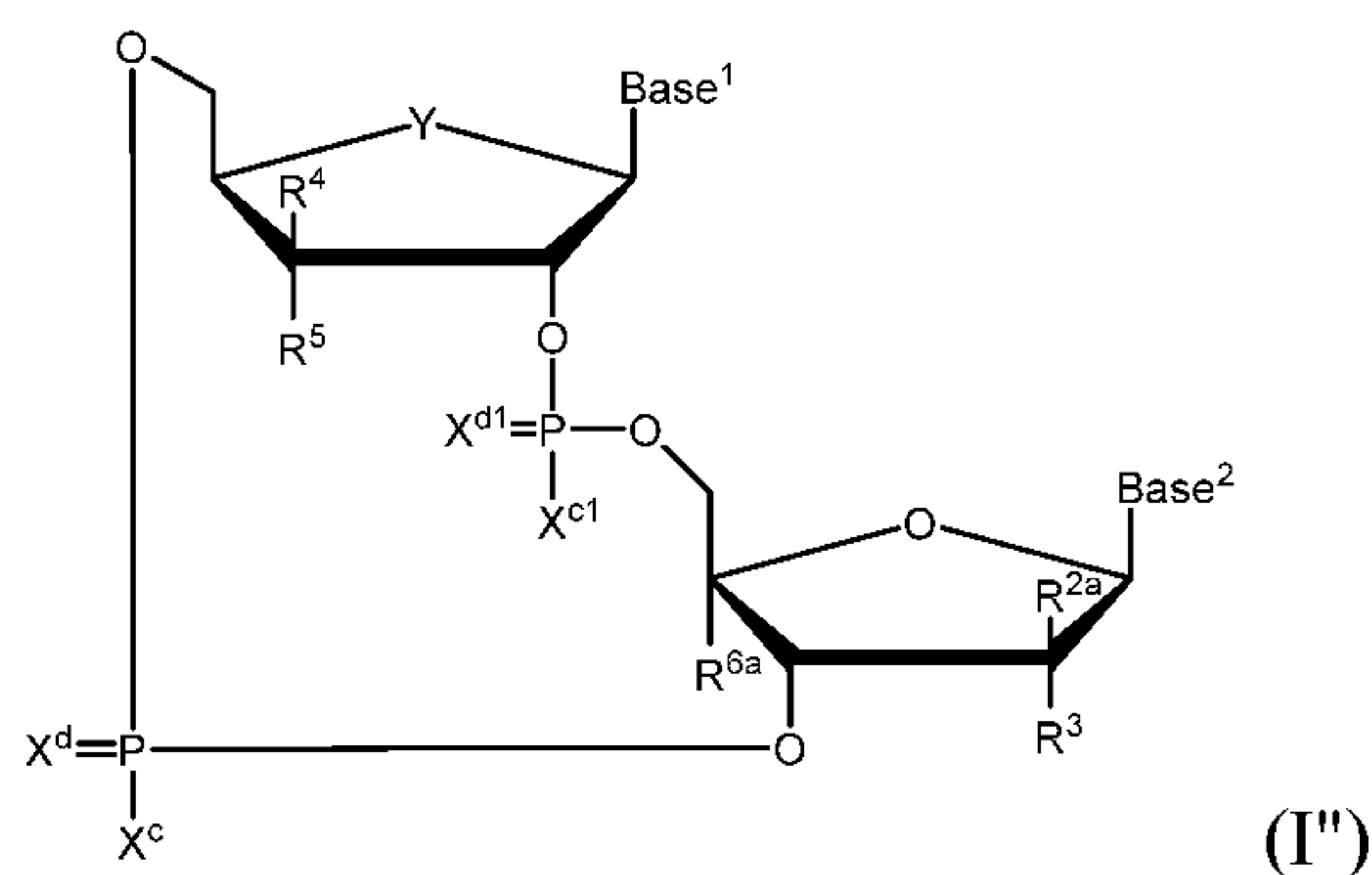


- 5 and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂; X^c and X^{cl} are each independently selected from the group consisting of OR⁹, SR⁹, and NR⁹R⁹; R³ is selected from the group
- 10 consisting of H, F, Cl, Br, I, OH, CN, N₃, C_{1-C6} alkyl, and C_{1-C6} haloalkyl, where said R³ C_{1-C6} alkyl or C_{1-C6} haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, and OH; R⁴ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C_{1-C6} alkyl, and C_{1-C6} haloalkyl, where said R⁴ C_{1-C6} alkyl or C_{1-C6} haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, and OH; R⁵ is selected from
- 15 the group consisting of H, F, Cl, Br, I, OH, CN, NH₂, N₃, C_{1-C6} alkyl, and C_{1-C6} haloalkyl, where said R⁵ C_{1-C6} alkyl or C_{1-C6} haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, and OH; R^{6a} is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C_{1-C6} alkyl, and C_{1-C6} haloalkyl, where said R^{6a} C_{1-C6} alkyl or C_{1-C6} haloalkyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, and OH; each R⁹ is independently selected from the group consisting of H, C_{2-C3} alkyl,
- 20



, where each R^9 C_2 - C_3 alkyl is optionally substituted by 1 to 2 substituents independently selected from the group consisting of OH, -O- C_1 - C_{20} alkyl, -S-C(O) C_1 - C_6 alkyl, and -C(O)OC $_1$ - C_6 alkyl; and optionally R^4 and R^5 are connected by C_1 - C_6 alkylene, -O- C_1 - C_6 alkylene, -O- C_2 - C_6 alkenylene, or -O- C_2 - C_6 alkynylene, such that where R^4 and R^5 are connected to form -O- C_1 - C_6 alkylene, -O- C_2 - C_6 alkenylene, or -O- C_2 - C_6 alkynylene, said O is bound at the R^5 position. In this aspect, all other groups are as provided in the formula (I') of the second embodiment above.

[0158] In another embodiment, the CDN STING agonist is selected from cyclic dinucleotide compounds of formula (I''):



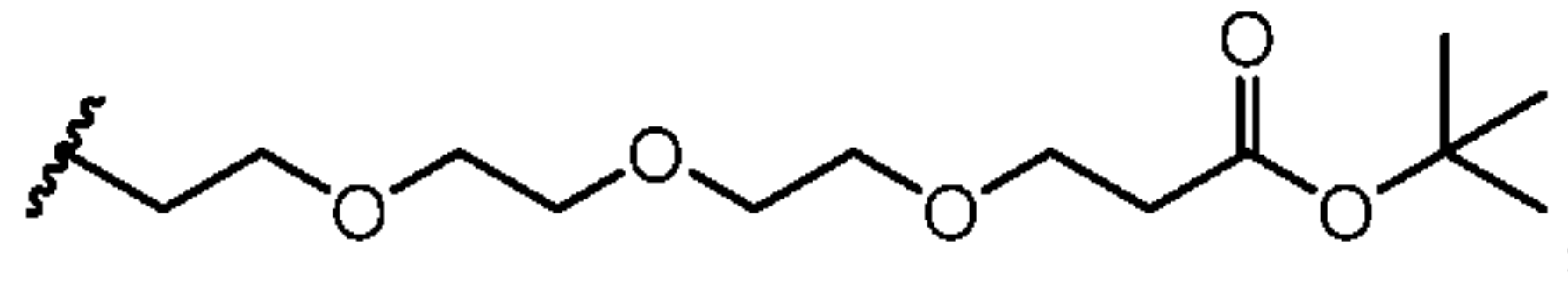
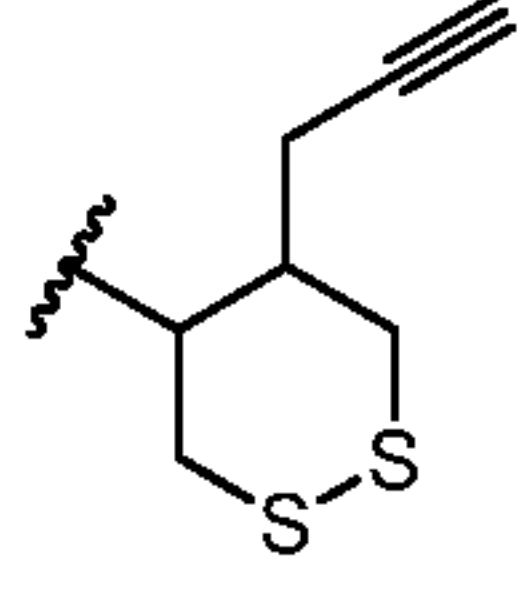
or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein Base¹ and

Base² are each independently selected from the group consisting of ,

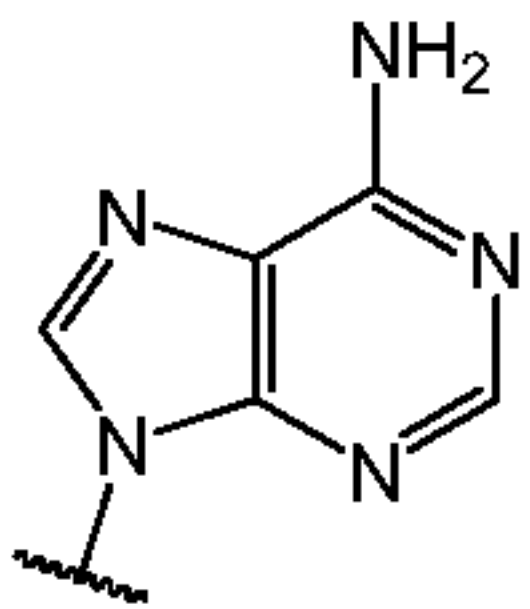
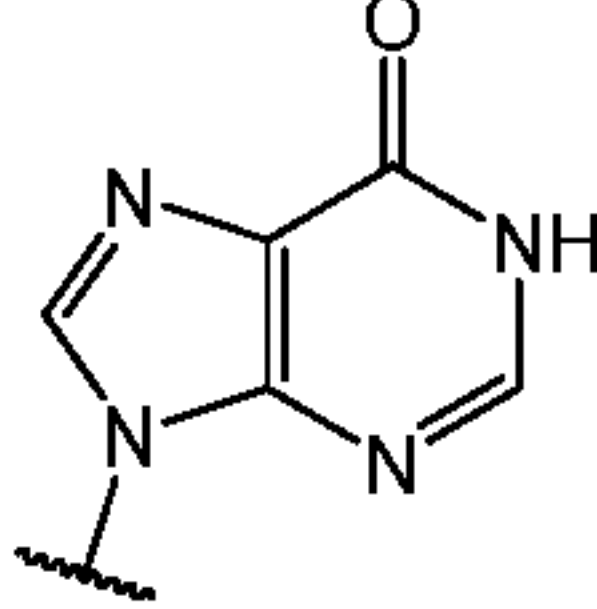
; Y is selected

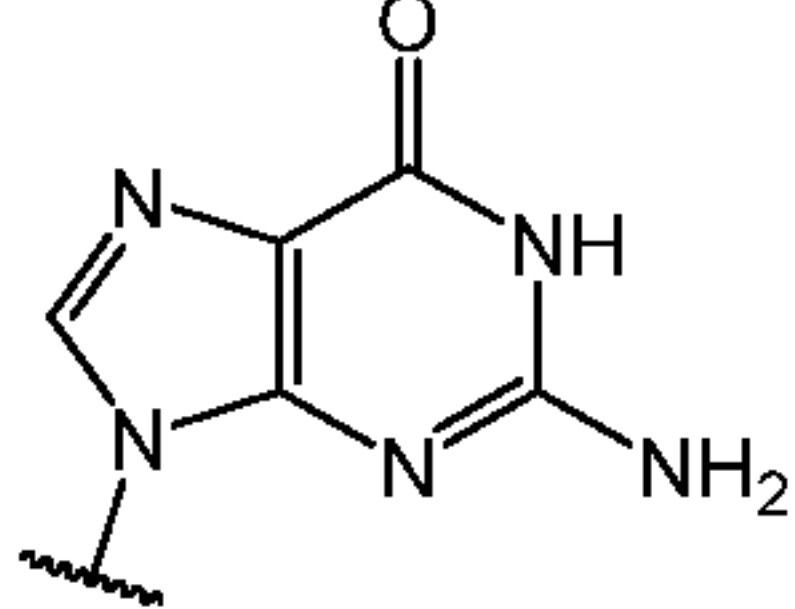
from the group consisting of -O- and -S-; X^c and X^{c1} are each independently selected from the group consisting of OR^9 and SR^9 ; X^d and X^{d1} are each independently selected from the group consisting of O and S; R^{2a} is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N_3 , C_1 - C_6 alkyl, C_2 - C_6 alkenyl, C_2 - C_6 alkynyl, C_1 - C_6 haloalkyl, C_2 - C_6 haloalkenyl, C_2 - C_6 haloalkynyl, -O- C_1 - C_6 alkyl, -O- C_2 - C_6 alkenyl, and -O- C_2 - C_6 alkynyl; R^3 is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N_3 , C_1 - C_6 alkyl, C_2 - C_6 alkenyl, C_2 - C_6 alkynyl, C_1 - C_6

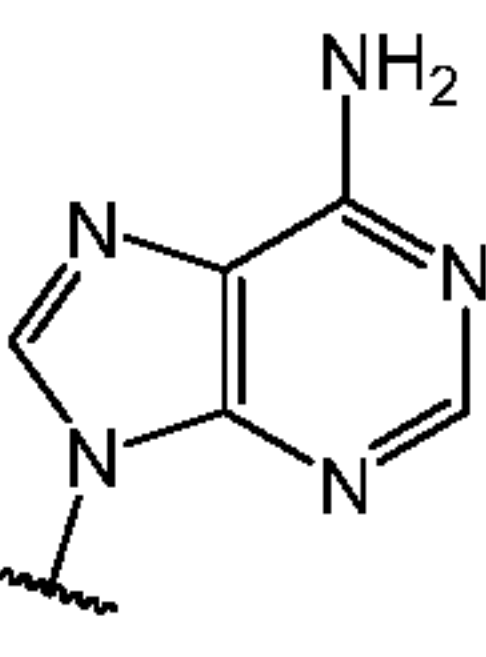
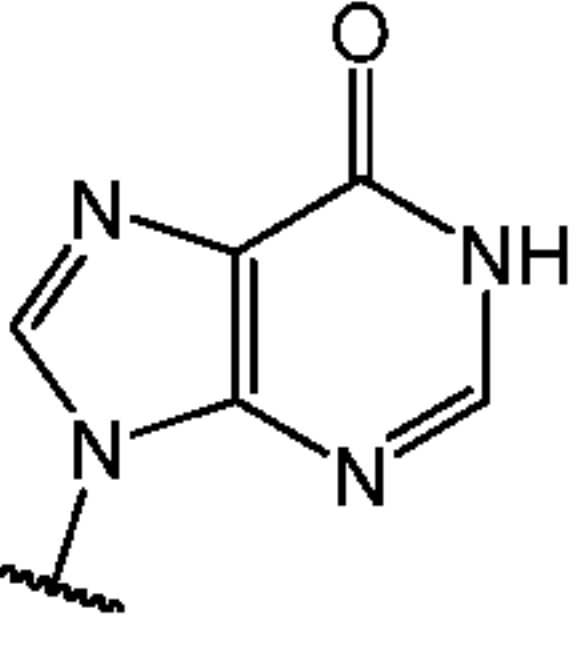
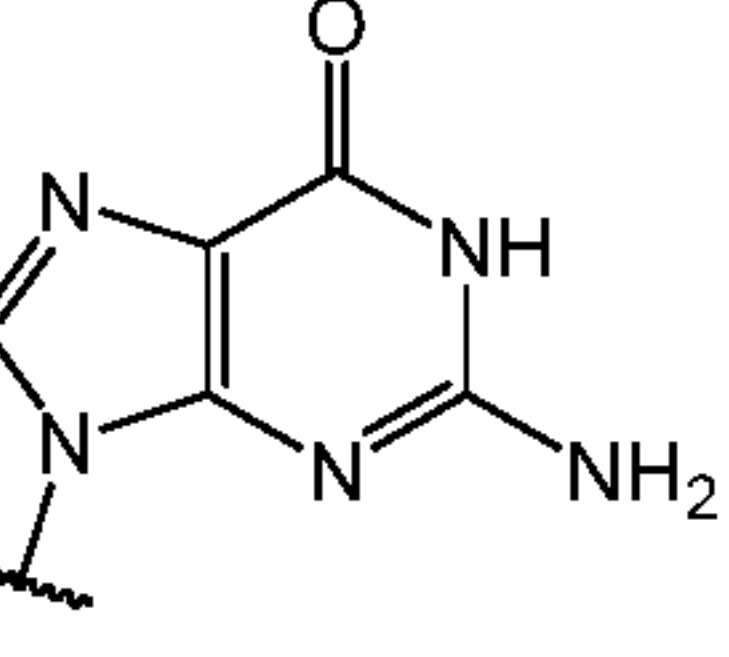
haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and
 -O-C₂-C₆ alkynyl; R⁴ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆
 alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl,
 -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl; R⁵ is selected from the group
 5 consisting of H, F, Cl, Br, I, OH, CN, NH₂, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆
 haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and
 -O-C₂-C₆ alkynyl; R^{6a} is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆
 alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl,
 -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl; each R⁹ is independently selected from

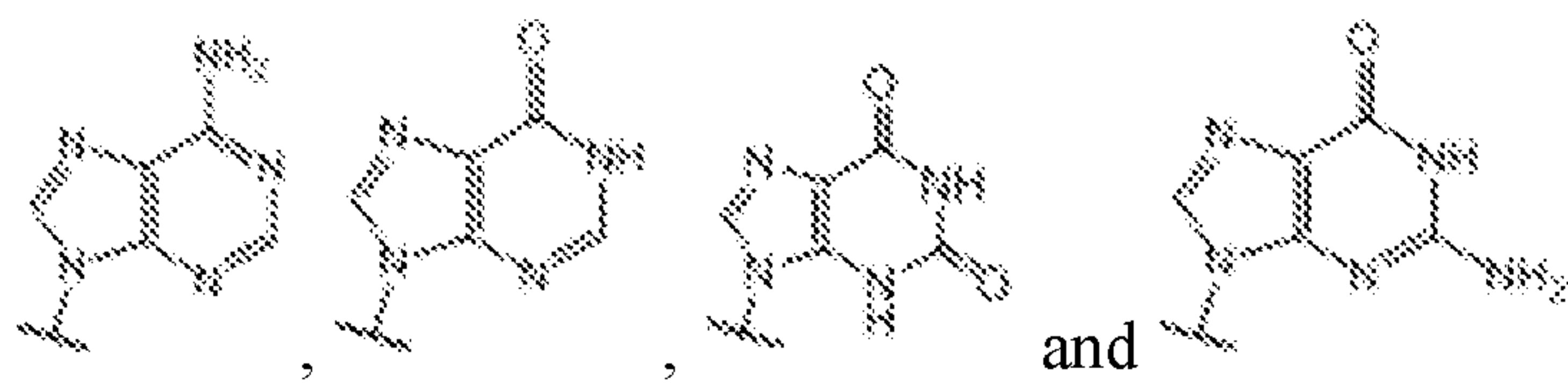
10 the group consisting of H, C₁-C₂₀ alkyl, , and ,
 where each R⁹ C₁-C₂₀ alkyl is optionally substituted by 0 to 3 substituents independently selected
 from the group consisting of OH, -O-C₁-C₂₀ alkyl, -S-C(O)C₁-C₆ alkyl, and -C(O)OC₁-C₆ alkyl;
 and optionally R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or
 -O-C₂-C₆ alkynylene, such that where R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene,
 15 -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position.

[0159] In specific aspects of this embodiment, when Y and Y^a are each O, X^a and X^{a1} are
 each O, X^b and X^{b1} are each O, and X^c and X^{c1} are each OH or SH, X^d and X^{d1} are each O, R¹
 and R^{1a} are each H, R² is H, R⁶ and R^{6a} are each H, R⁷ and R^{7a} are each H, R⁸ and R^{8a} are each H,

and Base¹ and Base² are each selected from the group consisting of , , and

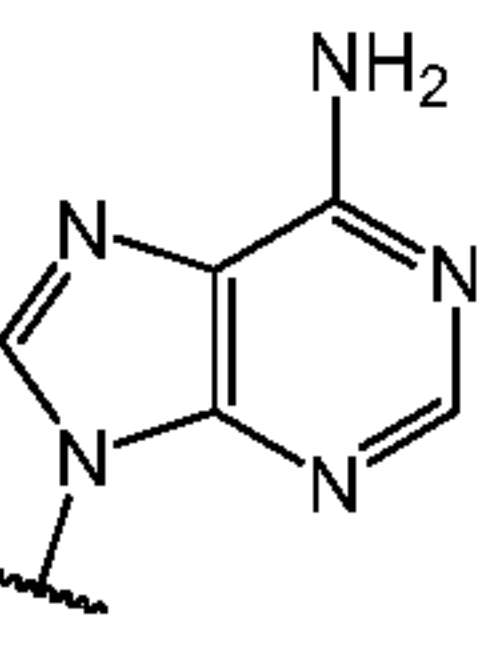
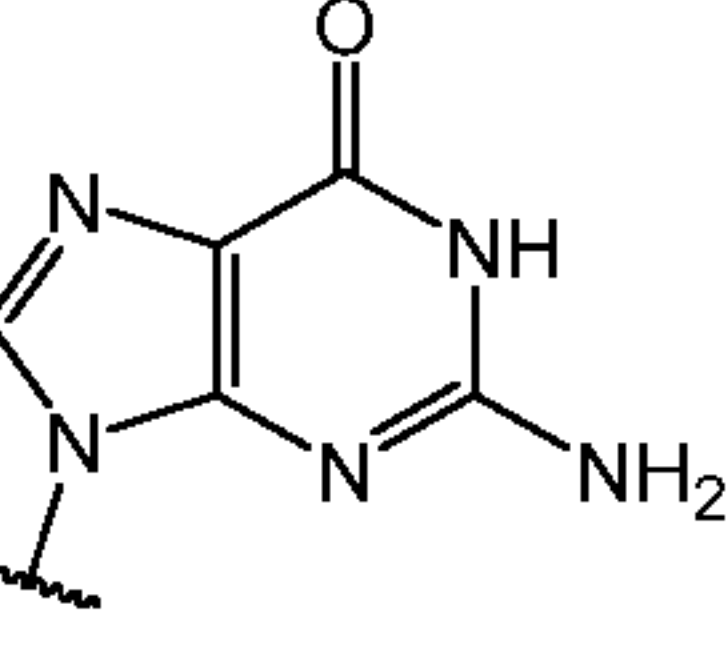
20 , R⁵ and R³ are not both selected from the group consisting of H, F and OH. That
 is, when Y and Y^a are each O, X^a and X^{a1} are each O, X^b and X^{b1} are each O, and X^c and X^{c1} are
 each OH or SH, X^d and X^{d1} are each O, R¹ and R^{1a} are each H, R² is H, R⁶ and R^{6a} are each H, R⁷
 and R^{7a} are each H, R⁸ and R^{8a} are each H, and Base¹ and Base² are each selected from the group

consisting of , , and , either only one of R⁵ and R³ is selected from the group consisting of H, F, and OH, or neither R⁵ and R³ is selected from the group consisting of H, F, and OH. In further specific instances of this aspect, when Y and Y^a are each O, X^a and X^{a1} are each O, X^b and X^{b1} are each O, and X^c and X^{c1} are each OH, X^d and X^{d1} are each O or S, R¹ and R^{1a} are each H, R² is H, R⁶ and R^{6a} are each H, R⁷ and R^{7a} are each H, R⁸ and R^{8a} are each H, and Base¹ and Base² are each selected from the group consisting of

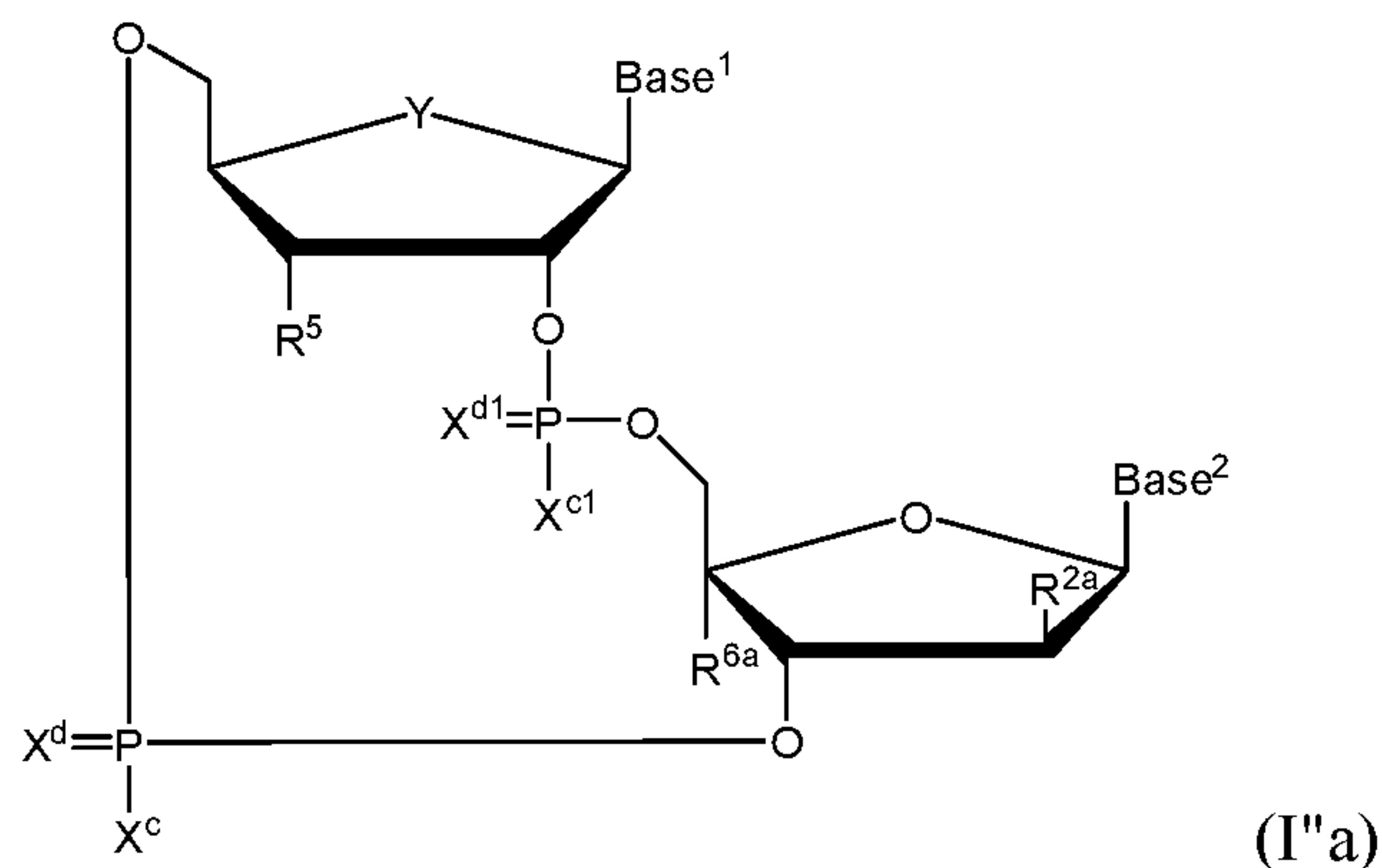


, R⁵ and R³ are not both selected from the group consisting of H, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, where said C₁-C₆ alkyl, C₂-C₆ alkenyl and C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I and OH.

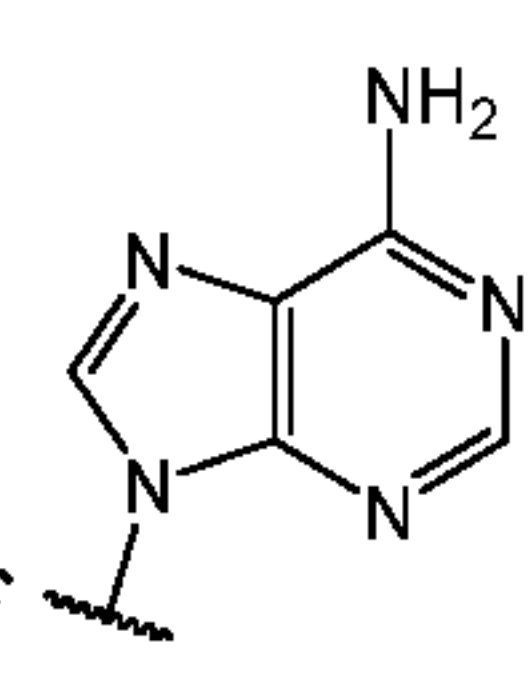
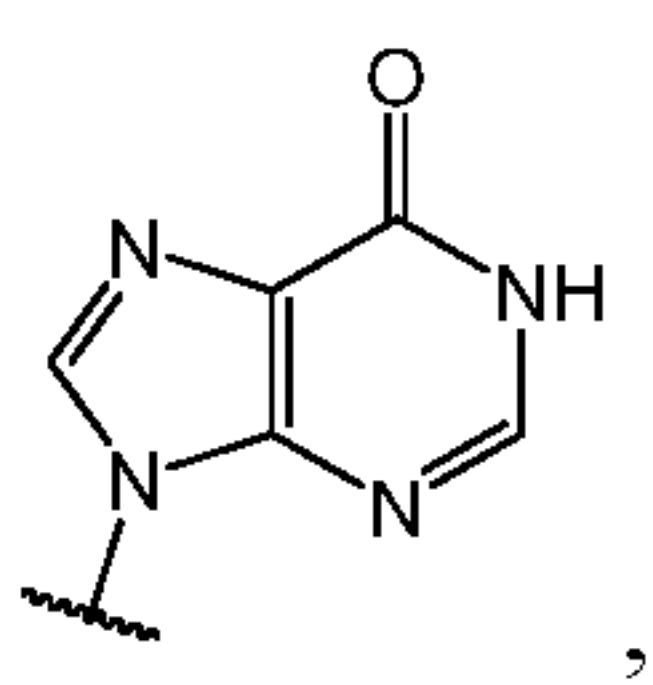
[0160] In further aspects of this embodiment, when Base¹ and Base² are each selected

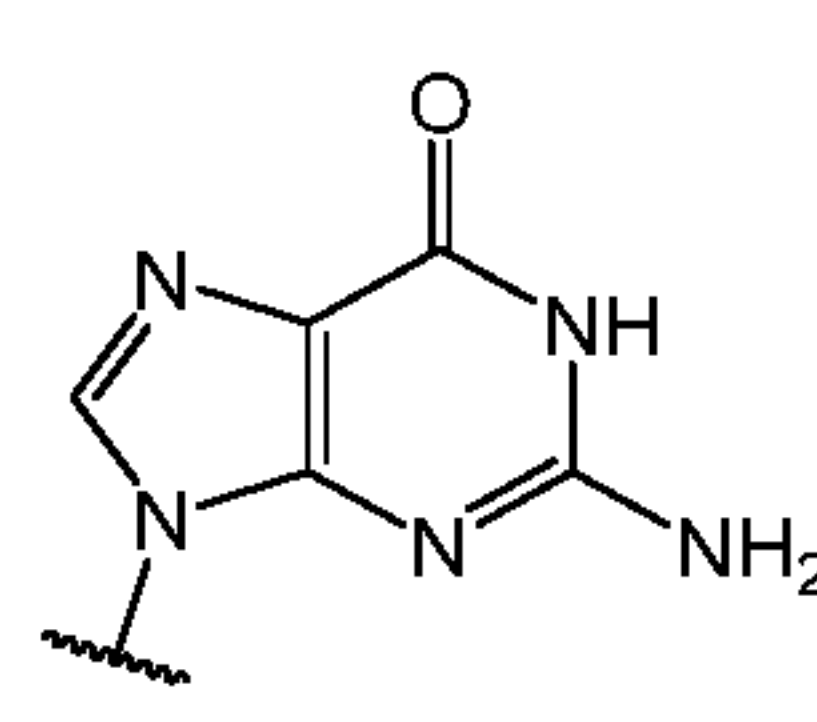
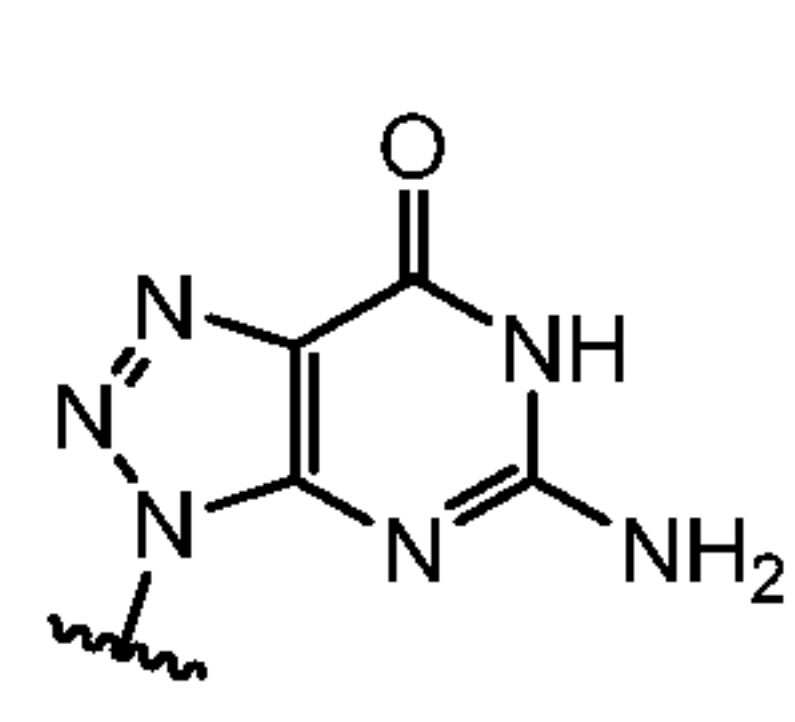
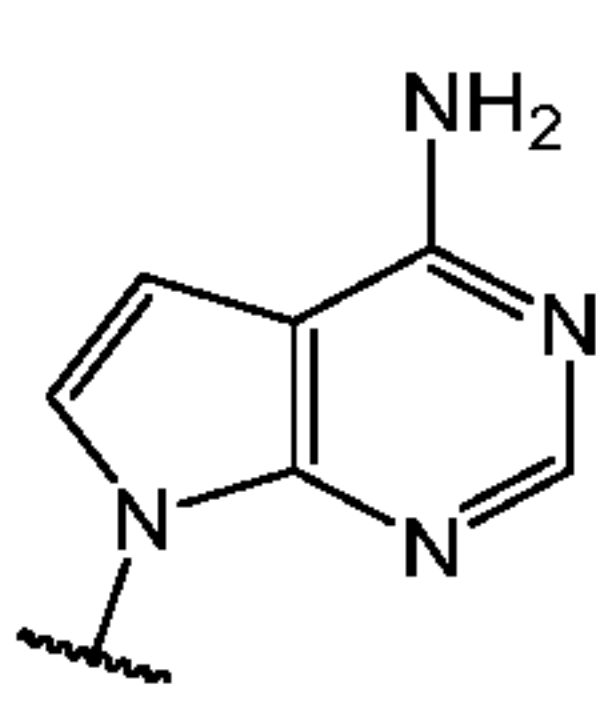
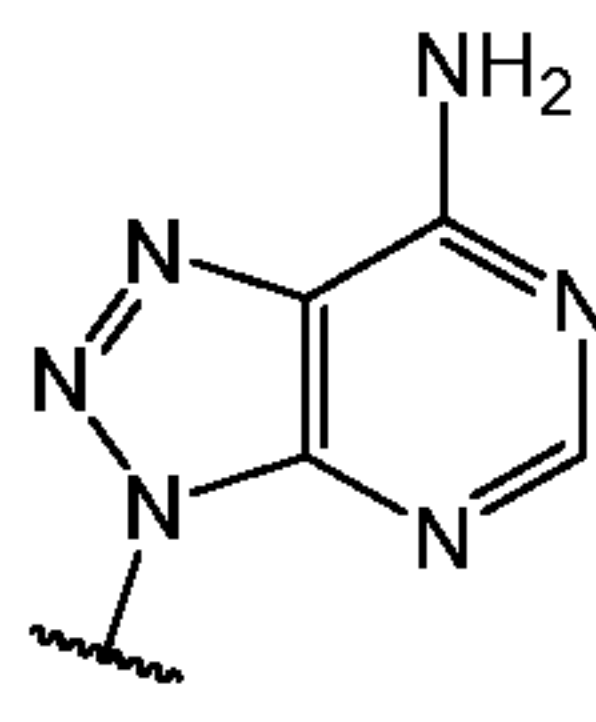
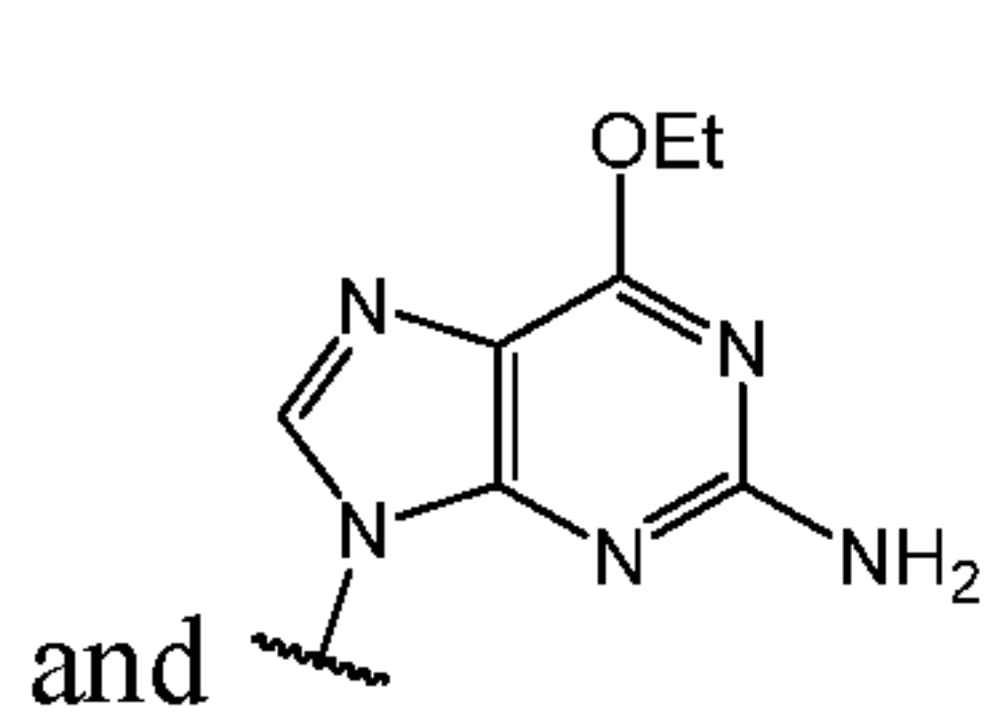
from the group consisting of  and , and R^{2a} is F and R⁵ is F, at least one of X^c and X^{c1} is SR⁹.

[0161] In aspects of this embodiment, the compound of formula (I'') is a compound of formula (I'a):



or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein Base¹ and

Base² are each independently selected from the group consisting of , ,

, , , , and ; Y is selected

from the group consisting of -O- and -S-; X^c and X^{c1} are each independently selected from the

5 group consisting of OR⁹ and SR⁹; X^d and X^{d1} are each independently selected from the group

consisting of O and S; R^{2a} is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆

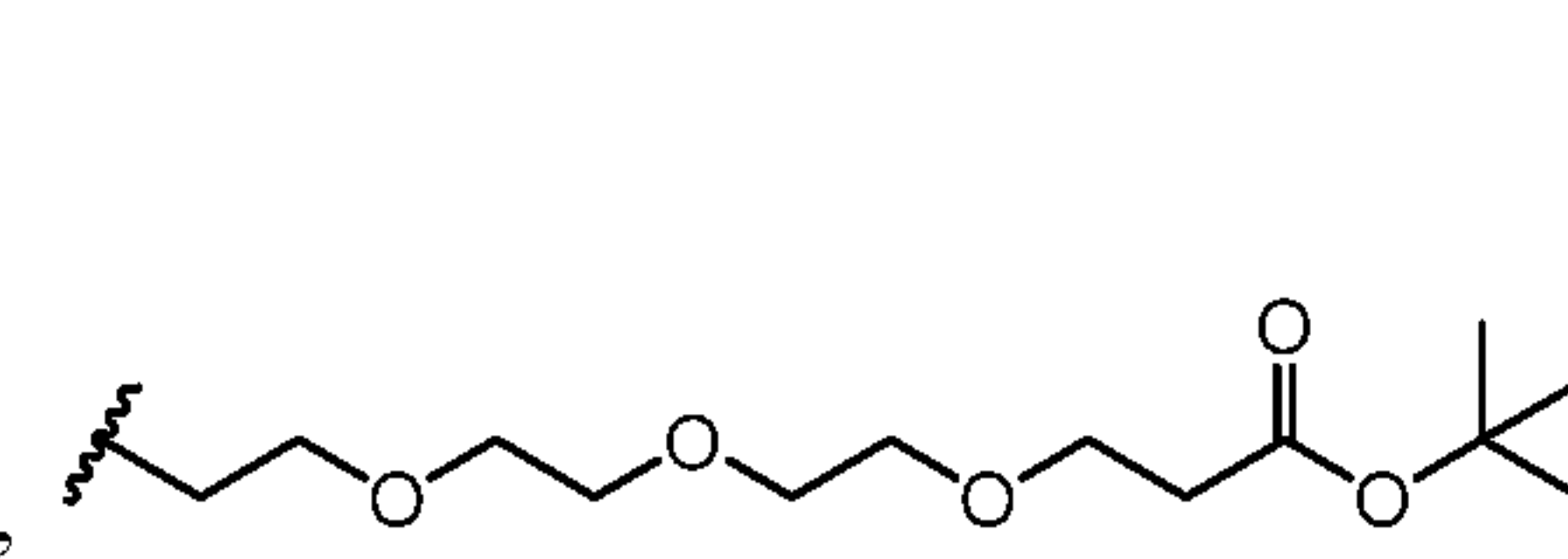
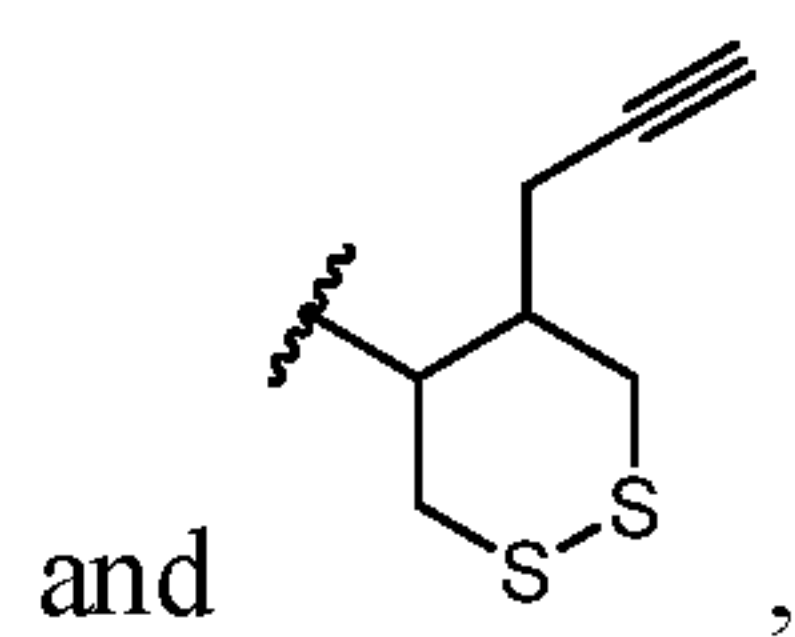
haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl; R⁵ is selected from the

group consisting of H, F, Cl, Br, I, OH, CN, NH₂, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl,

10 C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and

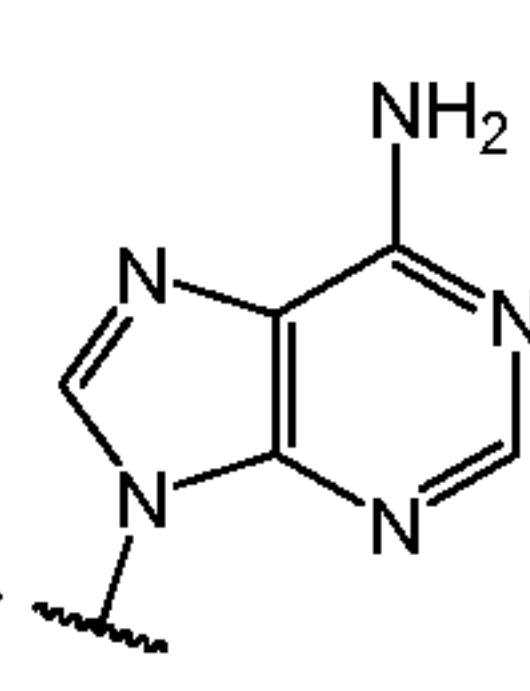
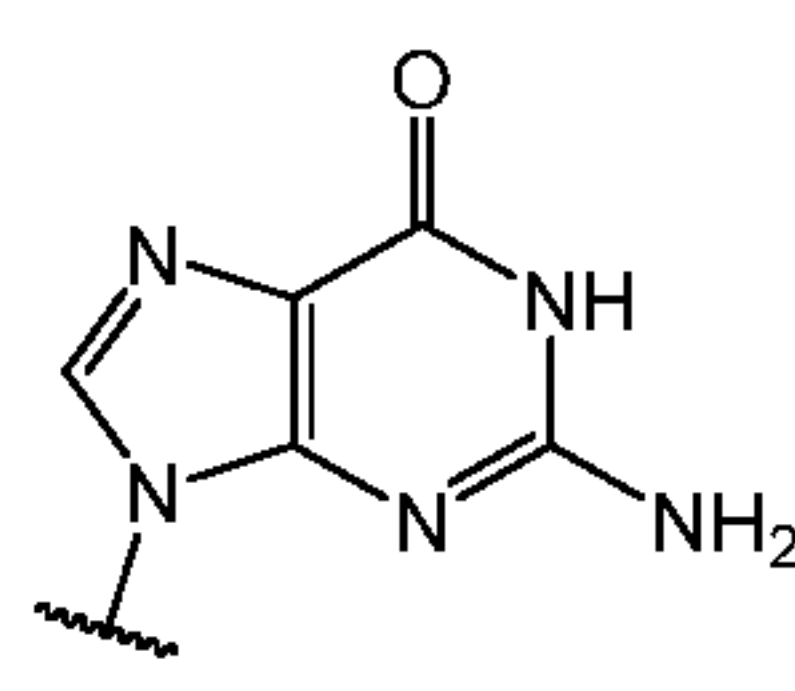
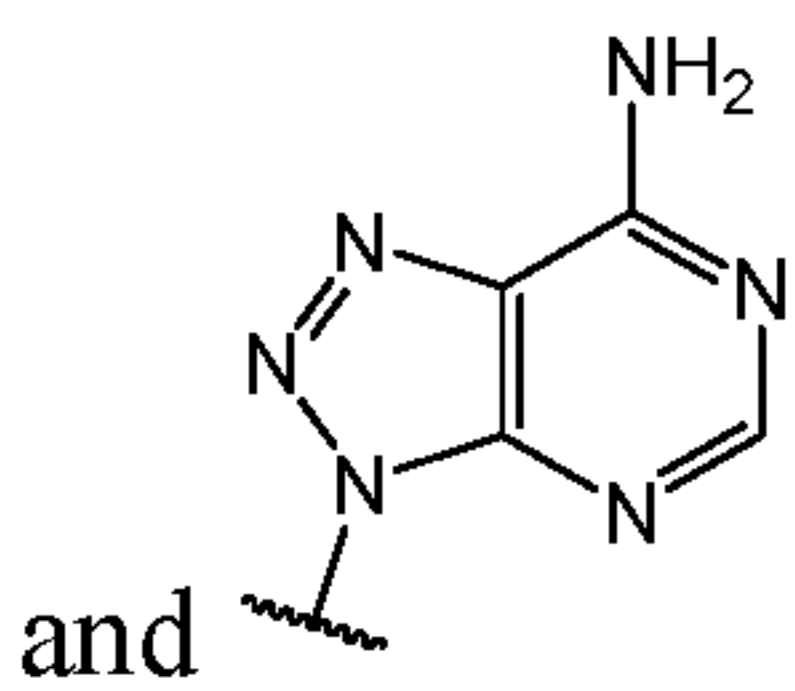
-O-C₂-C₆ alkynyl; R^{6a} is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl,

-O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl; and each R⁹ is independently selected

from the group consisting of H, C₁-C₂₀ alkyl, , and ,

15 where each R⁹ C₁-C₂₀ alkyl is optionally substituted by 0 to 3 substituents independently selected from the group consisting of OH, -O-C₁-C₂₀ alkyl, -S-C(O)C₁-C₆ alkyl, and -C(O)OC₁-C₆ alkyl.

In instances of this aspect, Base¹ and Base² are each independently selected from the group

consisting of , , and ; Y is selected from the group

consisting of -O- and -S-; X^c and X^{c1} are each independently selected from the group consisting

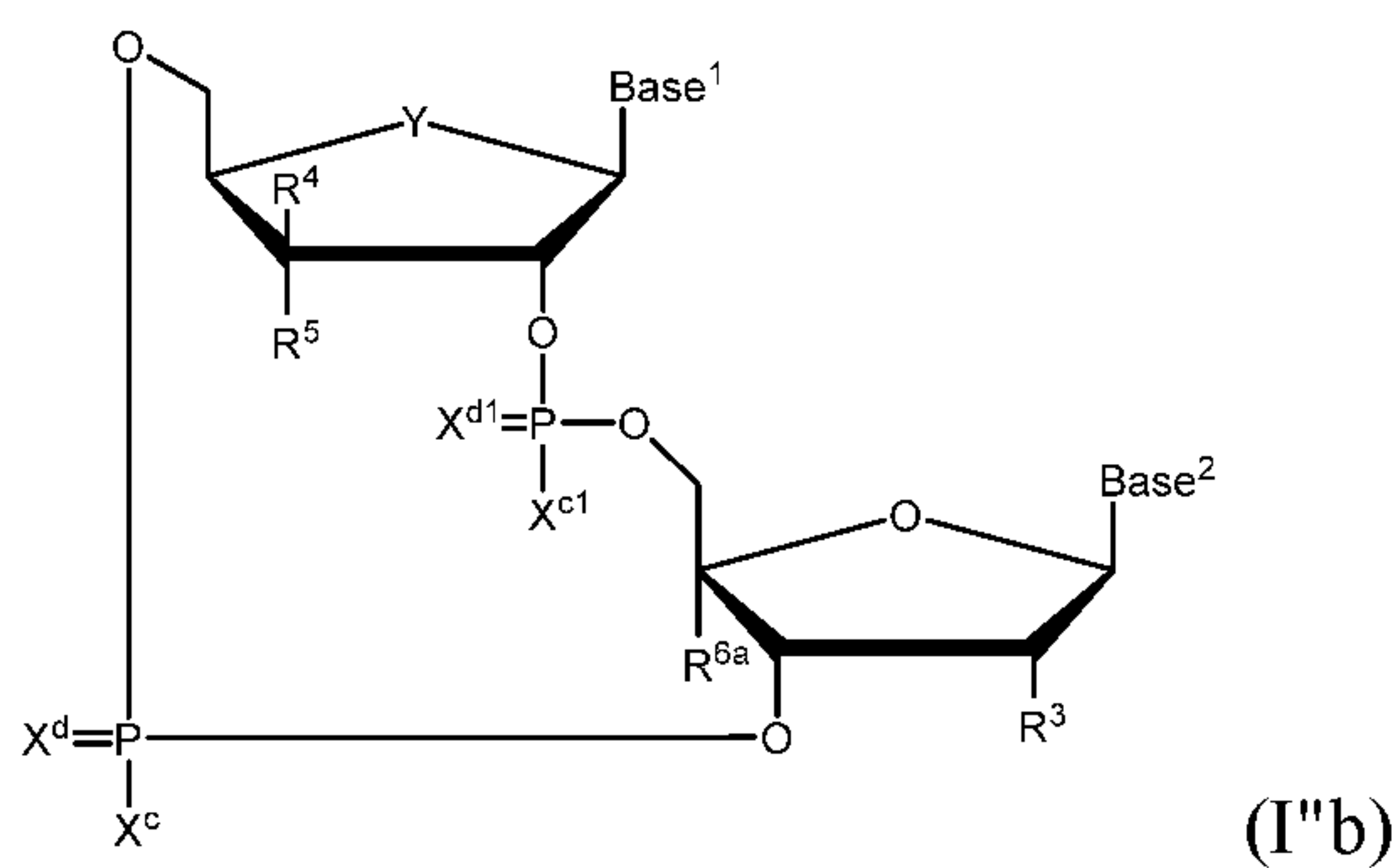
20 of OR⁹ and SR⁹; X^d and X^{d1} are each independently selected from the group consisting of O and

S; R^{2a} is F; R⁵ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, NH₂, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl,

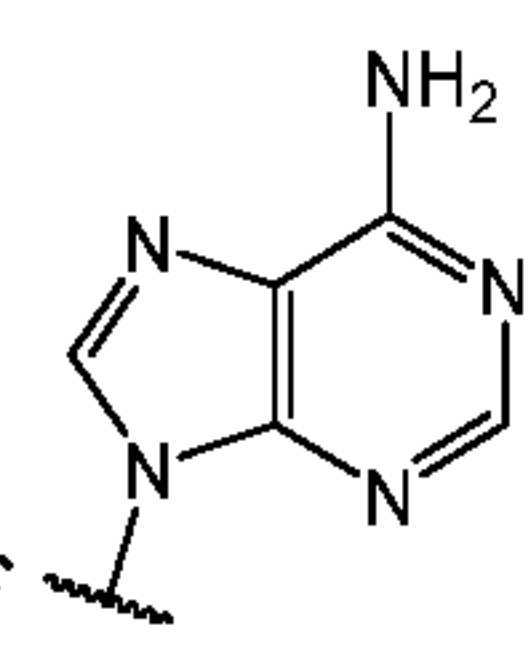
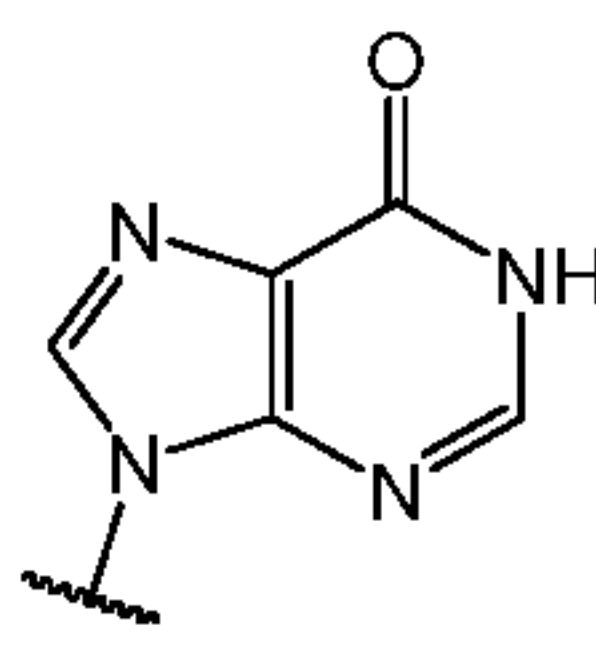
-O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl; R^{6a} is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl; and each R⁹ is independently H.

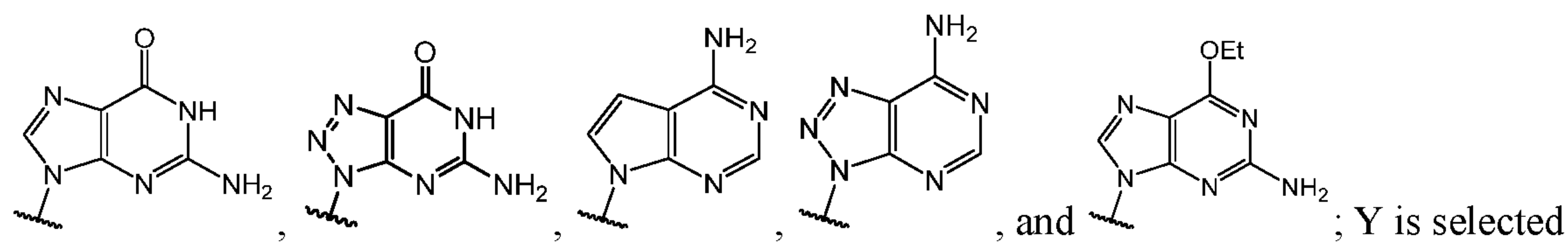
5 [0162] In aspects of this embodiment, the compound of formula (I'') is a compound wherein R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position.

10 [0163] In aspects of this embodiment, the compound of formula (I'') is a compound of formula (I''b):



or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein Base¹ and

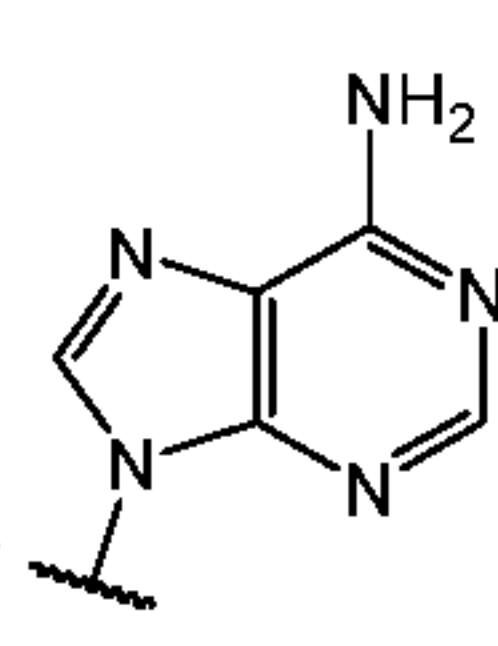
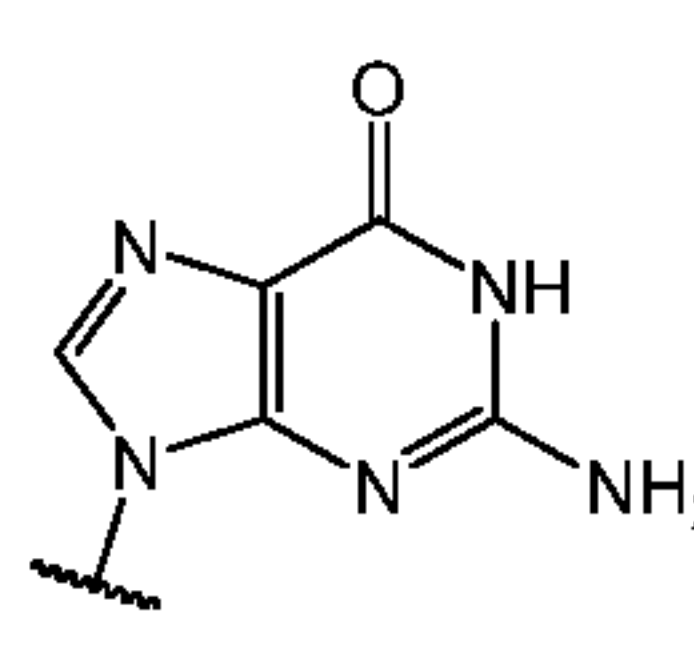
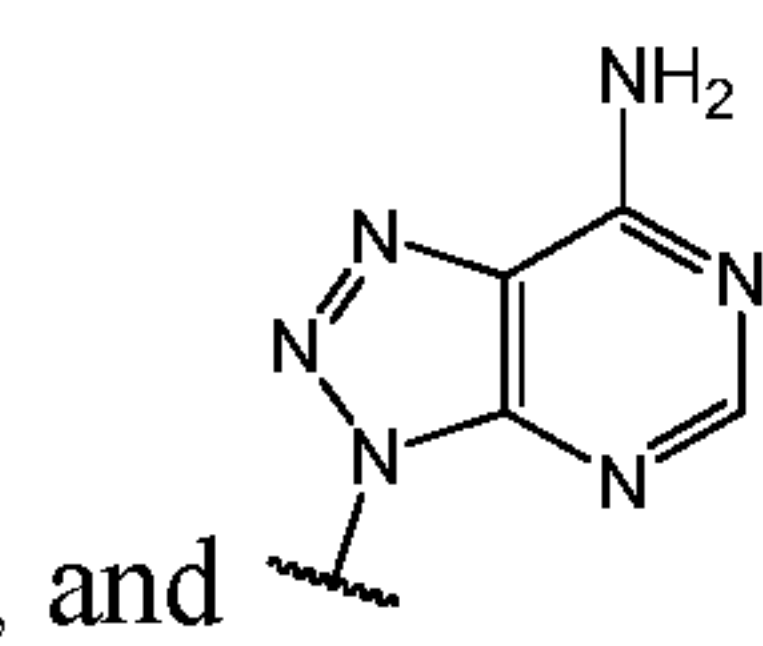
Base² are each independently selected from the group consisting of , , and



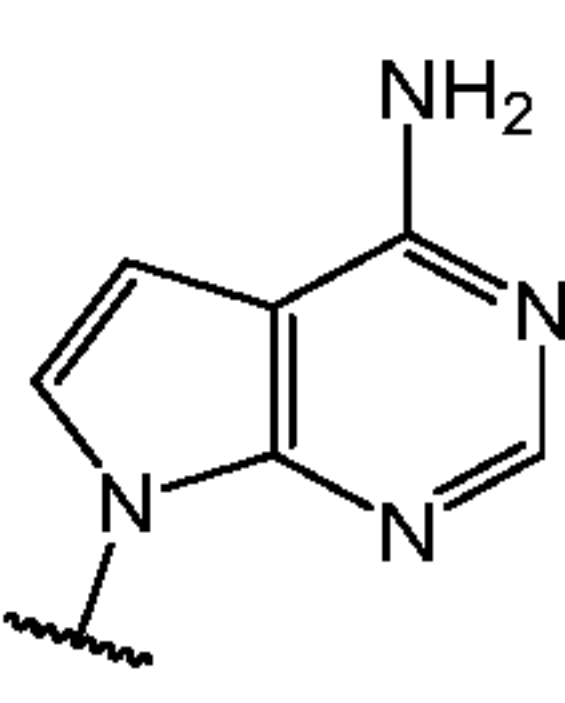
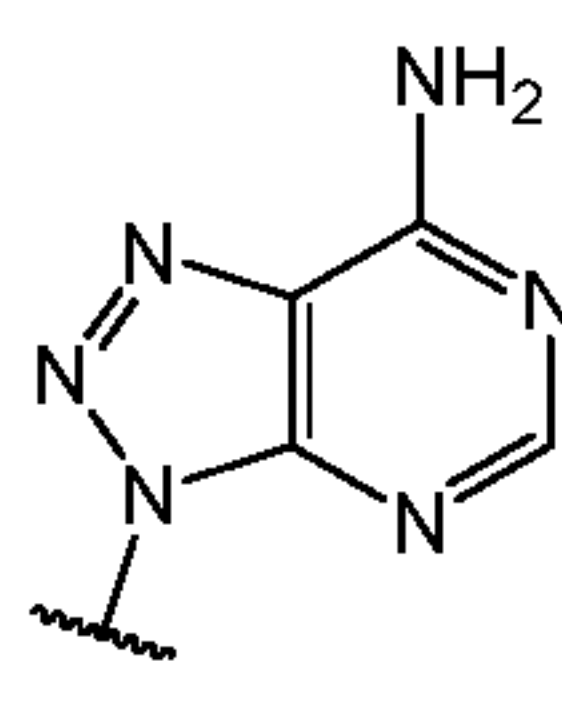
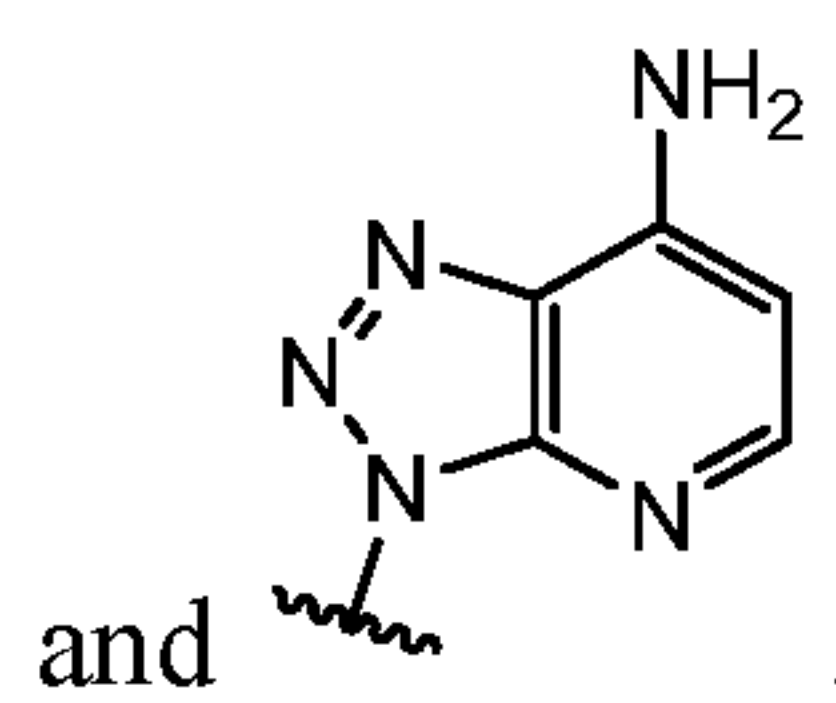
15 from the group consisting of -O- and -S-; X^c and X^{c1} are each independently selected from the group consisting of OR⁹ and SR⁹; X^d and X^{d1} are each independently selected from the group consisting of O and S; R³ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl; R⁴ is selected from the

20 group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and

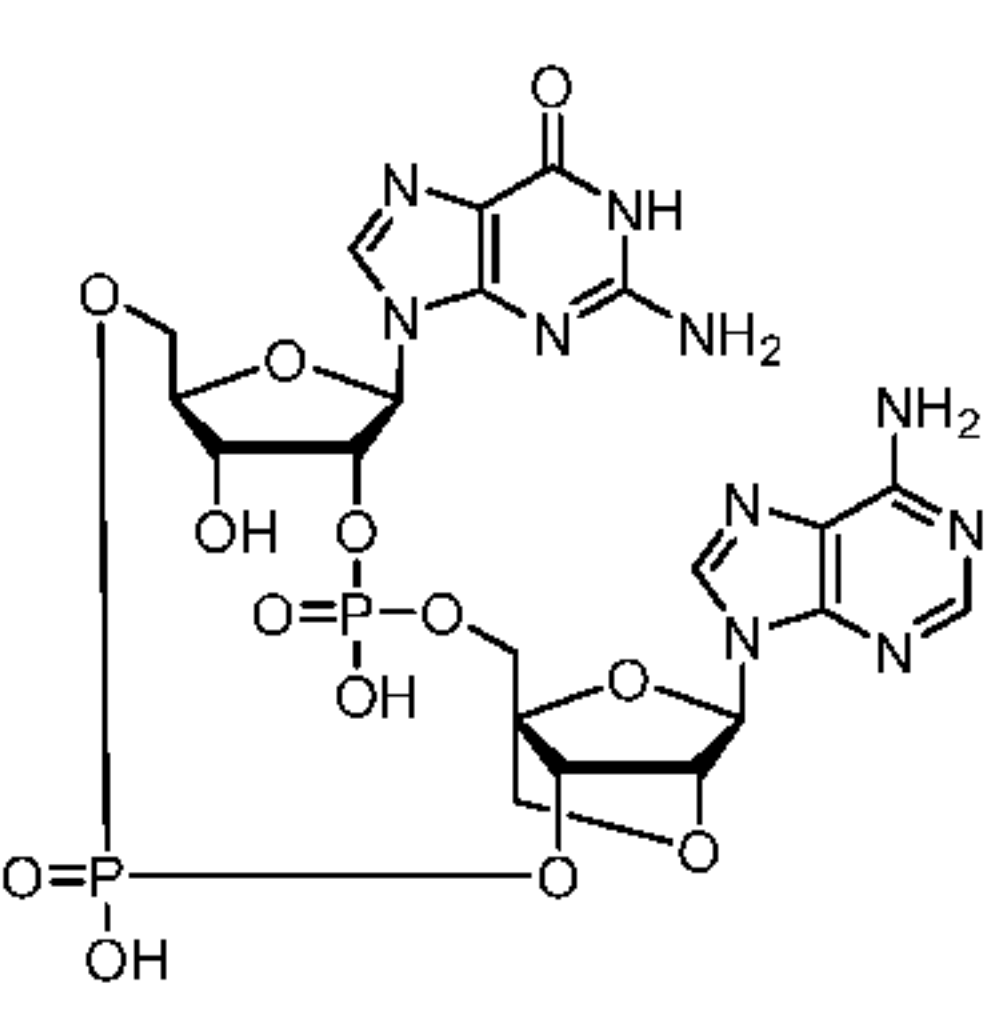
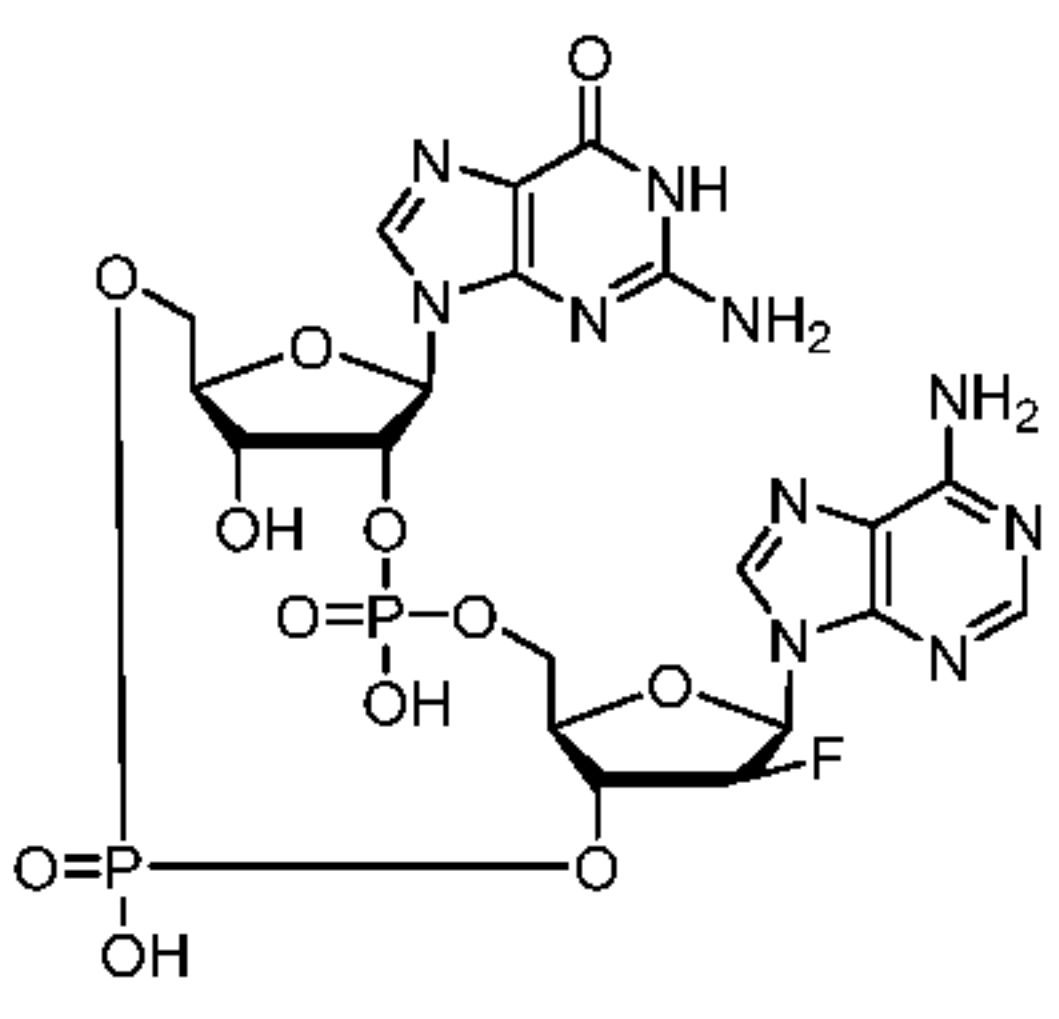
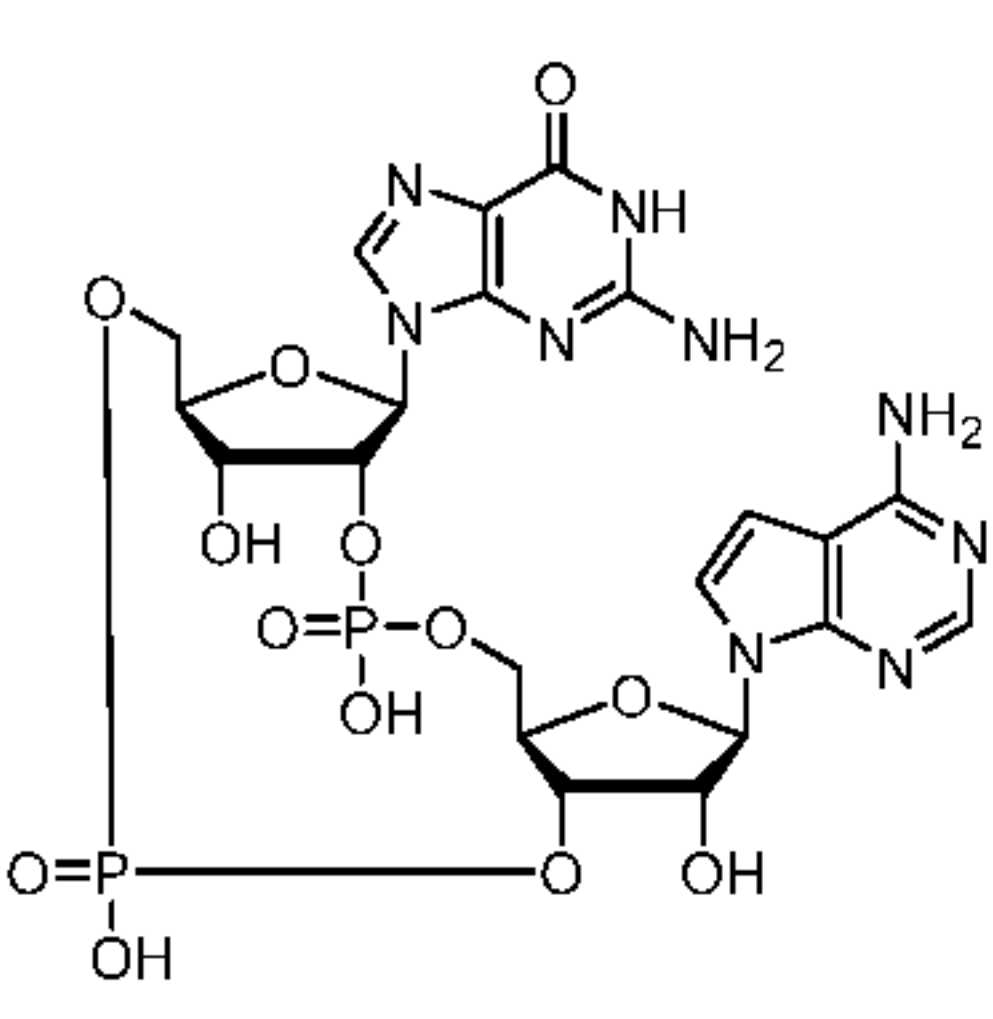
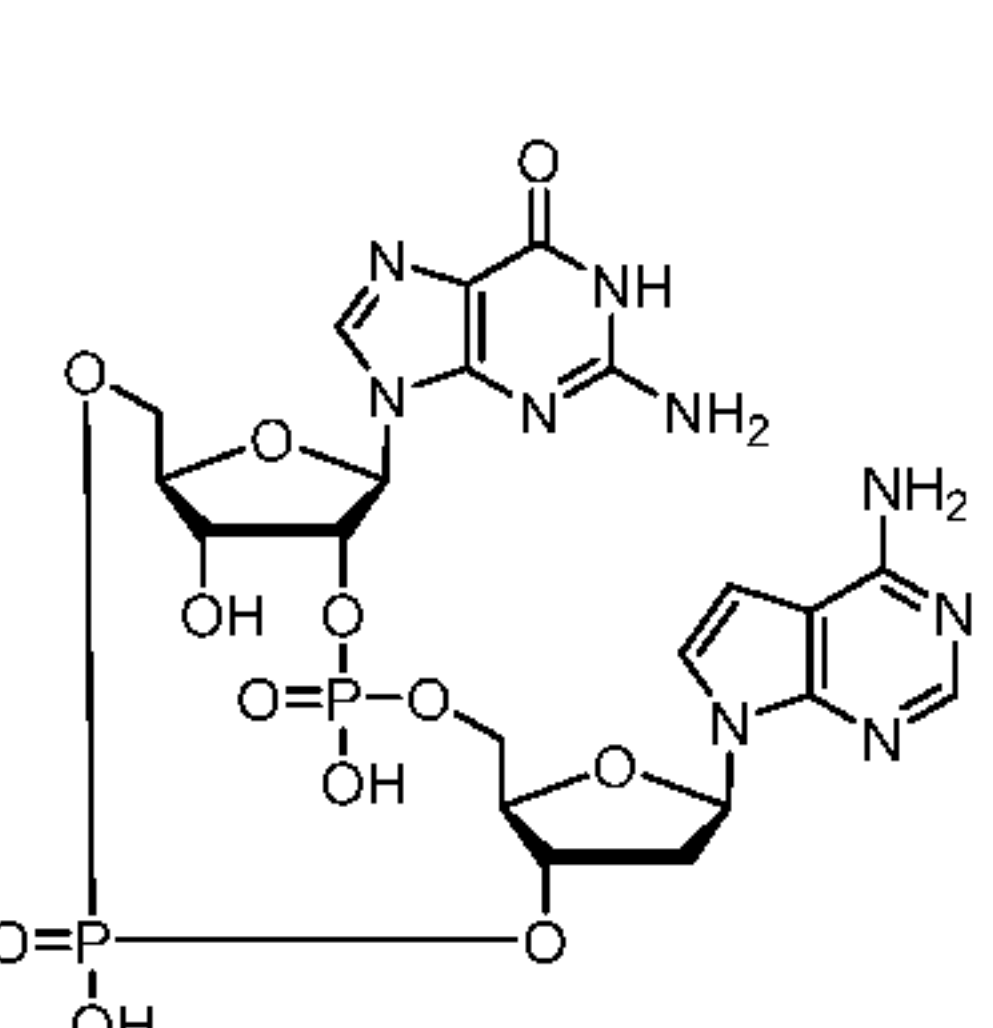
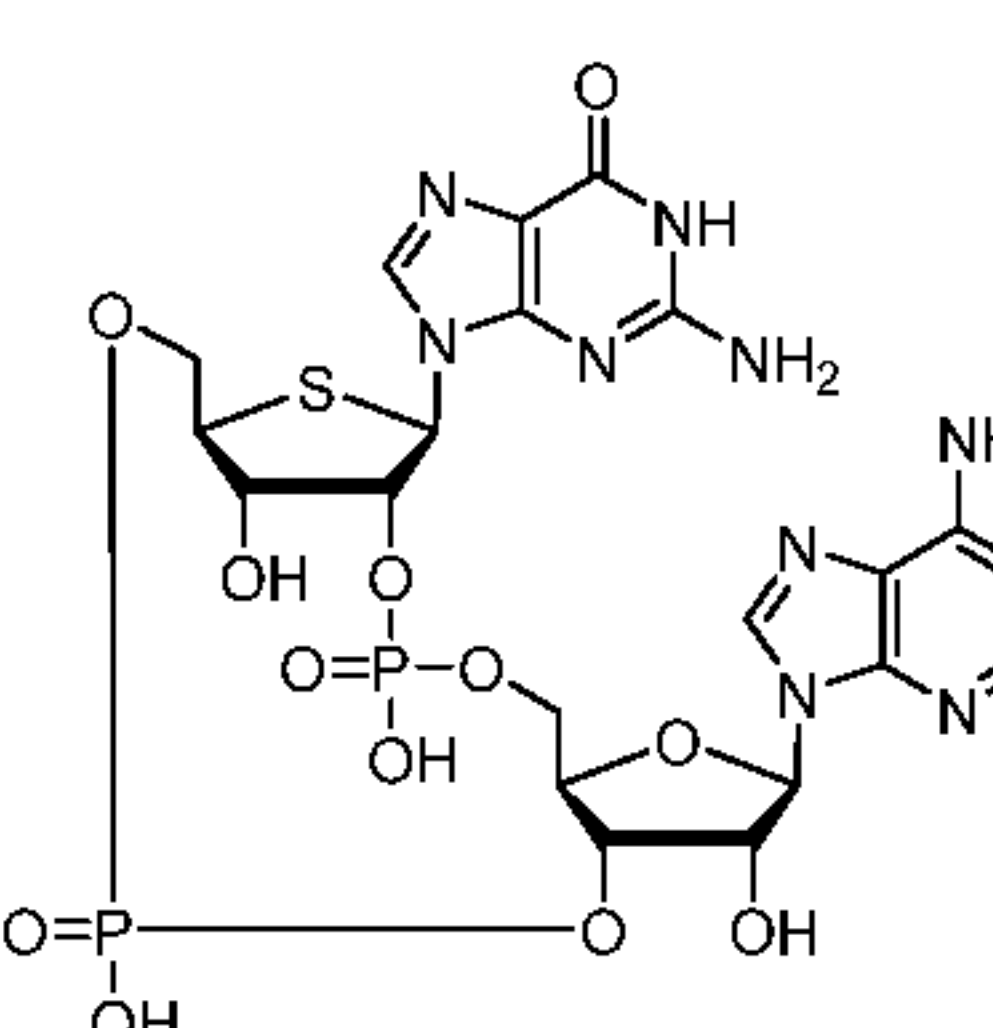
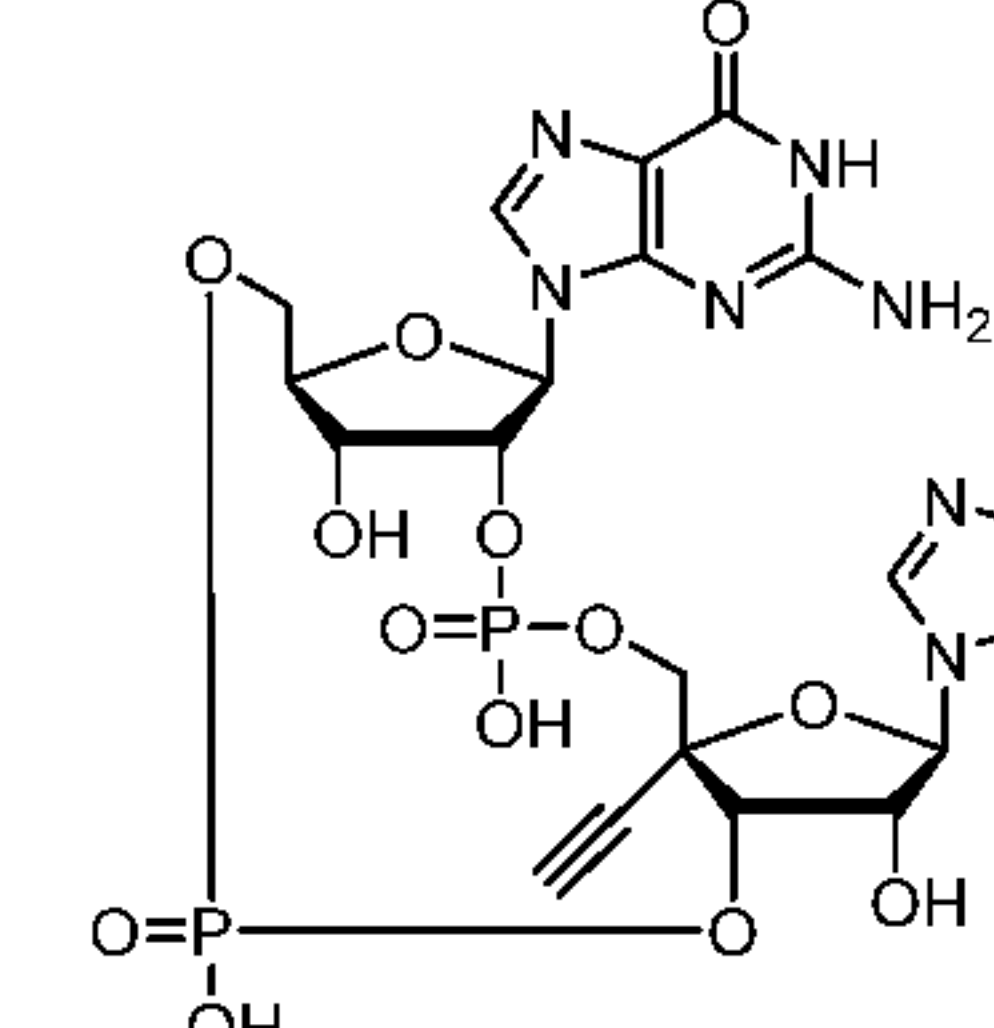
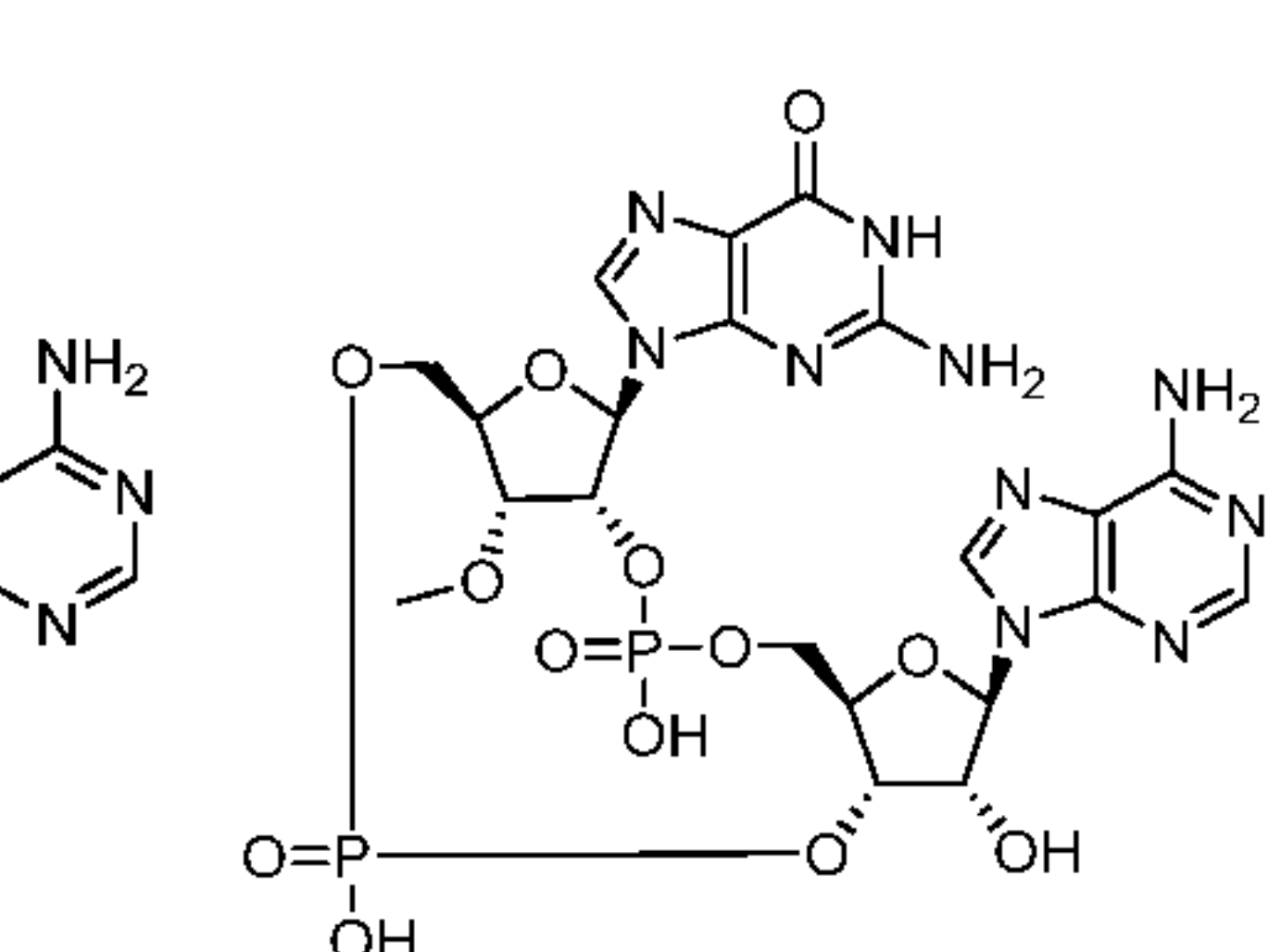
-O-C₂-C₆ alkynyl; R⁵ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, NH₂, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl; R^{6a} is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl; each R⁹ is independently H; and R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position. In instances of this aspect, Base¹ and Base² are each independently

selected from the group consisting of , , and .

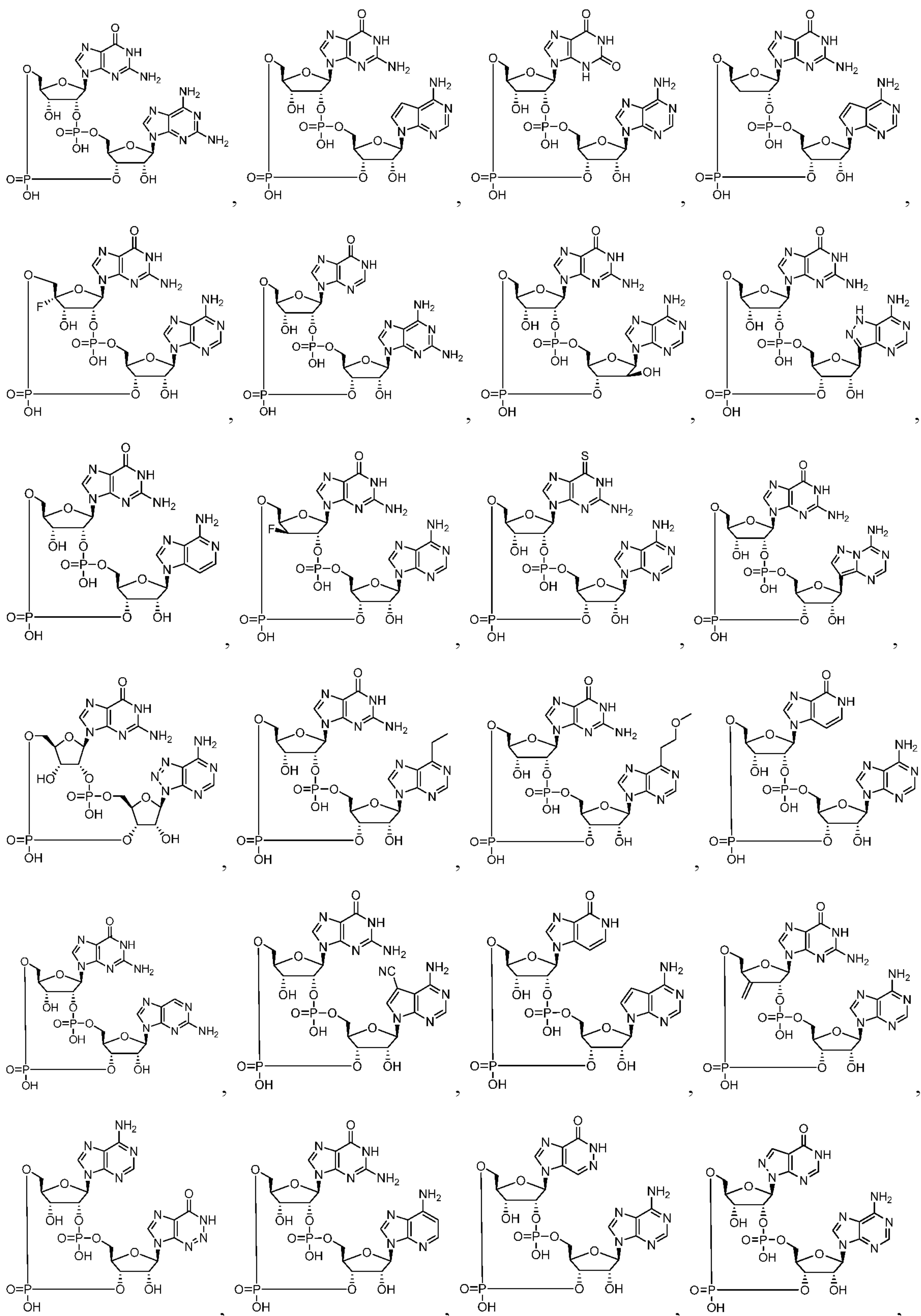
[0164] In aspects of this embodiment, the compound of formula (I'') is a compound wherein at least one of Base¹ and Base² are each independently selected from the group

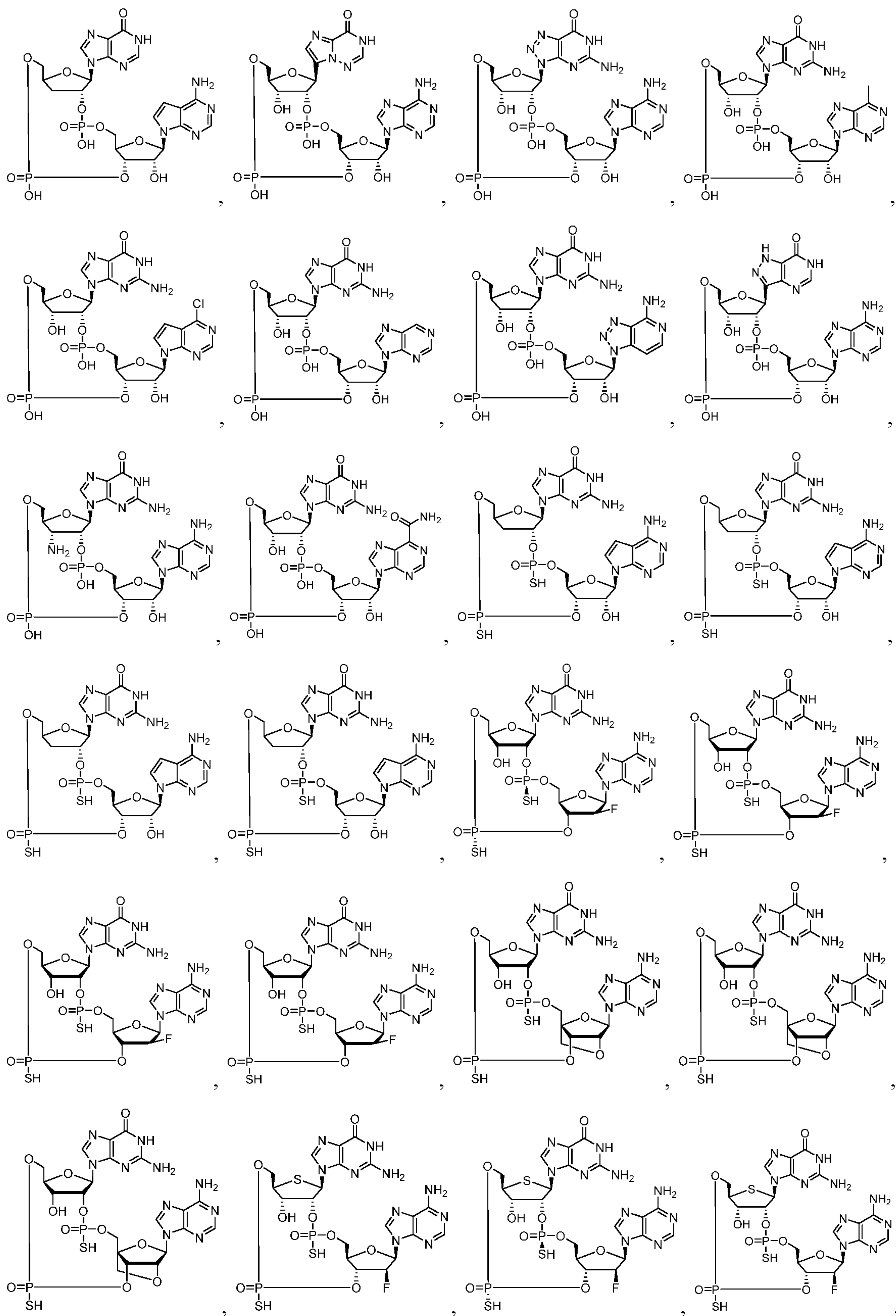
consisting of , , and .

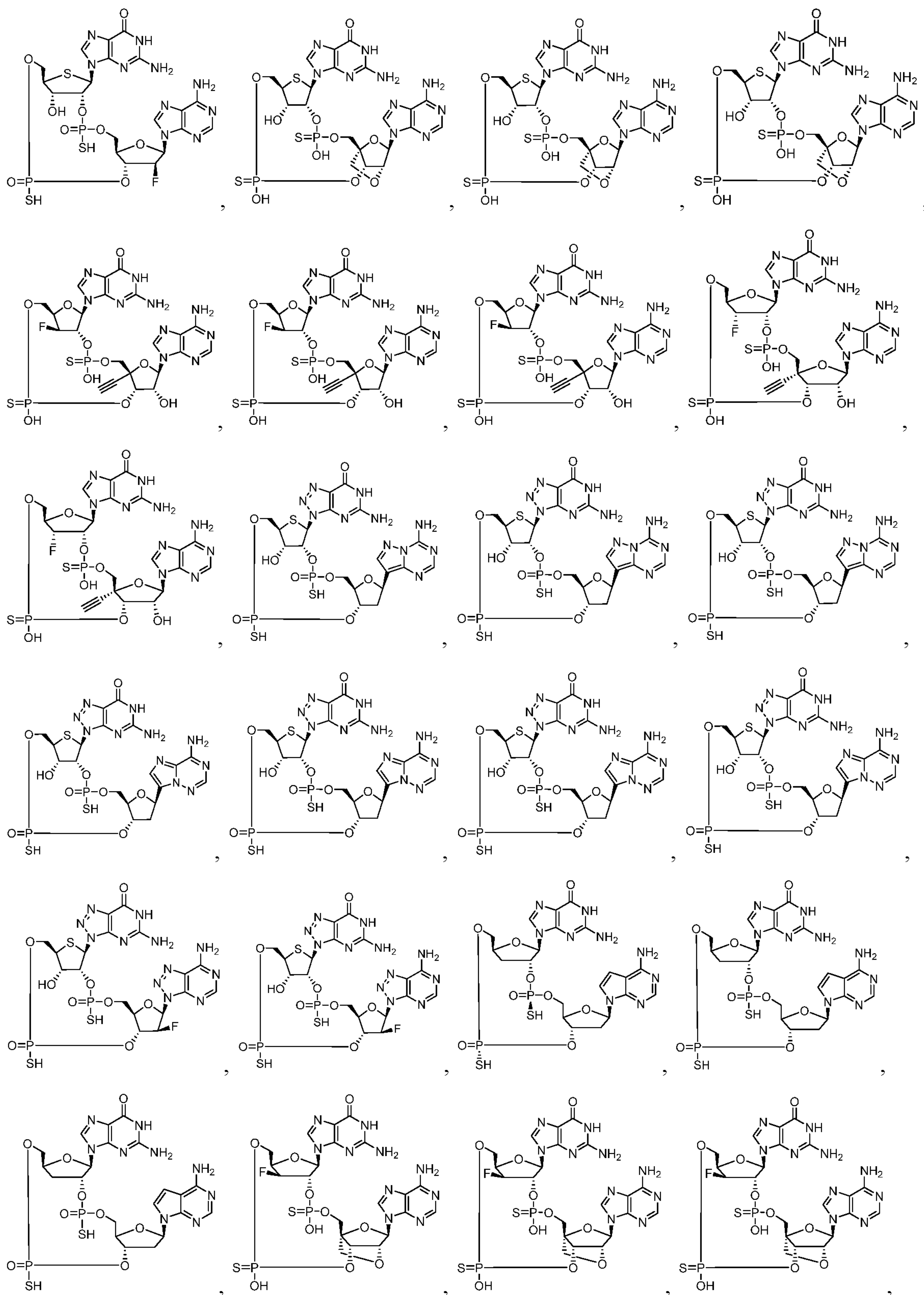
[0165] In an additional embodiment, the CDN STING agonist is selected from the group

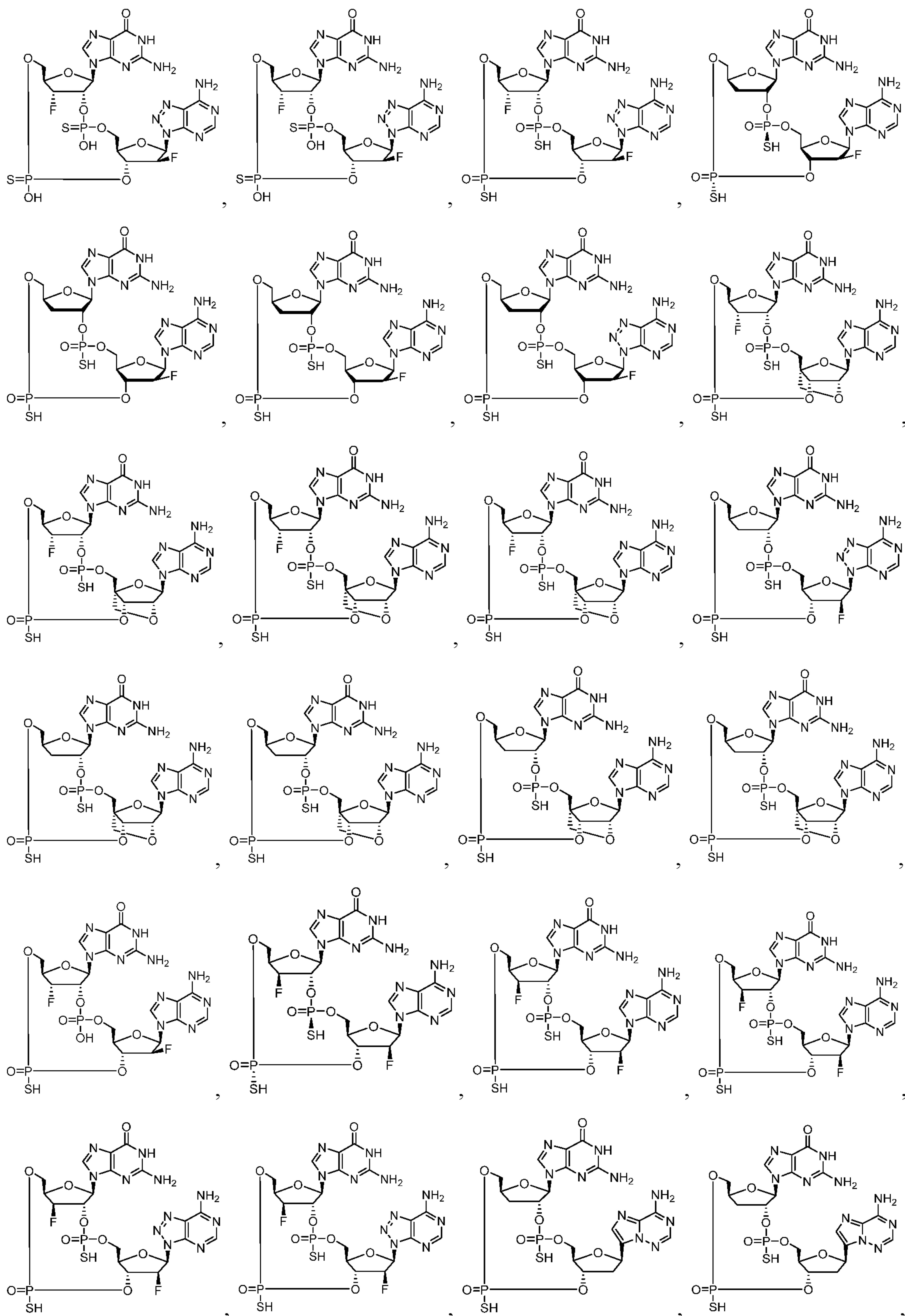
consisting of , , , , , , and .

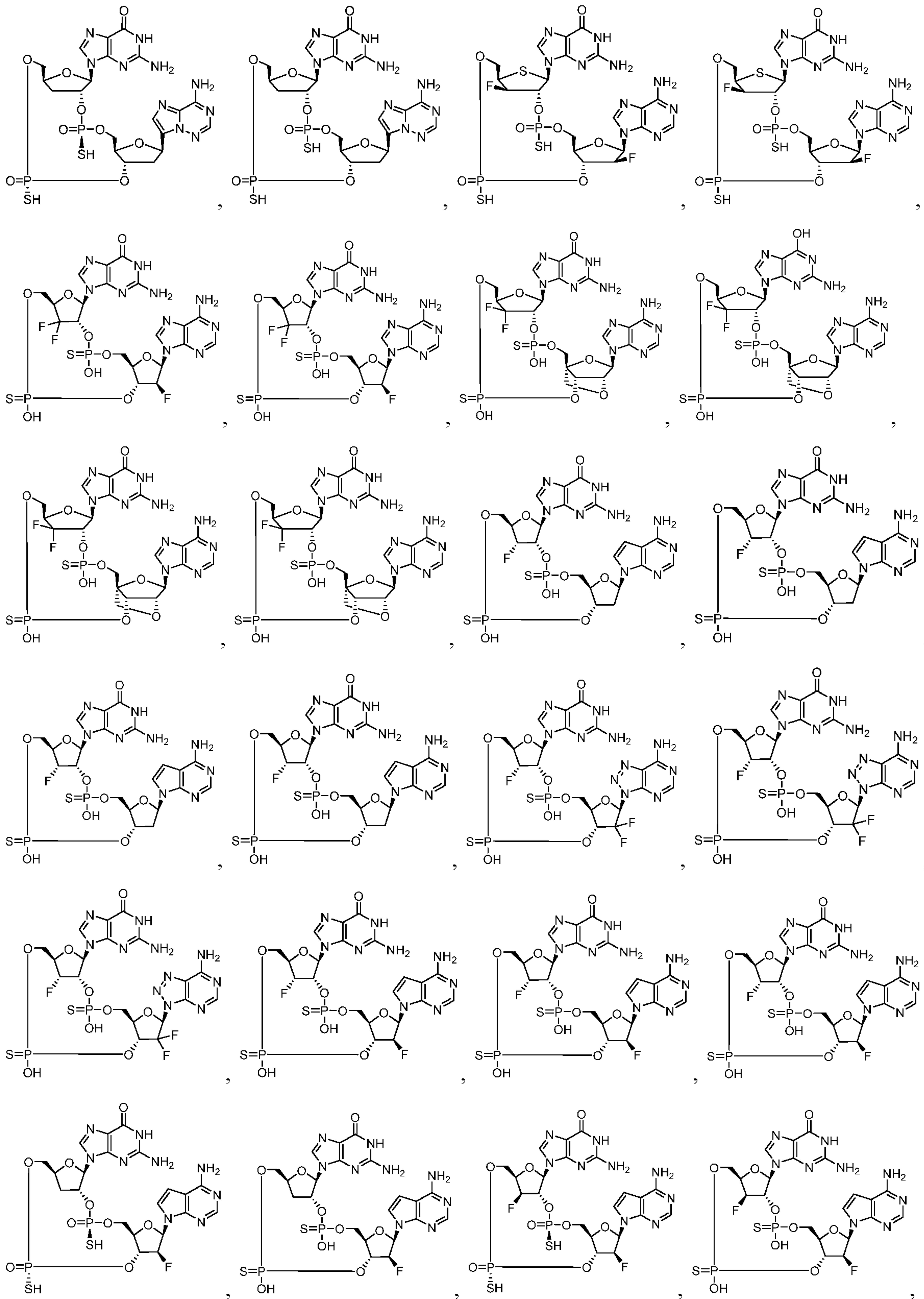


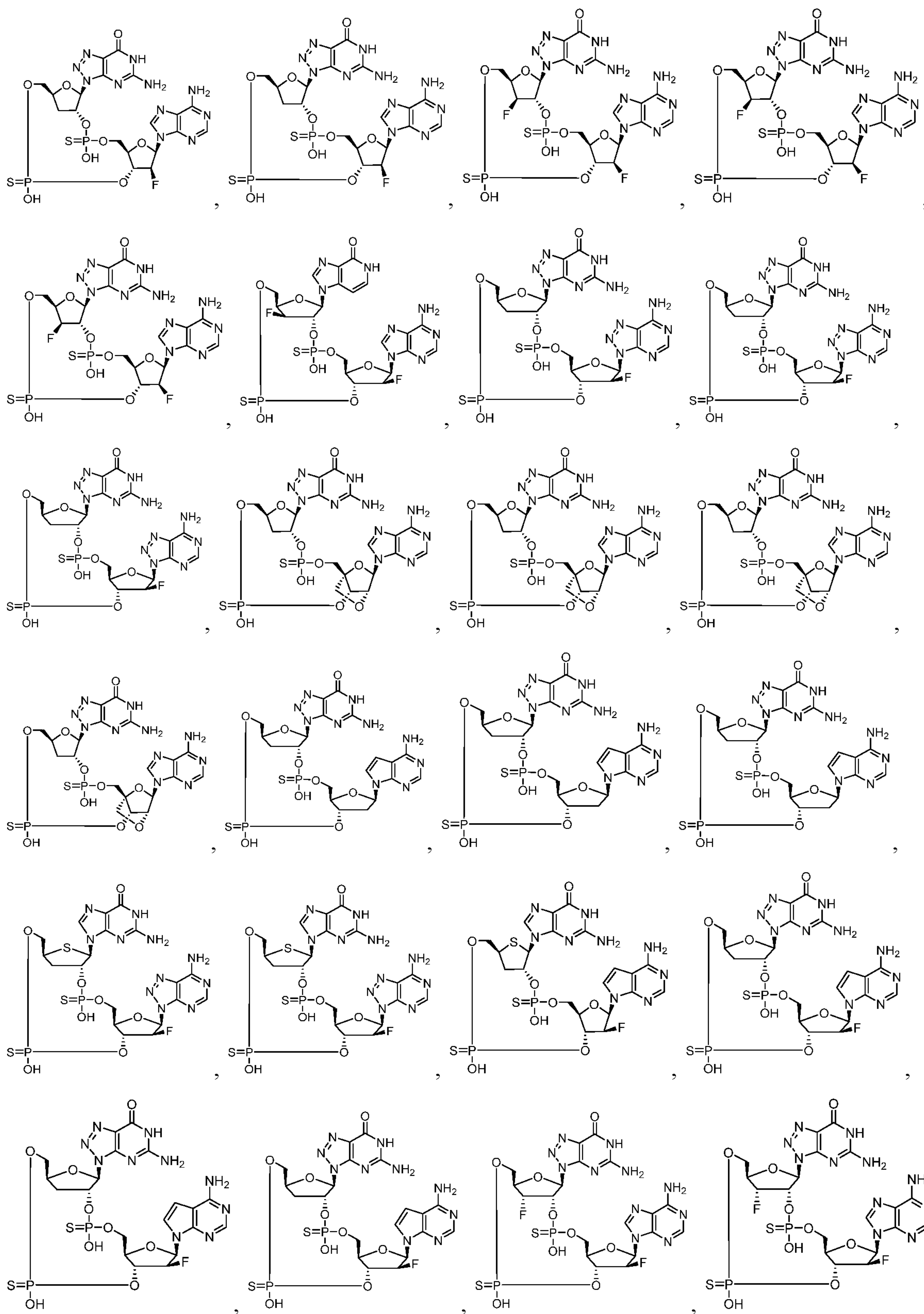


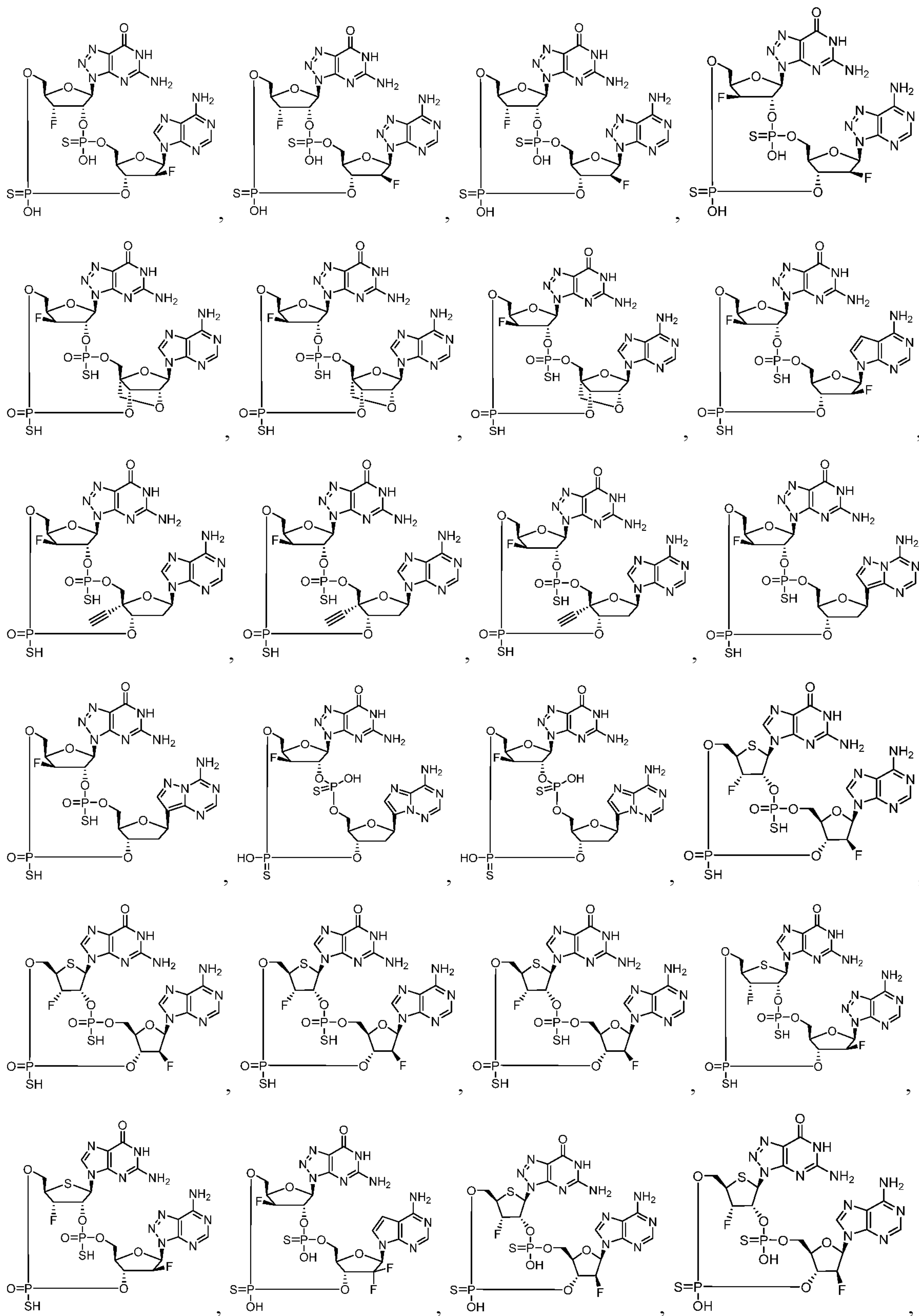


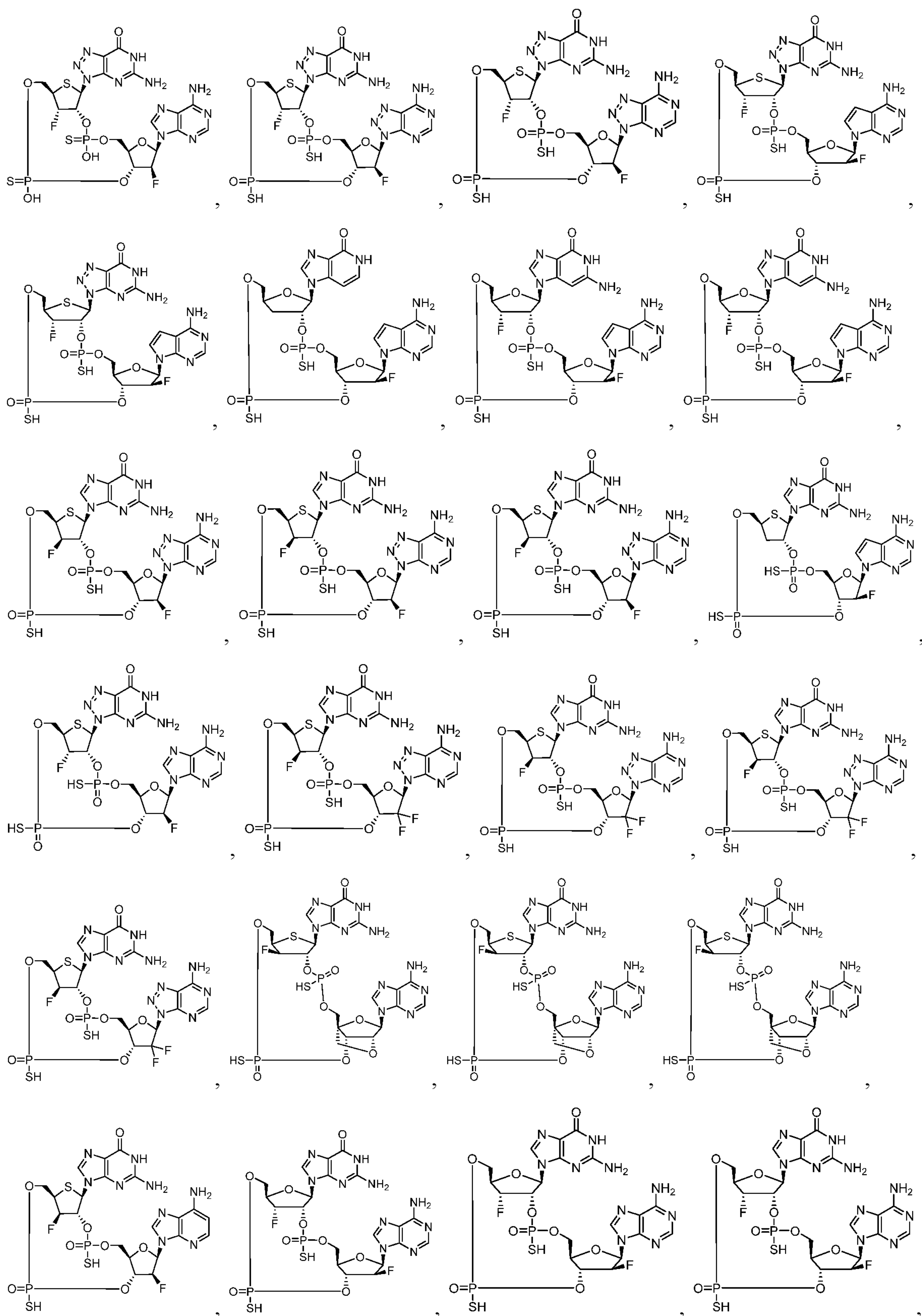


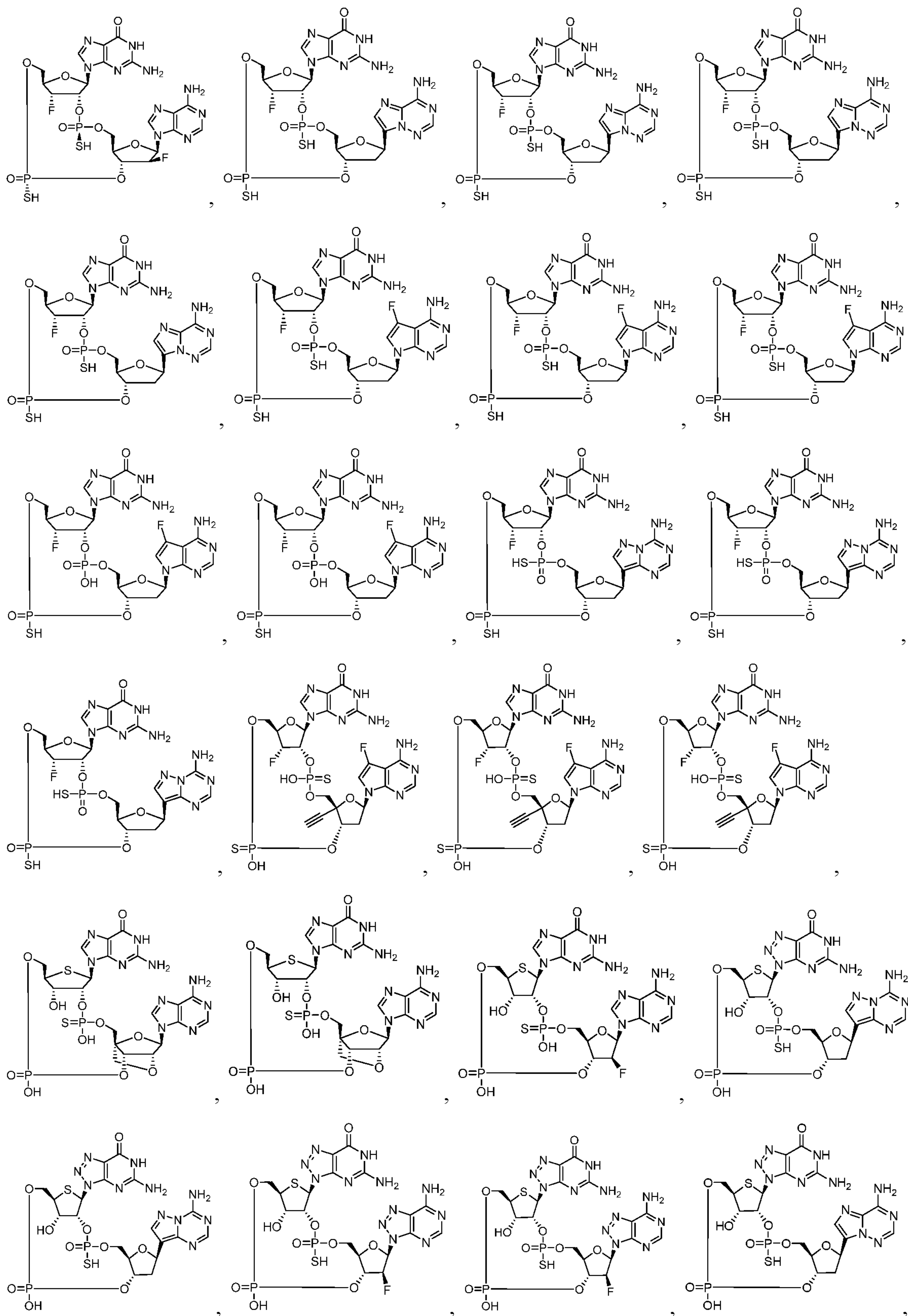


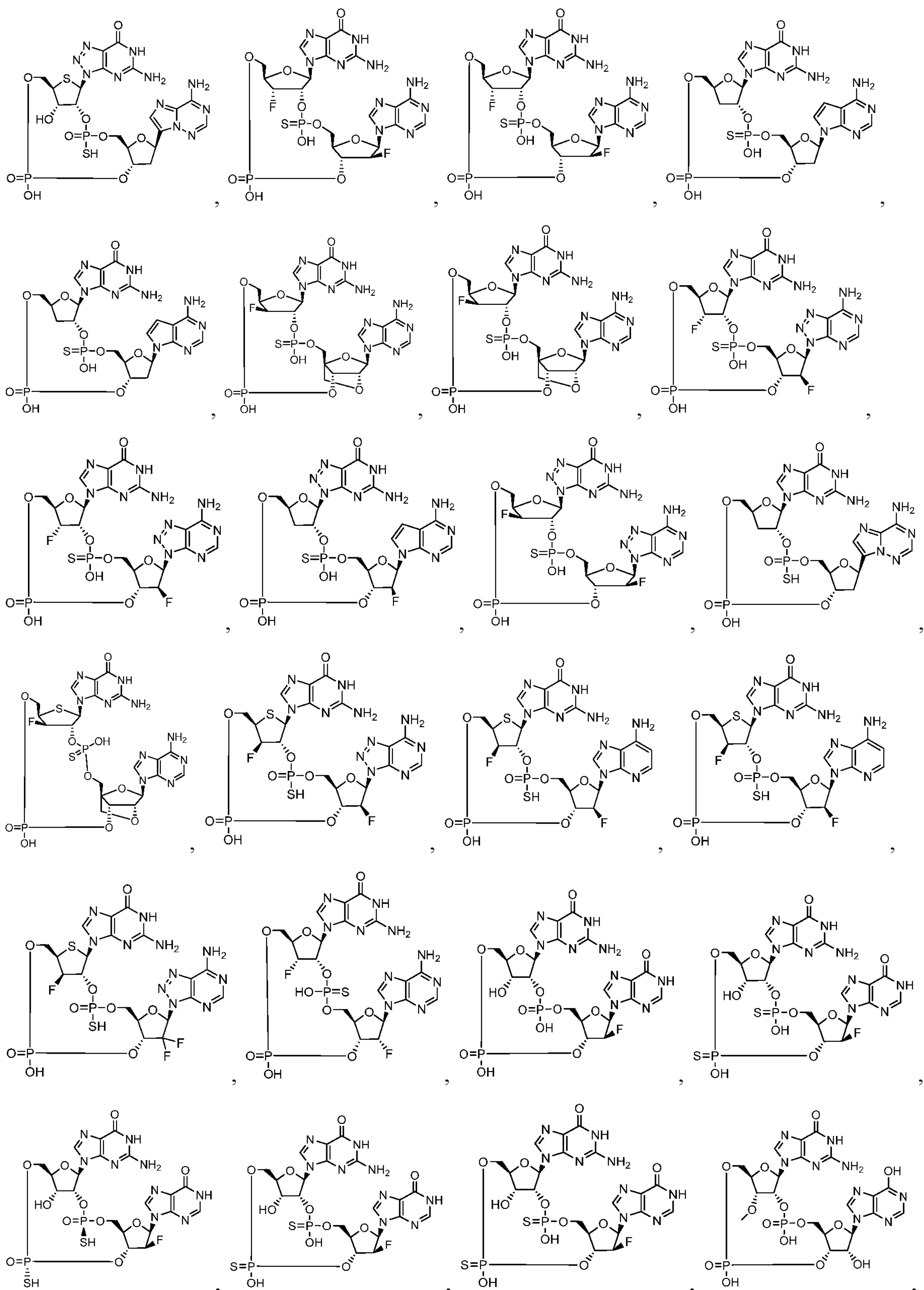


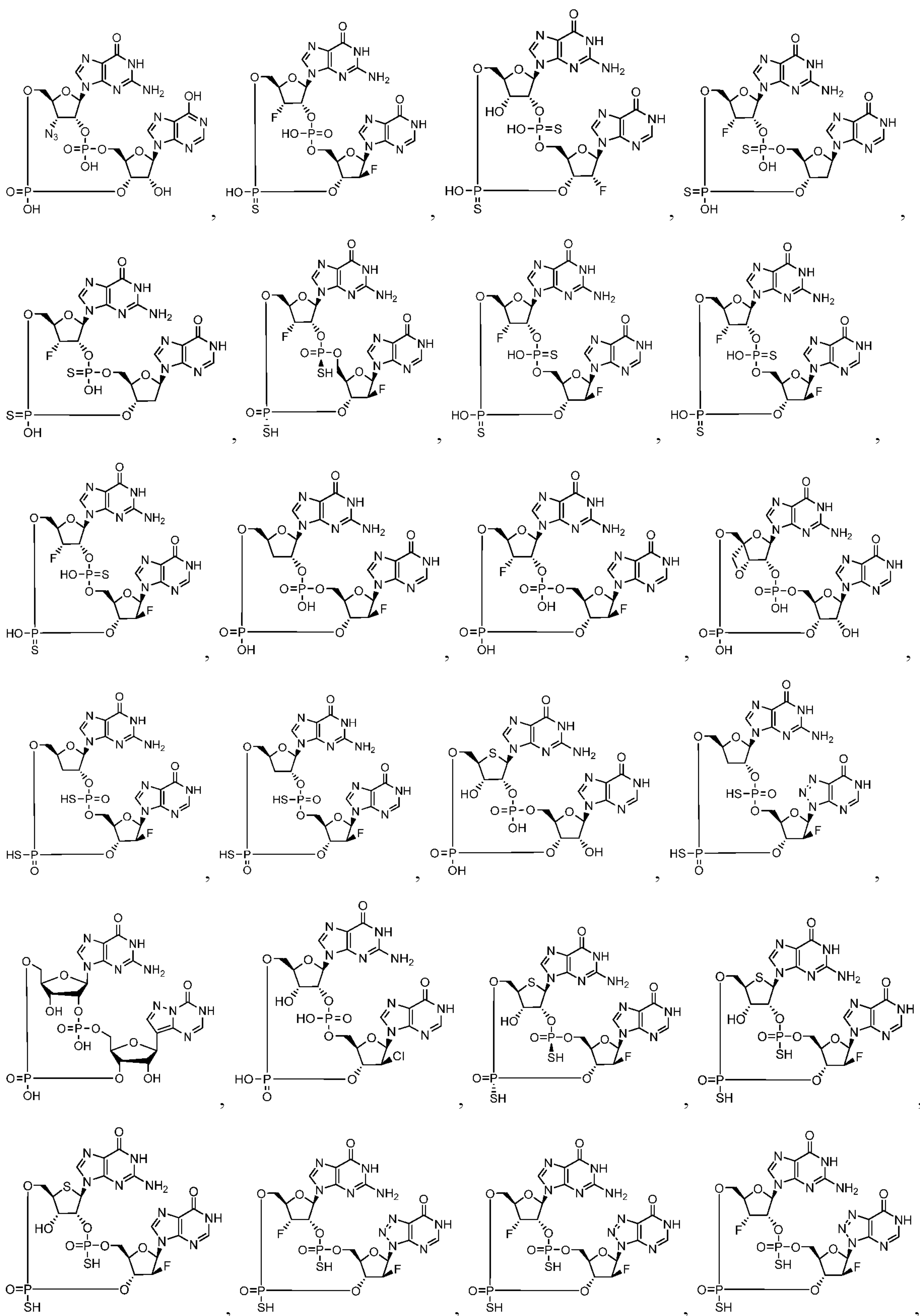


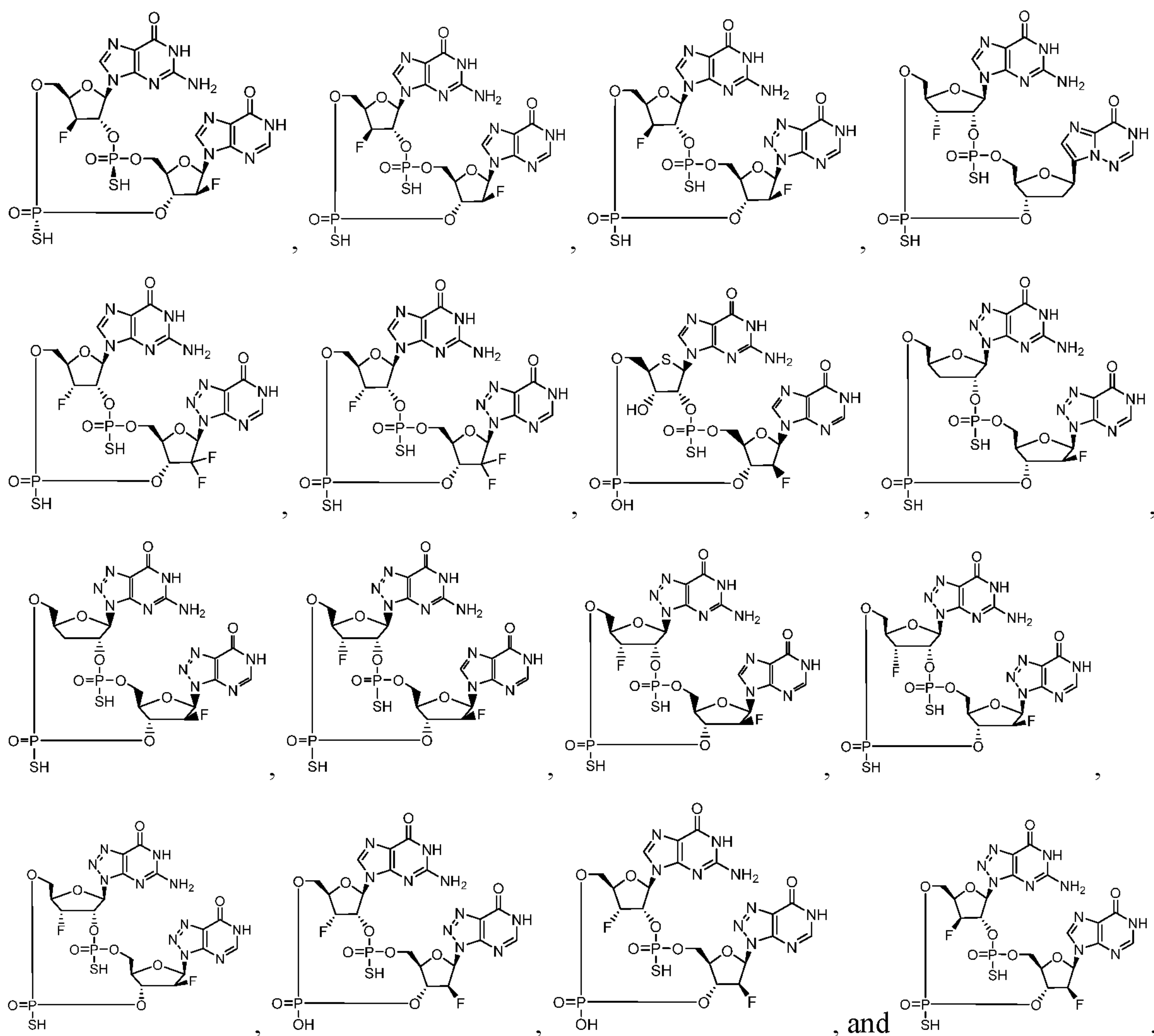




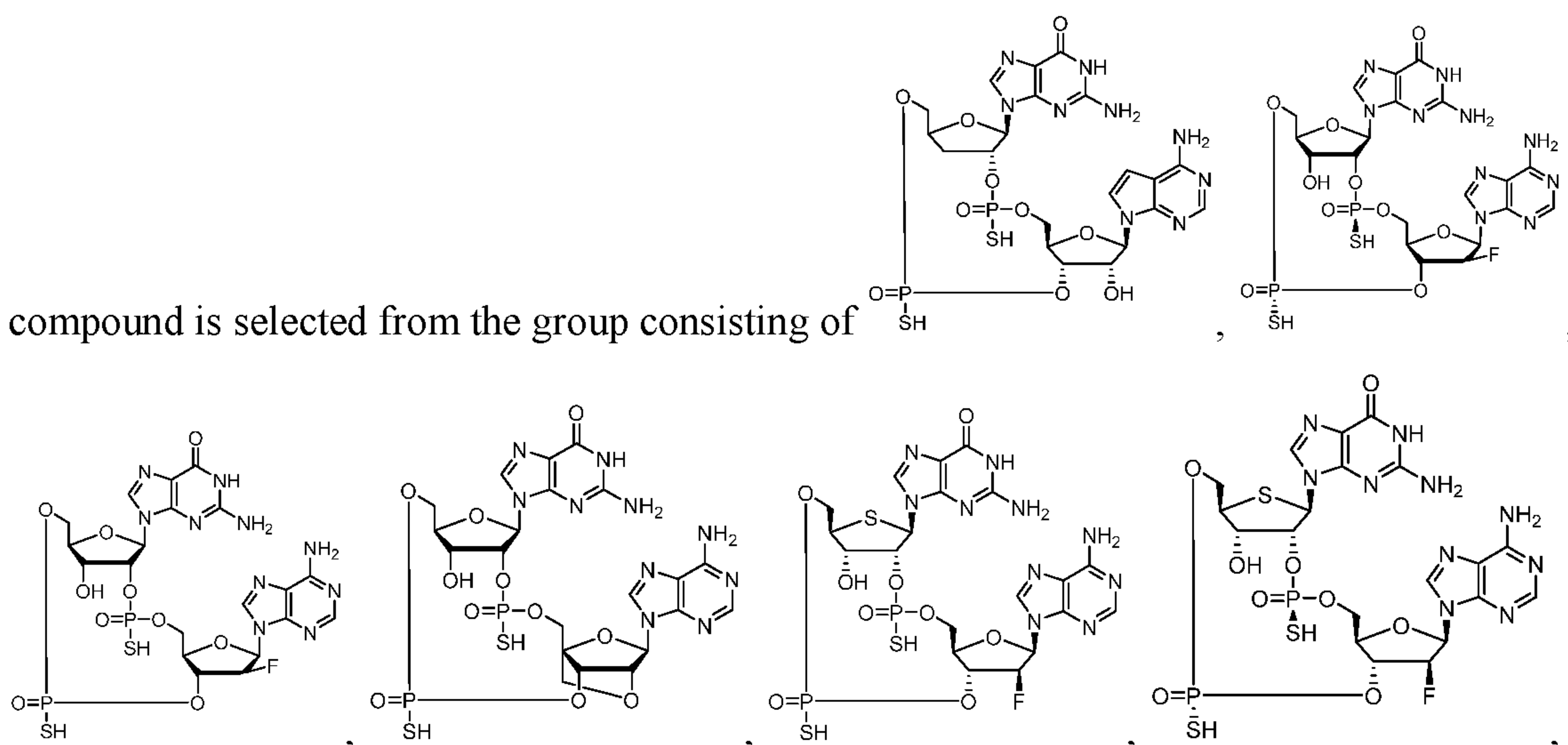




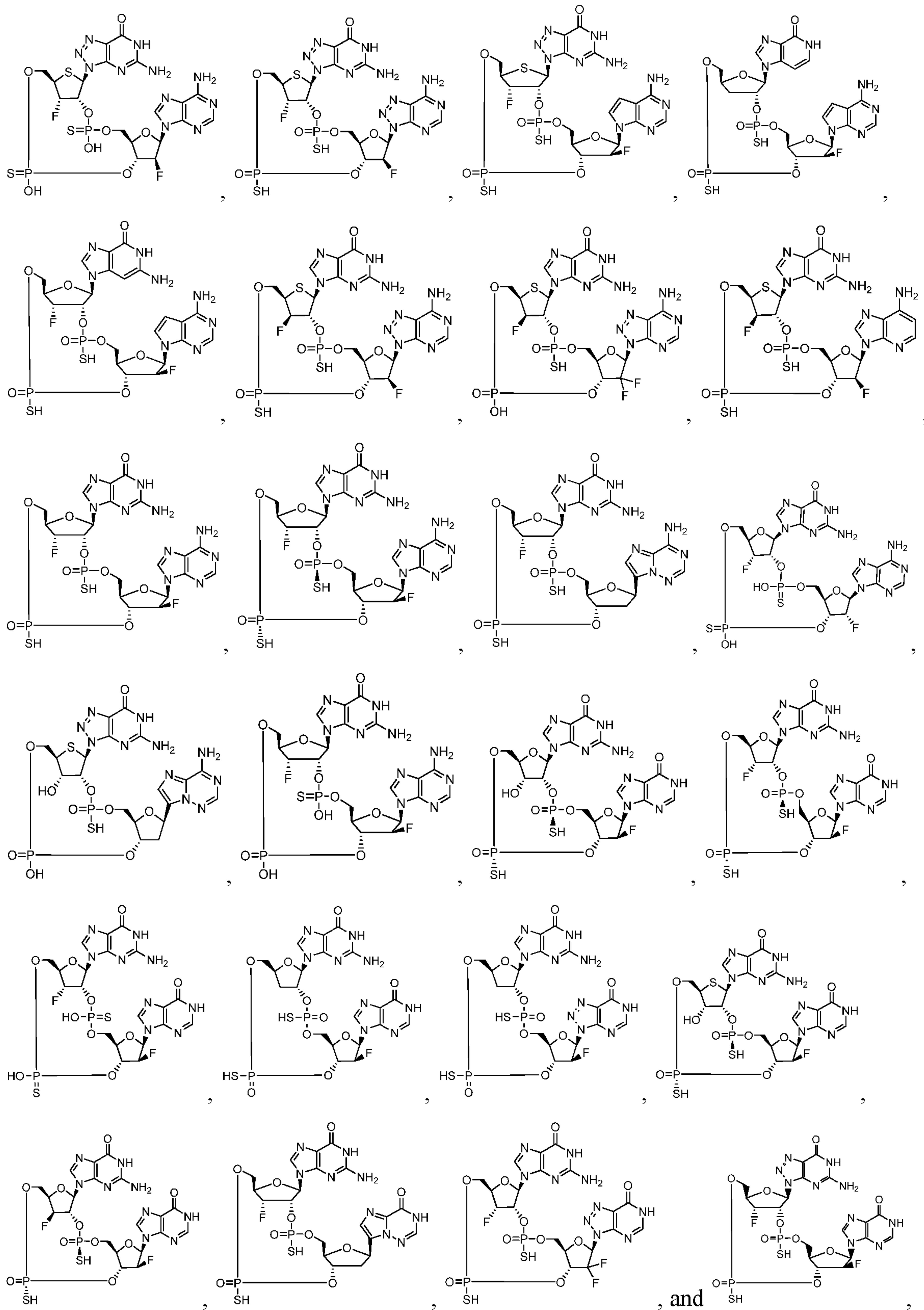




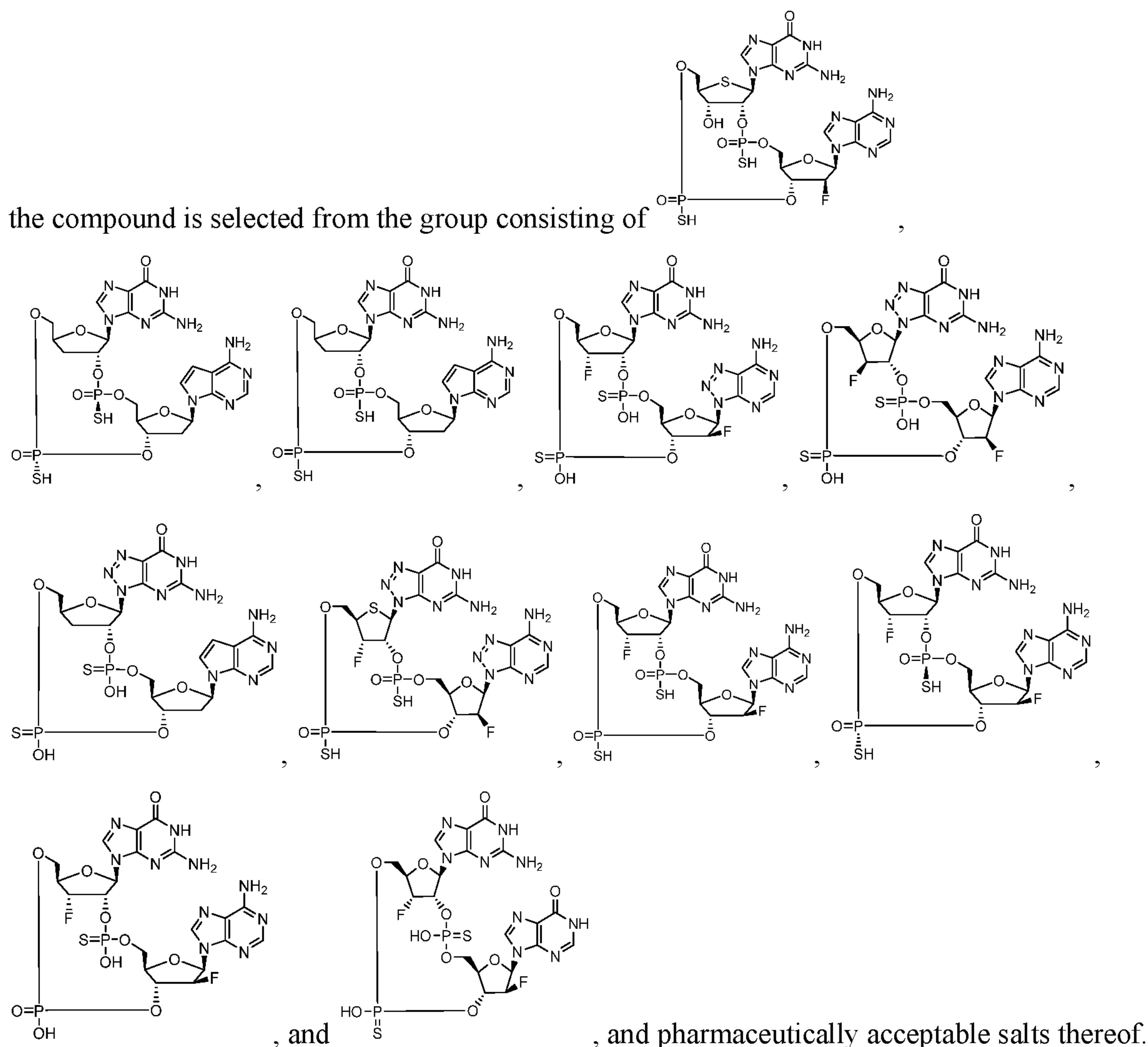
compound is selected from the group consisting of







and pharmaceutically acceptable salts thereof. In more particular aspects of this embodiment,



Methods of Preparing Compounds

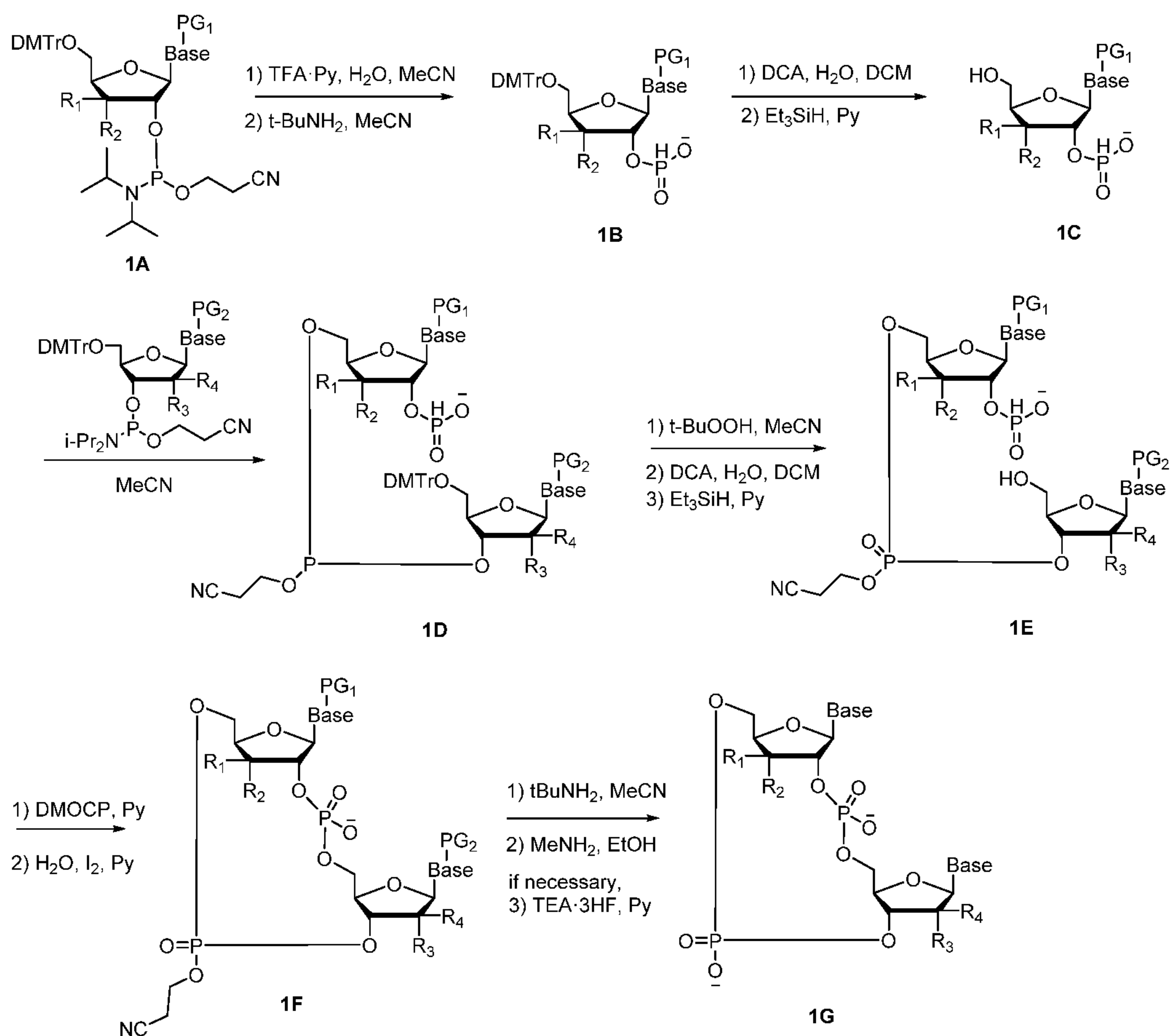
[0166] The CDN STING agonists of the disclosure may be prepared according to the methods disclosed in PCT International Patent Application No. PCT/US2016/046444, which published as PCT International Patent Application Publication No. WO2017/027646, and United States Patent Application No. 15/234,182, which published as U.S. Patent Application Publication No. US2017/0044206. In particular, several methods for preparing the compounds of general formula (I), compounds of general formula (I'), and compounds of general formula (I''), or pharmaceutically acceptable salts, hydrates, solvates, or prodrugs thereof, are described in the following Schemes. Starting materials and intermediates are purchased from commercial

sources, made from known procedures, or are otherwise illustrated. In some cases the order of carrying out the steps of the reaction schemes may be varied to facilitate the reaction or to avoid unwanted reaction products.

Method 1

- 5 [0167] One method for the preparation of examples of the disclosure is detailed in Scheme 1. This procedure was adequately modified from the previously reported procedure for cyclic dinucleotide synthesis (Barbara L. Gaffney *et al.*, *One-Flask Syntheses of c-di-GMP and the [Rp,Rp] and [Rp,Sp] Thiophosphate Analogues*, 12 ORG. LETT. 3269-3271 (2010)). The sequence starts with modified ribo-nucleoside with a nucleobase of which amino group was
- 10 appropriately protected with an alkyl or phenyl carbonyl group, a phosphoramidite functionality at 2'-O position, and DMTr ether at 5'-O position. It was treated with aqueous TFA/pyridine condition and subsequently t-butylamine to convert the 2'-phosphoramidite moiety to an H-phosphonate. Then, DMTr ether was removed under acidic condition. The resulting 5'-hydroxyl group was reacted with 3'-phosphoramidites of fully protected second modified
- 15 ribo-nucleoside to give a cyclized compound. It was immediately oxidized with t-butyl hydroperoxide. Then, the 5'-hydroxyl group of the second ribo-nucleoside was deprotected with dichloroacetic acid. Using 2-chloro-5,5-dimethyl-1,3,2-dioxaphosphinane 2-oxide as a coupling reagent, the H-phosphonate at 2'-O of the first ribo-nucleoside was reacted with 5'-OH of the second ribo-nucleoside to give a cyclic product. It was immediately oxidized with aqueous
- 20 iodine. Treatment with t-butylamine and methylamine plus fluoride anion in case silyl protection was used provided the desired cyclic dinucleotide 1G.

SCHEME 1



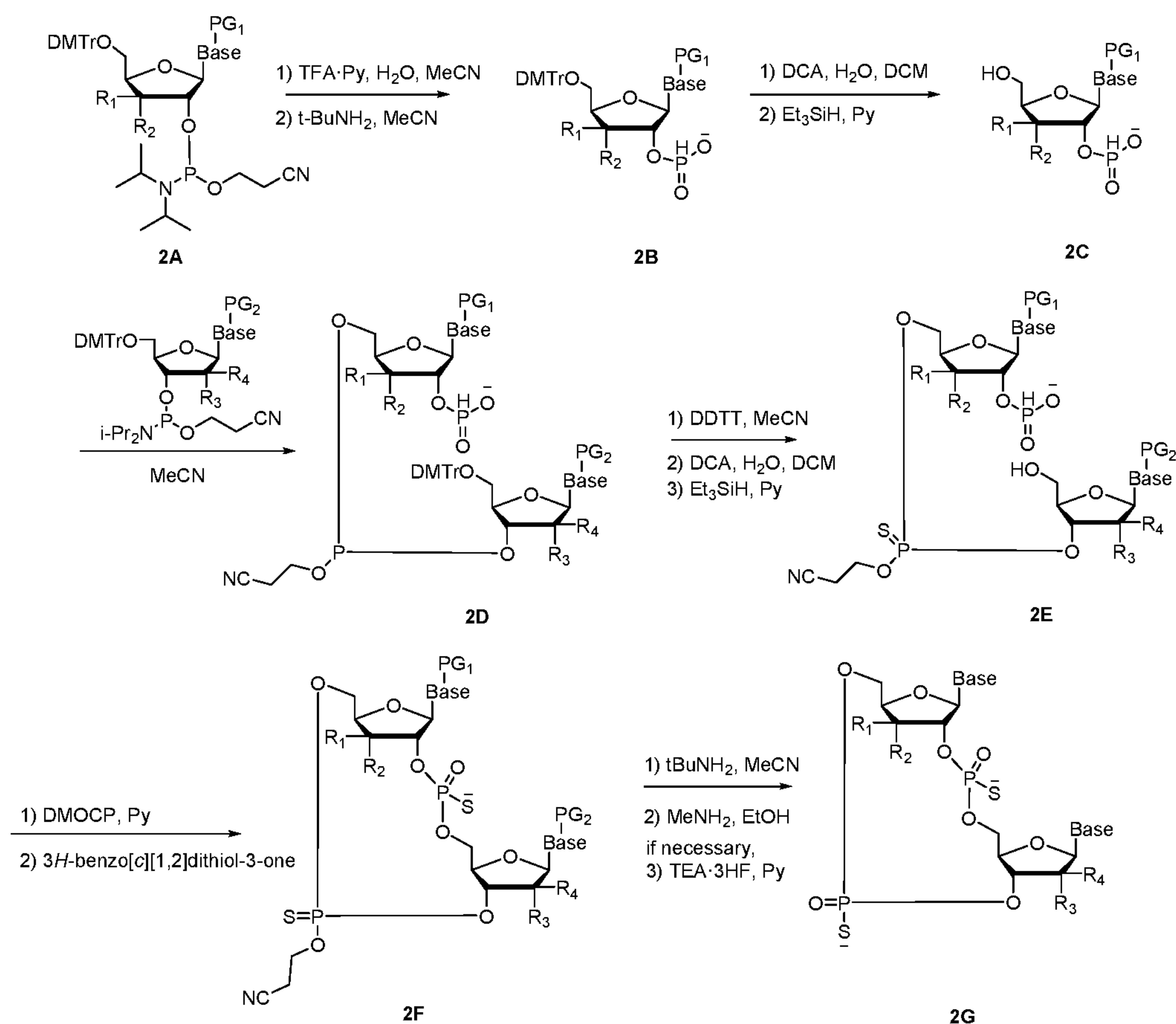
Method 2

[0168] Another method for the preparation of examples of the disclosure is detailed in Scheme 2. This procedure was modified from Scheme 1. The sequence starts with modified ribo-nucleoside with a nucleobase of which amino group was appropriately protected with an alkyl or phenyl carbonyl group, a phosphoramidite functionality at 2'-O position, and DMTr ether at 5'-O position. It was treated with aqueous TFA/pyridine condition and subsequently t-butylamine to convert the 2'-phosphoramidite moiety to an H-phosphonate. Then, DMTr ether was removed under acidic condition. The resulting 5'-hydroxyl group was reacted with 3'-phosphoramidites of fully protected second modified ribo-nucleoside to give a cyclized compound. It was immediately thioated with (E)-N,N-dimethyl-N'-(3-thioxo-3H-1,2,4-dithiazol-5-yl)formimidamide. Then, the 5'-hydroxyl group of the second ribo-nucleoside was deprotected with dichloroacetic acid. Using 2-chloro-5,5-dimethyl-1,3,2-dioxaphosphinane 2-oxide as a coupling reagent, the H-phosphonate at 2'-O of the first ribo-nucleoside was reacted

with 5'-OH of the second ribo-nucleoside to give a cyclic product. It was immediately thioated with 3H-benzo[c][1,2]dithiol-3-one. Treatment with t-butylamine and methylamine plus fluoride anion in case silyl protection was used provided the desired cyclic dinucleotide diphosphorothioate 2G.

5

SCHEME 2



[0169] The CDN STING agonists and a pharmaceutically acceptable carrier or excipient(s) will typically be formulated into a dosage form adapted for administration to a subject by a desired route of administration. For example, dosage forms include those adapted for (1) oral administration, such as tablets, capsules, caplets, pills, troches, powders, syrups, elixirs, suspensions, solutions, emulsions, sachets, and cachets; and (2) parenteral administration, such as sterile solutions, suspensions, and powders for reconstitution. Suitable pharmaceutically acceptable carriers or excipients will vary depending upon the particular dosage form chosen. In addition, suitable pharmaceutically acceptable carriers or excipients may be chosen for a particular function that they may serve in the composition. In embodiments, the CDN STING

agonist may be formulated into a dosage form that allows for systemic use, i.e., distribution of the CDN STING agonist throughout the body of the subject; examples of such systemic administration include oral administration and intravenous administration. In additional embodiments, the CDN STING agonist may be formulated into a dosage form that allows for
 5 targeted or isolated use, i.e., administration of the CDN STING agonist only to the portion of the subject's body to be treated; examples of such targetted administration include intratumoral injection.

[0170] The cyclic dinucleotide STING agonist is administered once every 1 to 30 days. In embodiments, the cyclic dinucleotide STING agonist is administered once every 3 to 28 days.
 10 In particular embodiments, the cyclic dinucleotide STING agonist is administered once every 3, 7, 14, 21, or 28 days.

[0171] In embodiments of such methods, the cyclic dinucleotide STING agonist is administered for from 2 to 36 months. In specific embodiments, the cyclic dinucleotide STING agonist is administered for up to 3 months.

15 [0172] In additional embodiments of such methods, the cyclic dinucleotide STING agonist is administered once every 3, 7, 14, 21, or 28 days for from 2 to 36 months. In further embodiments, the cyclic dinucleotide STING agonist is administered once every 3, 7, 14, 21, or 28 days for up to 3 months. In specific embodiments, the cyclic dinucleotide STING agonist is administered once every 3, 7, 14, 21, or 28 days for up to 3 months, followed by a period, lasting
 20 at least 2 months, in which the time interval between doses is increased by at least two-fold. In more specific embodiments, the cyclic dinucleotide STING agonist is administered once every 3, 7, 14, 21, or 28 days for up to 3 months, followed by a period, lasting at least 2 months, in which the time interval between doses is increased by at least three-fold. For example, if the cyclic dinucleotide STING agonist is administered once every 7 days for up to 3 months, it may be
 25 followed by a period in which the cyclic dinucleotide STING agonist is administered once every 14 or 21 days for up to two years.

[0173] In some embodiments, at least one of the therapeutic agents (the PD-1 antagonist and the cyclic dinucleotide STING agonist) in the combination therapy is administered using the same dosage regimen (dose, frequency, and duration of treatment) that is typically employed
 30 when the agent is used as monotherapy for treating the same condition. In other embodiments, the patient receives a lower total amount of at least one of the therapeutic agents in the

combination therapy than when the agent is used as monotherapy, *e.g.*, smaller doses, less frequent doses, and/or shorter treatment duration.

[0174] A combination therapy of the invention may be used prior to or following surgery to remove a tumor and may be used prior to, during, or after radiation treatment.

5 [0175] In some embodiments, a combination therapy of the invention is administered to a patient who has not previously been treated with a biotherapeutic or chemotherapeutic agent, targeted therapy, or hormonal therapy, *i.e.*, is treatment-naïve. In other embodiments, the combination therapy is administered to a patient who failed to achieve a sustained response after prior therapy with the biotherapeutic or chemotherapeutic agent, *i.e.*, is treatment-experienced.

10 [0176] The present disclosure further relates to methods of treating a cell-proliferation disorder, said method comprising administering to a subject in need thereof a combination therapy that comprises (a) a PD-1 antagonist; and (b) a cyclic dinucleotide STING agonist; wherein the PD-1 antagonist is administered once every 21 days; and the cyclic dinucleotide STING agonist is administered once every 1 to 30 days for 3 to 90 days, then optionally once
15 every 1 to 30 days for up to 1050 days. In embodiments, the CDN STING agonist is administered at least three times.

[0177] In specific embodiments, the cyclic dinucleotide STING agonist is administered once every 3 to 30 days for 9 to 90 days, then optionally once every 3 to 30 days for up to 1050 days. In specific embodiments, the cyclic dinucleotide STING agonist is administered once
20 every 3 to 21 days for 9 to 63 days, then optionally once every 3 to 21 days for up to 735 days. In further specific embodiments, the cyclic dinucleotide STING agonist is administered once every 7 to 21 days for 21 to 63 days, then optionally once every 7 to 21 days for up to 735 days. In still further embodiments, the cyclic dinucleotide STING agonist is administered once every 7
25 to 10 days for 21 to 30 days, then optionally once every 21 days for up to 735 days. In still further embodiments, the cyclic dinucleotide STING agonist is administered once every 7 days for 21 days, then optionally once every 21 days for up to 735 days. In additional embodiments, the cyclic dinucleotide STING agonist is administered once every 21 days for 63 days, then optionally once every 21 days for up to 735 days. In specific embodiments of the foregoing, the CDN STING agonist is administered at least three times.

30 [0178] In some embodiments, one or more optional “rest” periods, during which the CDN STING agonist is not administered, may be included in the treatment period. In specific embodiments, the optional rest period may be for from 3 to 30 days, from 7 to 21 days, or from 7

to 14 days. Following the rest period, dosing of the CDN STING agonist may be resumed as described above.

CELL-PROLIFERATION DISORDERS

[0179] The combination therapies disclosed herein are potentially useful in treating diseases or disorders including, but not limited to, cell-proliferation disorders. Cell-proliferation disorders include, but are not limited to, cancers, benign papillomatosis, gestational trophoblastic diseases, and benign neoplastic diseases, such as skin papilloma (warts) and genital papilloma. The terms “cancer”, “cancerous”, or “malignant” refer to or describe the physiological condition in mammals that is typically characterized by unregulated cell growth. A variety of cancers where PD-L1 or PD-L2 are implicated, whether malignant or benign and whether primary or secondary, may be treated or prevented with a method provided by the disclosure. Particularly preferred cancers that may be treated in accordance with the present disclosure include those characterized by elevated expression of one or both of PD-L1 and PD-L2 in tested tissue samples.

[0180] In specific embodiments, the disease or disorder to be treated is a cell-proliferation disorder. In certain embodiments, the cell-proliferation disorder is cancer. In particular embodiments, the cancer is selected from brain and spinal cancers, cancers of the head and neck, leukemia and cancers of the blood, skin cancers, cancers of the reproductive system, cancers of the gastrointestinal system, liver and bile duct cancers, kidney and bladder cancers, bone cancers, lung cancers, malignant mesothelioma, sarcomas, lymphomas, glandular cancers, thyroid cancers, heart tumors, germ cell tumors, malignant neuroendocrine (carcinoid) tumors, midline tract cancers, and cancers of unknown primary (*i.e.*, cancers in which a metastasized cancer is found but the original cancer site is not known). In particular embodiments, the cancer is present in an adult patient; in additional embodiments, the cancer is present in a pediatric patient. In particular embodiments, the cancer is AIDS-related.

[0181] In specific embodiments, the cancer is selected from brain and spinal cancers. In particular embodiments, the brain and spinal cancer is selected from the group consisting of anaplastic astrocytomas, glioblastomas, astrocytomas, and esthesioneuroblastomas (also known as olfactory blastomas). In particular embodiments, the brain cancer is selected from the group consisting of astrocytic tumor (e.g., pilocytic astrocytoma, subependymal giant-cell astrocytoma, diffuse astrocytoma, pleomorphic xanthoastrocytoma, anaplastic astrocytoma, astrocytoma, giant cell glioblastoma, glioblastoma, secondary glioblastoma, primary adult glioblastoma, and

primary pediatric glioblastoma), oligodendroglial tumor (e.g., oligodendroglioma, and anaplastic oligodendroglioma), oligoastrocytic tumor (e.g., oligoastrocytoma, and anaplastic oligoastrocytoma), ependymoma (e.g., myxopapillary ependymoma, and anaplastic ependymoma); medulloblastoma, primitive neuroectodermal tumor, schwannoma, meningioma, atypical meningioma, anaplastic meningioma, pituitary adenoma, brain stem glioma, cerebellar astrocytoma, cerebral astrocytoma/malignant glioma, visual pathway and hypothalamic glioma, and primary central nervous system lymphoma. In specific instances of these embodiments, the brain cancer is selected from the group consisting of glioma, glioblastoma multiforme, paraganglioma, and supratentorial primordial neuroectodermal tumors (sPNET).

[0182] In specific embodiments, the cancer is selected from cancers of the head and neck, including recurrent or metastatic head and neck squamous cell carcinoma (HNSCC), nasopharyngeal cancers, nasal cavity and paranasal sinus cancers, hypopharyngeal cancers, oral cavity cancers (e.g., squamous cell carcinomas, lymphomas, and sarcomas), lip cancers, oropharyngeal cancers, salivary gland tumors, cancers of the larynx (e.g., laryngeal squamous cell carcinomas, rhabdomyosarcomas), and cancers of the eye or ocular cancers. In particular embodiments, the ocular cancer is selected from the group consisting of intraocular melanoma and retinoblastoma.

[0183] In specific embodiments, the cancer is selected from leukemia and cancers of the blood. In particular embodiments, the cancer is selected from the group consisting of myeloproliferative neoplasms, myelodysplastic syndromes, myelodysplastic/myeloproliferative neoplasms, acute myeloid leukemia (AML), myelodysplastic syndrome (MDS), chronic myelogenous leukemia (CML), myeloproliferative neoplasm (MPN), post-MPN AML, post-MDS AML, del(5q)-associated high risk MDS or AML, blast-phase chronic myelogenous leukemia, angioimmunoblastic lymphoma, acute lymphoblastic leukemia, Langerans cell histiocytosis, hairy cell leukemia, and plasma cell neoplasms including plasmacytomas and multiple myelomas. Leukemias referenced herein may be acute or chronic.

[0184] In specific embodiments, the cancer is selected from skin cancers. In particular embodiments, the skin cancer is selected from the group consisting of melanoma, squamous cell cancers, and basal cell cancers. In specific embodiments, the skin cancer is unresectable or metastatic melanoma.

[0185] In specific embodiments, the cancer is selected from cancers of the reproductive system. In particular embodiments, the cancer is selected from the group consisting of breast

cancers, cervical cancers, vaginal cancers, ovarian cancers, endometrial cancers, prostate cancers, penile cancers, and testicular cancers. In specific instances of these embodiments, the cancer is a breast cancer selected from the group consisting of ductal carcinomas and phyllodes tumors. In specific instances of these embodiments, the breast cancer may be male breast cancer
 5 or female breast cancer. In more specific instances of these embodiments, the breast cancer is triple-negative breast cancer. In specific instances of these embodiments, the cancer is a cervical cancer selected from the group consisting of squamous cell carcinomas and adenocarcinomas. In specific instances of these embodiments, the cancer is an ovarian cancer selected from the group consisting of epithelial cancers.

10 [0186] In specific embodiments, the cancer is selected from cancers of the gastrointestinal system. In particular embodiments, the cancer is selected from the group consisting of esophageal cancers, gastric cancers (also known as stomach cancers), gastrointestinal carcinoid tumors, pancreatic cancers, gallbladder cancers, colorectal cancers, and anal cancer. In instances of these embodiments, the cancer is selected from the group consisting
 15 of esophageal squamous cell carcinomas, esophageal adenocarcinomas, gastric adenocarcinomas, gastrointestinal carcinoid tumors, gastrointestinal stromal tumors, gastric lymphomas, gastrointestinal lymphomas, solid pseudopapillary tumors of the pancreas, pancreatoblastoma, islet cell tumors, pancreatic carcinomas including acinar cell carcinomas and ductal adenocarcinomas, gallbladder adenocarcinomas, colorectal adenocarcinomas, and anal
 20 squamous cell carcinomas.

[0187] In specific embodiments, the cancer is selected from liver and bile duct cancers. In particular embodiments, the cancer is liver cancer (also known as hepatocellular carcinoma). In particular embodiments, the cancer is bile duct cancer (also known as cholangiocarcinoma); in instances of these embodiments, the bile duct cancer is selected from the group consisting of
 25 intrahepatic cholangiocarcinoma and extrahepatic cholangiocarcinoma.

[0188] In specific embodiments, the cancer is selected from kidney and bladder cancers. In particular embodiments, the cancer is a kidney cancer selected from the group consisting of renal cell cancer, Wilms tumors, and transitional cell cancers. In particular embodiments, the cancer is a bladder cancer selected from the group consisting of urothelial carcinoma (a
 30 transitional cell carcinoma), squamous cell carcinomas, and adenocarcinomas.

[0189] In specific embodiments, the cancer is selected from bone cancers. In particular embodiments, the bone cancer is selected from the group consisting of osteosarcoma, malignant fibrous histiocytoma of bone, Ewing sarcoma, chordoma (cancer of the bone along the spine).

[0190] In specific embodiments, the cancer is selected from lung cancers. In particular
5 embodiments, the lung cancer is selected from the group consisting of non-small cell lung cancer, small cell lung cancers, bronchial tumors, and pleuropulmonary blastomas.

[0191] In specific embodiments, the cancer is selected from malignant mesothelioma. In particular embodiments, the cancer is selected from the group consisting of epithelial mesothelioma and sarcomatoids.

10 [0192] In specific embodiments, the cancer is selected from sarcomas. In particular embodiments, the sarcoma is selected from the group consisting of central chondrosarcoma, central and periosteal chondroma, fibrosarcoma, clear cell sarcoma of tendon sheaths, and Kaposi's sarcoma.

[0193] In specific embodiments, the cancer is selected from lymphomas. In particular
15 embodiments, the cancer is selected from the group consisting of Hodgkin lymphoma (e.g., Reed-Sternberg cells), non-Hodgkin lymphoma (e.g., diffuse large B-cell lymphoma, follicular lymphoma, mycosis fungoides, Sezary syndrome, primary central nervous system lymphoma), cutaneous T-cell lymphomas, and primary central nervous system lymphomas.

[0194] In specific embodiments, the cancer is selected from glandular cancers. In
20 particular embodiments, the cancer is selected from the group consisting of adrenocortical cancer (also known as adrenocortical carcinoma or adrenal cortical carcinoma), pheochromocytomas, paragangliomas, pituitary tumors, thymoma, and thymic carcinomas.

[0195] In specific embodiments, the cancer is selected from thyroid cancers. In particular
25 embodiments, the thyroid cancer is selected from the group consisting of medullary thyroid carcinomas, papillary thyroid carcinomas, and follicular thyroid carcinomas.

[0196] In specific embodiments, the cancer is selected from germ cell tumors. In particular
embodiments, the cancer is selected from the group consisting of malignant extracranial germ cell tumors and malignant extragonadal germ cell tumors. In specific instances of these embodiments, the malignant extragonadal germ cell tumors are selected from the group
30 consisting of nonseminomas and seminomas.

[0197] In specific embodiments, the cancer is selected from heart tumors. In particular
embodiments, the heart tumor is selected from the group consisting of malignant teratoma,

lymphoma, rhabdomyosarcoma, angiosarcoma, chondrosarcoma, infantile fibrosarcoma, and synovial sarcoma.

[0198] In specific embodiments, the cell-proliferation disorder is selected from benign papillomatosis, benign neoplastic diseases and gestational trophoblastic diseases. In particular
5 embodiments, the benign neoplastic disease is selected from skin papilloma (warts) and genital papilloma. In particular embodiments, the gestational trophoblastic disease is selected from the group consisting of hydatidiform moles, and gestational trophoblastic neoplasia (*e.g.*, invasive moles, choriocarcinomas, placental-site trophoblastic tumors, and epithelioid trophoblastic tumors).

10 [0199] In embodiments, the cell-proliferation disorder is a cancer that has metastasized, for example, a liver metastases from colorectal cancer.

[0200] In embodiments, the cell-proliferation disorder is selected from the group consisting of solid tumors and lymphomas. In particular embodiments, the cell-proliferation disorder is selected from the group consisting of advanced or metastatic solid tumors and
15 lymphomas. In more particular embodiments, the cell-proliferation disorder is selected from the group consisting of malignant melanoma, head and neck squamous cell carcinoma, breast adenocarcinoma, and lymphomas. In aspects of such embodiments, the lymphomas are selected from the group consisting of diffuse large B-cell lymphoma, follicular lymphoma, mantle cell lymphoma, small lymphocytic lymphoma, mediastinal large B-cell lymphoma, splenic marginal
20 zone B-cell lymphoma, extranodal marginal zone B-cell lymphoma of mucosa-associated lymphoid tissue (malt), nodal marginal zone B-cell lymphoma, lymphoplasmacytic lymphoma, primary effusion lymphoma, Burkitt lymphoma, anaplastic large cell lymphoma (primary cutaneous type), anaplastic large cell lymphoma (systemic type), peripheral T-cell lymphoma, angioimmunoblastic T-cell lymphoma, adult T-cell lymphoma/leukemia, nasal type extranodal
25 NK/T-cell lymphoma, enteropathy-associated T-cell lymphoma, gamma/delta hepatosplenic T-cell lymphoma, subcutaneous panniculitis-like T-cell lymphoma, mycosis fungoides, and Hodgkin lymphoma.

[0201] In particular embodiments, the cell-proliferation disorder is classified as stage III cancer or stage IV cancer. In instances of these embodiments, the cancer is not surgically
30 resectable.

METHODS, USES, AND MEDICAMENTS

[0202] Products provided as therapeutic combinations may include a composition comprising a PD-1 antagonist and a CDN STING agonist together in the same pharmaceutical composition, or may include a composition comprising a PD-1 antagonist, and a composition comprising a CDN STING agonist in separate form, e.g., in the form of a kit or in any form
 5 designed to enable separate administration either concurrently or on separate dosing schedules.

[0203] The combination therapy may also comprise one or more additional therapeutic agents. The additional therapeutic agent may be, e.g., a chemotherapeutic, a biotherapeutic agent (including but not limited to antibodies to VEGF, VEGFR, EGFR, Her2/neu, other growth factor receptors, CD20, CD40, CD-40L, CTLA-4, OX-40, 4-1BB, and ICOS), an immunogenic agent
 10 (for example, attenuated cancerous cells, tumor antigens, antigen presenting cells such as dendritic cells pulsed with tumor derived antigen or nucleic acids, immune stimulating cytokines (for example, IL-2, IFN α 2, GM-CSF), and cells transfected with genes encoding immune stimulating cytokines such as but not limited to GM-CSF). The one or more additional active agents may be co-administered either with the PD-1 antagonist or with the CDN STING agonist.
 15 The additional active agent(s) may be administered in a single dosage form with one or more co-administered agent selected from the PD-1 antagonist and the CDN STING agonist, or the additional active agent(s) may be administered in separate dosage form(s) from the dosage forms containing the PD-1 antagonist and/or the CDN STING agonist.

[0204] The therapeutic combination disclosed herein may be used in combination with
 20 one or more other active agents, including but not limited to, other anti-cancer agents that are used in the prevention, treatment, control, amelioration, or reduction of risk of a particular disease or condition (e.g., cell-proliferation disorders). In one embodiment, a compound disclosed herein is combined with one or more other anti-cancer agents for use in the prevention, treatment, control amelioration, or reduction of risk of a particular disease or condition for which
 25 the compounds disclosed herein are useful. Such other active agents may be administered, by a route and in an amount commonly used therefor, contemporaneously or sequentially with a compound of the present disclosure.

[0205] The additional active agent(s) may be one or more agents selected from the group consisting of STING agonists, anti-viral compounds, antigens, adjuvants, anti-cancer agents,
 30 CTLA-4, LAG-3, and PD-1 pathway antagonists, lipids, liposomes, peptides, cytotoxic agents, chemotherapeutic agents, immunomodulatory cell lines, checkpoint inhibitors, vascular endothelial growth factor (VEGF) receptor inhibitors, topoisomerase II inhibitors, smoothen

inhibitors, alkylating agents, anti-tumor antibiotics, anti-metabolites, retinoids, and immunomodulatory agents including but not limited to anti-cancer vaccines. It will be understood the descriptions of the above additional active agents may be overlapping. It will also be understood that the treatment combinations are subject to optimization, and it is understood that the best combination to use of the PD-1 antagonist and/or the CDN STING agonist, and one or more additional active agents will be determined based on the individual patient needs.

[0206] When the therapeutic combination disclosed herein is used contemporaneously with one or more other active agents, the PD-1 antagonist and/or the CDN STING agonist may be administered either simultaneously with, or before or after, one or more other active agent(s). Either of the PD-1 antagonist and/or the CDN STING agonist may be administered separately, by the same or different route of administration, or together in the same pharmaceutical composition as the other agent(s).

[0207] The weight ratio of the PD-1 antagonist to the CDN STING agonist may be varied and will depend upon the therapeutically effective dose of each agent. Generally, a therapeutically effective dose of each will be used. Combinations including at least one PD-1 antagonist, at least one CDN STING agonist, and other active agents will generally include a therapeutically effective dose of each active agent. In such combinations, the PD-1 antagonist and/or the CDN STING agonist disclosed herein and other active agents may be administered separately or in conjunction. In addition, the administration of one element may be prior to, concurrent with, or subsequent to the administration of other agent(s).

[0208] In one embodiment, this disclosure provides a PD-1 antagonist and/or a CDN STING agonist, and at least one other active agent as a combined preparation for simultaneous, separate or sequential use in therapy. In one embodiment, the therapy is the treatment of a cell-proliferation disorder, such as cancer.

[0209] In one embodiment, the disclosure provides a kit comprising two or more separate pharmaceutical compositions, one of which contains a PD-1 antagonist and another of which contains a CDN STING agonist. In one embodiment, the kit comprises means for separately retaining said compositions, such as a container, divided bottle, or divided foil packet. A kit of this disclosure may be used for administration of different dosage forms, for example, oral and parenteral, for administration of the separate compositions at different dosage intervals, or for

titration of the separate compositions against one another. To assist with compliance, a kit of the disclosure typically comprises directions for administration.

[0210] The disclosure also provides the use of a CDN STING agonist for treating a cell-proliferation disorder, where the patient has previously (e.g., within 24 hours) been treated with a PD-1 antagonist. The disclosure also provides the use of a PD-1 antagonist for treating a cell-proliferation disorder, where the patient has previously (e.g., within 24 hours) been treated with a CDN STING agonist.

[0211] Anti-viral compounds that may be used in combination with the therapeutic combinations disclosed herein include hepatitis B virus (HBV) inhibitors, hepatitis C virus (HCV) protease inhibitors, HCV polymerase inhibitors, HCV NS4A inhibitors, HCV NS5A inhibitors, HCV NS5b inhibitors, and human immunodeficiency virus (HIV) inhibitors.

[0212] Antigens and adjuvants that may be used in combination with the therapeutic combinations disclosed herein include B7 costimulatory molecule, interleukin-2, interferon- γ , GM-CSF, CTLA-4 antagonists, OX-40/OX-40 ligand, CD40/CD40 ligand, sargramostim, levamisole, vaccinia virus, Bacille Calmette-Guerin (BCG), liposomes, alum, Freund's complete or incomplete adjuvant, detoxified endotoxins, mineral oils, surface active substances such as lipolecithin, pluronic polyols, polyanions, peptides, and oil or hydrocarbon emulsions. Adjuvants, such as aluminum hydroxide or aluminum phosphate, can be added to increase the ability of the vaccine to trigger, enhance, or prolong an immune response. Additional materials, such as cytokines, chemokines, and bacterial nucleic acid sequences, like CpG, a toll-like receptor (TLR) 9 agonist as well as additional agonists for TLR 2, TLR 4, TLR 5, TLR 7, TLR 8, TLR9, including lipoprotein, lipopolysaccharide (LPS), monophosphoryllipid A, lipoteichoic acid, imiquimod, resiquimod, and in addition retinoic acid-inducible gene I (RIG-I) agonists such as poly I:C, used separately or in combination are also potential adjuvants.

[0213] Examples of cytotoxic agents that may be used in combination with the therapeutic combinations disclosed herein include, but are not limited to, arsenic trioxide (sold under the tradename TRISENOX[®]), asparaginase (also known as L-asparaginase, and Erwinia L-asparaginase, sold under the tradenames ELSPAR[®] and KIDROLASE[®]).

[0214] Chemotherapeutic agents that may be used in combination with the therapeutic combinations disclosed herein include abiraterone acetate, altretamine, anhydrovinblastine, auristatin, bexarotene, bicalutamide, BMS 184476, 2,3,4,5,6-pentafluoro-N-(3-fluoro-4-methoxyphenyl)benzene sulfonamide, bleomycin, N,N-dimethyl-L-valyl-L-valyl-N-methyl-L-

valyl-L-prolyl- 1-L-proline-t-butylamide, cachectin, cemadotin, chlorambucil, cyclophosphamide, 3',4'-didehydro-4'deoxy-8'-norvin-caleukoblastine, docetaxol, doxetaxel, cyclophosphamide, carboplatin, carmustine, cisplatin, cryptophycin, cyclophosphamide, cytarabine, dacarbazine (DTIC), dactinomycin, daunorubicin, decitabine dolastatin, doxorubicin (adriamycin), etoposide, 5 5-fluorouracil, finasteride, flutamide, hydroxyurea and hydroxyurea and taxanes, ifosfamide, liarozole, lonidamine, lomustine (CCNU), MDV3100, mechlorethamine (nitrogen mustard), melphalan, mivobulin isethionate, rhizoxin, sertenef, streptozocin, mitomycin, methotrexate, taxanes, nilutamide, nivolumab, onapristone, paclitaxel, pembrolizumab, prednimustine, procarbazine, RPR109881, stramustine phosphate, tamoxifen, tasonermin, taxol, tretinoin, 10 vinblastine, vincristine, vindesine sulfate, and vinflunine, and pharmaceutically acceptable salts thereof.

[0215] Examples of vascular endothelial growth factor (VEGF) receptor inhibitors include, but are not limited to, bevacizumab (sold under the trademark AVASTIN by Genentech/Roche), axitinib (described in PCT International Patent Publication No.

15 WO01/002369), Brivanib Alaninate ((S)-((R)-1-(4-(4-Fluoro-2-methyl-1H-indol-5-yloxy)-5-methylpyrrolo[2,1-f][1,2,4]triazin-6-yloxy)propan-2-yl)2-aminopropanoate, also known as BMS-582664), motesanib (N-(2,3-dihydro-3,3-dimethyl-1H-indol-6-yl)-2-[(4-pyridinylmethyl)amino]-3-pyridinecarboxamide. and described in PCT International Patent Application Publication No. WO02/068470), pasireotide (also known as SO 230, and described in PCT International Patent 20 Publication No. WO02/010192), and sorafenib (sold under the tradename NEXAVAR).

[0216] Examples of topoisomerase II inhibitors, include but are not limited to, etoposide (also known as VP-16 and Etoposide phosphate, sold under the tradenames TOPOSAR, VEPESID, and ETOPOPHOS), and teniposide (also known as VM-26, sold under the tradename VUMON).

25 [0217] Examples of hypomethylating agents and alkylating agents, include but are not limited to, 5-azacytidine (sold under the trade name VIDAZA), decitabine (sold under the trade name of DECOGEN), temozolomide (sold under the trade names TEMODAR and TEMODAL), dactinomycin (also known as actinomycin-D and sold under the tradename COSMEGEN), melphalan (also known as L-PAM, L-sarcolysin, and phenylalanine mustard, sold under the 30 tradename ALKERAN), altretamine (also known as hexamethylmelamine (HMM), sold under the tradename HEXALEN), carmustine (sold under the tradename BCNU), bendamustine (sold under the tradename TREANDA), busulfan (sold under the tradenames BUSULFEX[®] and

MYLERAN[®]), carboplatin (sold under the tradename PARAPLATIN[®]), lomustine (also known as CCNU, sold under the tradename CEENU[®]), cisplatin (also known as CDDP, sold under the tradenames PLATINOL[®] and PLATINOL[®]-AQ), chlorambucil (sold under the tradename LEUKERAN[®]), cyclophosphamide (sold under the tradenames CYTOXAN[®] and NEOSAR[®]),
 5 dacarbazine (also known as DTIC, DIC and imidazole carboxamide, sold under the tradename DTIC-DOME[®]), altretamine (also known as hexamethylmelamine (HMM) sold under the tradename HEXALEN[®]), ifosfamide (sold under the tradename IFEX[®]), procarbazine (sold under the tradename MATULANE[®]), mechlorethamine (also known as nitrogen mustard, mustine and mechloroethamine hydrochloride, sold under the tradename MUSTARGEN[®]), streptozocin (sold
 10 under the tradename ZANOSAR[®]), thiotepa (also known as thiophosphoamide, TESPA and TSPA, and sold under the tradename THIOPLEX[®], and pharmaceutically acceptable salts thereof.

[0218] Examples of anti-tumor antibiotics include, but are not limited to, doxorubicin (sold under the tradenames ADRIAMYCIN[®] and RUBEX[®]), bleomycin (sold under the tradename LENOXANE[®]), daunorubicin (also known as dauorubicin hydrochloride, daunomycin, and
 15 rubidomycin hydrochloride, sold under the tradename CERUBIDINE[®]), daunorubicin liposomal (daunorubicin citrate liposome, sold under the tradename DAUNOXOME[®]), mitoxantrone (also known as DHAD, sold under the tradename NOVANTRONE[®]), epirubicin (sold under the tradename ELLENCE[™]), idarubicin (sold under the tradenames IDAMYCIN[®], IDAMYCIN PFS[®]), and mitomycin C (sold under the tradename MUTAMYCIN[®]).

[0219] Examples of anti-metabolites include, but are not limited to, claribine (2-chlorodeoxyadenosine, sold under the tradename LEUSTATIN[®]), 5-fluorouracil (sold under the tradename ADRUCIL[®]), 6-thioguanine (sold under the tradename PURINETHOL[®]), pemetrexed (sold under the tradename ALIMTA[®]), cytarabine (also known as arabinosylcytosine (Ara-C), sold under the tradename CYTOSAR-U[®]), cytarabine liposomal (also known as Liposomal Ara-C, sold
 25 under the tradename DEPOCYT[™]), decitabine (sold under the tradename DACOGEN[®]), hydroxyurea and (sold under the tradenames HYDREA[®], DROXIA[™] and MYLOCEL[™]), fludarabine (sold under the tradename FLUDARA[®]), floxuridine (sold under the tradename FUDR[®]), cladribine (also known as 2-chlorodeoxyadenosine (2-CdA) sold under the tradename LEUSTATIN[™]), methotrexate (also known as amethopterin, methotrexate sodium (MTX), sold
 30 under the tradenames RHEUMATREX[®] and TREXALL[™]), and pentostatin (sold under the tradename NIPENT[®]).

[0220] Examples of retinoids include, but are not limited to, alitretinoin (sold under the tradename PANRETIN[®]), tretinoin (all-trans retinoic acid, also known as ATRA, sold under the tradename VESANOID[®]), Isotretinoin (13-c/s-retinoic acid, sold under the tradenames ACCUTANE[®], AMNESTEEM[®], CLARAVIS[®], CLARUS[®], DECUTAN[®], ISOTANE[®], IZOTECH[®],
 5 ORATANE[®], ISOTRET[®], and SOTRET[®]), and bexarotene (sold under the tradename TARGRETIN[®]).

ADDITIONAL EMBODIMENTS

[0221] The present disclosure further relates to methods of treating a cell-proliferation disorder, said method comprising administering to a subject in need thereof a combination therapy that comprises (a) a PD-1 antagonist; and (b) a cyclic dinucleotide STING agonist;
 10 wherein the PD-1 antagonist is administered once every 21 days; and the cyclic dinucleotide STING agonist is administered once every 1 to 30 days. In embodiments, the cyclic dinucleotide STING agonist is administered once every 3 to 28 days. In particular embodiments, the cyclic dinucleotide STING agonist is administered once every 3, 7, 14, 21, or 28 days.

[0222] In embodiments of such methods, the cyclic dinucleotide STING agonist is
 15 administered for from 2 to 36 months. In specific embodiments, the cyclic dinucleotide STING agonist is administered for up to 3 months.

[0223] In additional embodiments of such methods, the cyclic dinucleotide STING agonist is administered once every 3, 7, 14, 21, or 28 days for from 2 to 36 months. In further embodiments, the cyclic dinucleotide STING agonist is administered once every 3, 7, 14, 21, or
 20 28 days for up to 3 months. In specific embodiments, the cyclic dinucleotide STING agonist is administered once every 3, 7, 14, 21, or 28 days for up to 3 months, followed by a period, lasting at least 2 months, in which the time interval between doses is increased by at least two-fold. In more specific embodiments, the cyclic dinucleotide STING agonist is administered once every 3,
 7, 14, 21, or 28 days for up to 3 months, followed by a period, lasting at least 2 months, in which
 25 the time interval between doses is increased by at least three-fold. For example, if the cyclic dinucleotide STING agonist is administered once every 7 days for up to 3 months, it may be followed by a period in which the cyclic dinucleotide STING agonist is administered once every 14 or 21 days for up to two years.

[0224] The present disclosure further relates to methods of treating a cell-proliferation
 30 disorder, said method comprising administering to a subject in need thereof a combination therapy that comprises (a) a PD-1 antagonist; and (b) a cyclic dinucleotide STING agonist; wherein the PD-1 antagonist is administered once every 21 days; and the cyclic dinucleotide

STING agonist is administered once every 1 to 30 days for 3 to 90 days, then optionally once every 1 to 30 days for up to 1050 days. In embodiments, the CDN STING agonist is administered at least three times.

[0225] In specific embodiments, the cyclic dinucleotide STING agonist is administered once every 3 to 30 days for 9 to 90 days, then optionally once every 3 to 30 days for up to 1050 days. In specific embodiments, the cyclic dinucleotide STING agonist is administered once every 3 to 21 days for 9 to 63 days, then optionally once every 3 to 21 days for up to 735 days. In further specific embodiments, the cyclic dinucleotide STING agonist is administered once every 7 to 21 days for 21 to 63 days, then optionally once every 7 to 21 days for up to 735 days. In still further embodiments, the cyclic dinucleotide STING agonist is administered once every 7 to 10 days for 21 to 30 days, then optionally once every 21 days for up to 735 days. In still further embodiments, the cyclic dinucleotide STING agonist is administered once every 7 days for 21 days, then optionally once every 21 days for up to 735 days. In additional embodiments, the cyclic dinucleotide STING agonist is administered once every 21 days for 63 days, then optionally once every 21 days for up to 735 days. In specific embodiments of the foregoing, the CDN STING agonist is administered at least three times.

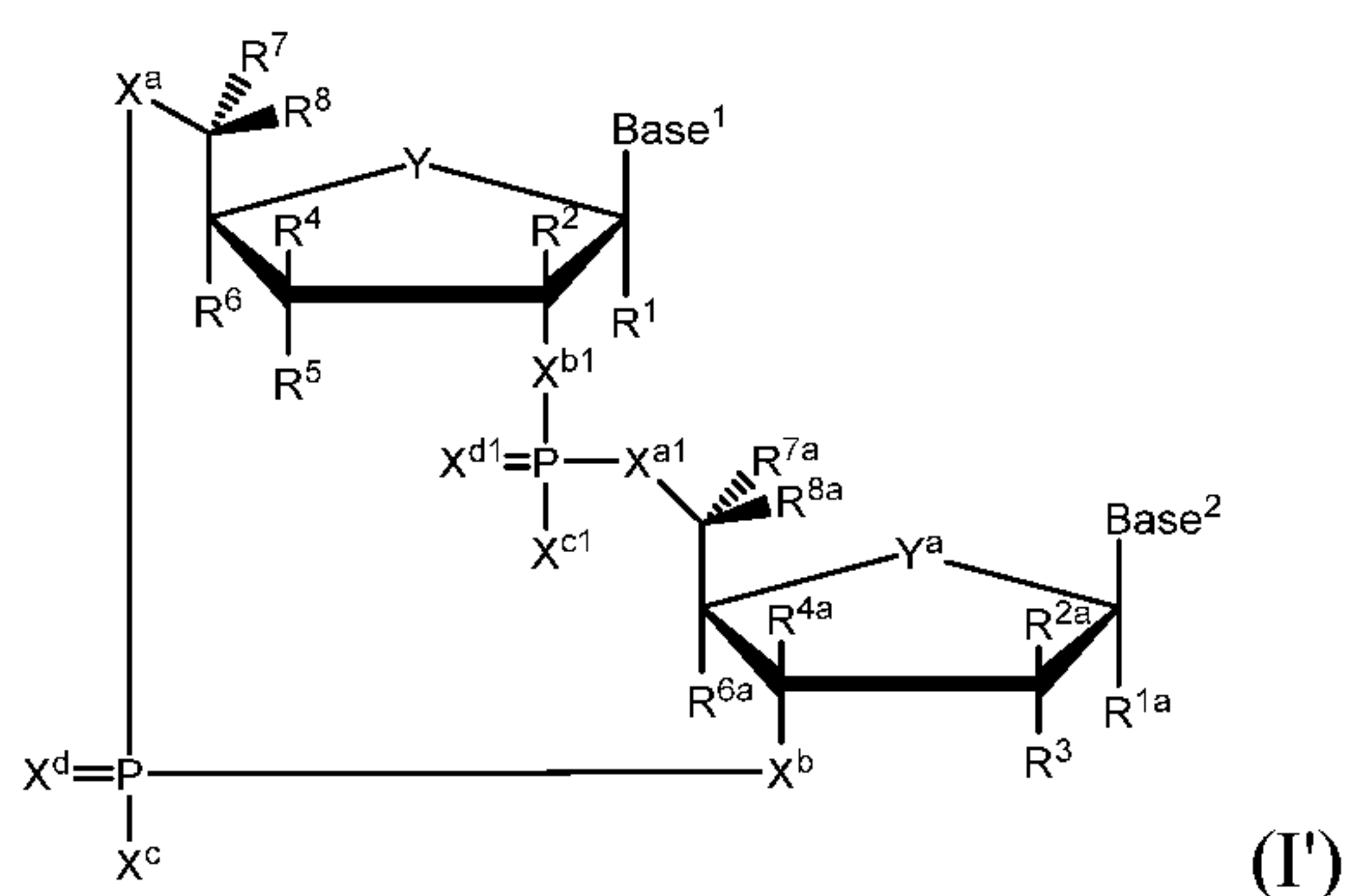
[0226] Additionally, the present disclosure relates to methods of treating a cell-proliferation disorder, said method comprising administering to a subject in need thereof a combination therapy that comprises (a) a PD-1 antagonist; and (b) a cyclic dinucleotide STING agonist; wherein the cell-proliferation disorder is cancer. In specific embodiments, the cancer occurs as one or more solid tumors or lymphomas. In further specific embodiments, the cancer is selected from the group consisting of advanced or metastatic solid tumors and lymphomas. In still further specific embodiments, the cancer is selected from the group consisting of malignant melanoma, head and neck squamous cell carcinoma, breast adenocarcinoma, and lymphomas. In additional embodiments, the lymphoma is selected from the group consisting of diffuse large B-cell lymphoma, follicular lymphoma, mantle cell lymphoma, small lymphocytic lymphoma, mediastinal large B-cell lymphoma, splenic marginal zone B-cell lymphoma, extranodal marginal zone B-cell lymphoma of mucosa-associated lymphoid tissue (malt), nodal marginal zone B-cell lymphoma, lymphoplasmacytic lymphoma, primary effusion lymphoma, Burkitt lymphoma, anaplastic large cell lymphoma (primary cutaneous type), anaplastic large cell lymphoma (systemic type), peripheral T-cell lymphoma, angioimmunoblastic T-cell lymphoma, adult T-cell lymphoma/leukemia, nasal type extranodal NK/T-cell lymphoma, enteropathy-

associated T-cell lymphoma, gamma/delta hepatosplenic T-cell lymphoma, subcutaneous panniculitis-like T-cell lymphoma, mycosis fungoides, and Hodgkin lymphoma. In particular embodiments, the cell-proliferation disorder is a cancer that has metastasized, for example, a liver metastases from colorectal cancer. In additional embodiments, the cell-proliferation disorder is a cancer is classified as stage III cancer or stage IV cancer. In instances of these

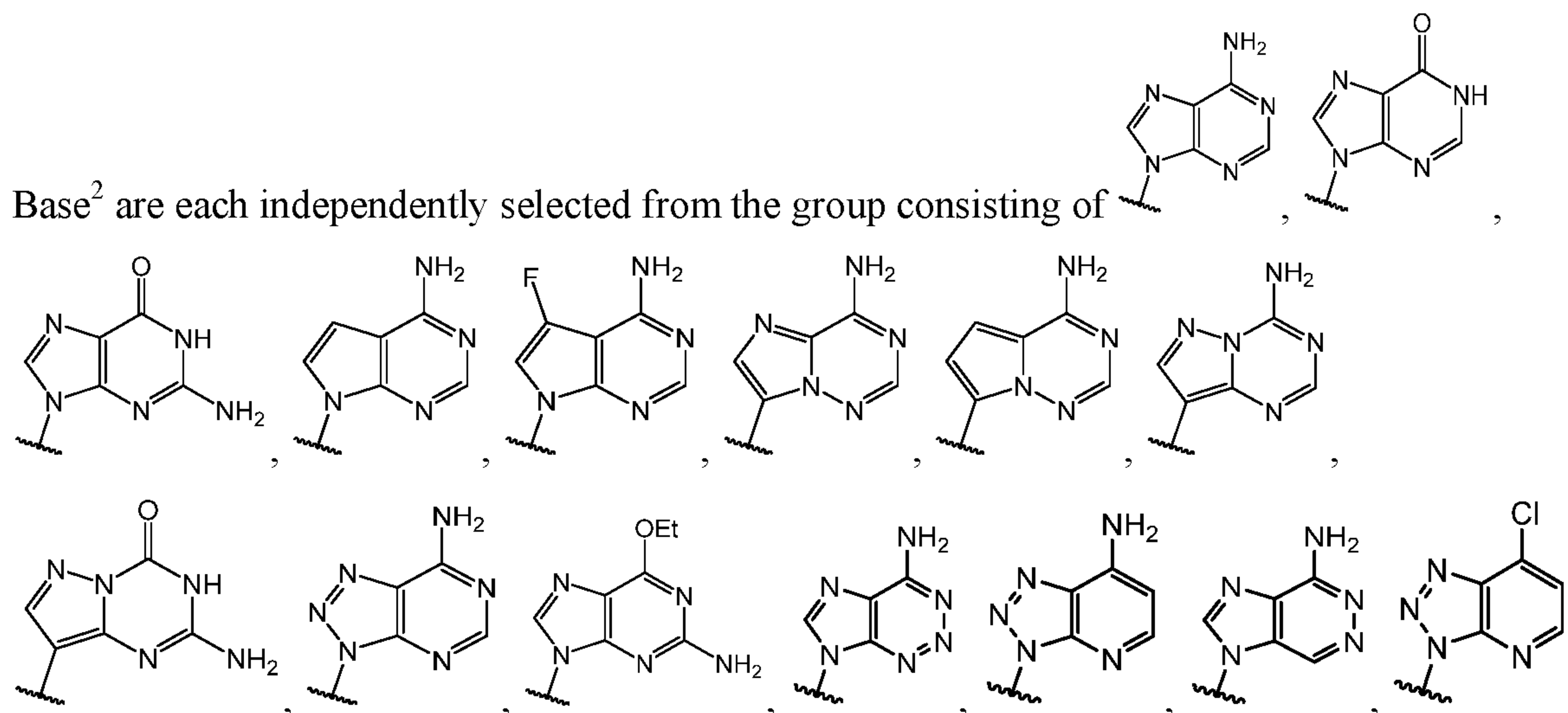
[0227] In embodiments of the methods disclosed herein, the PD-1 antagonist is an anti-PD-1 monoclonal antibody. In particular aspects of these embodiments, the PD-1 antagonist is selected from the group consisting of nivolumab, pembrolizumab, pidilizumab, and AMP-224.

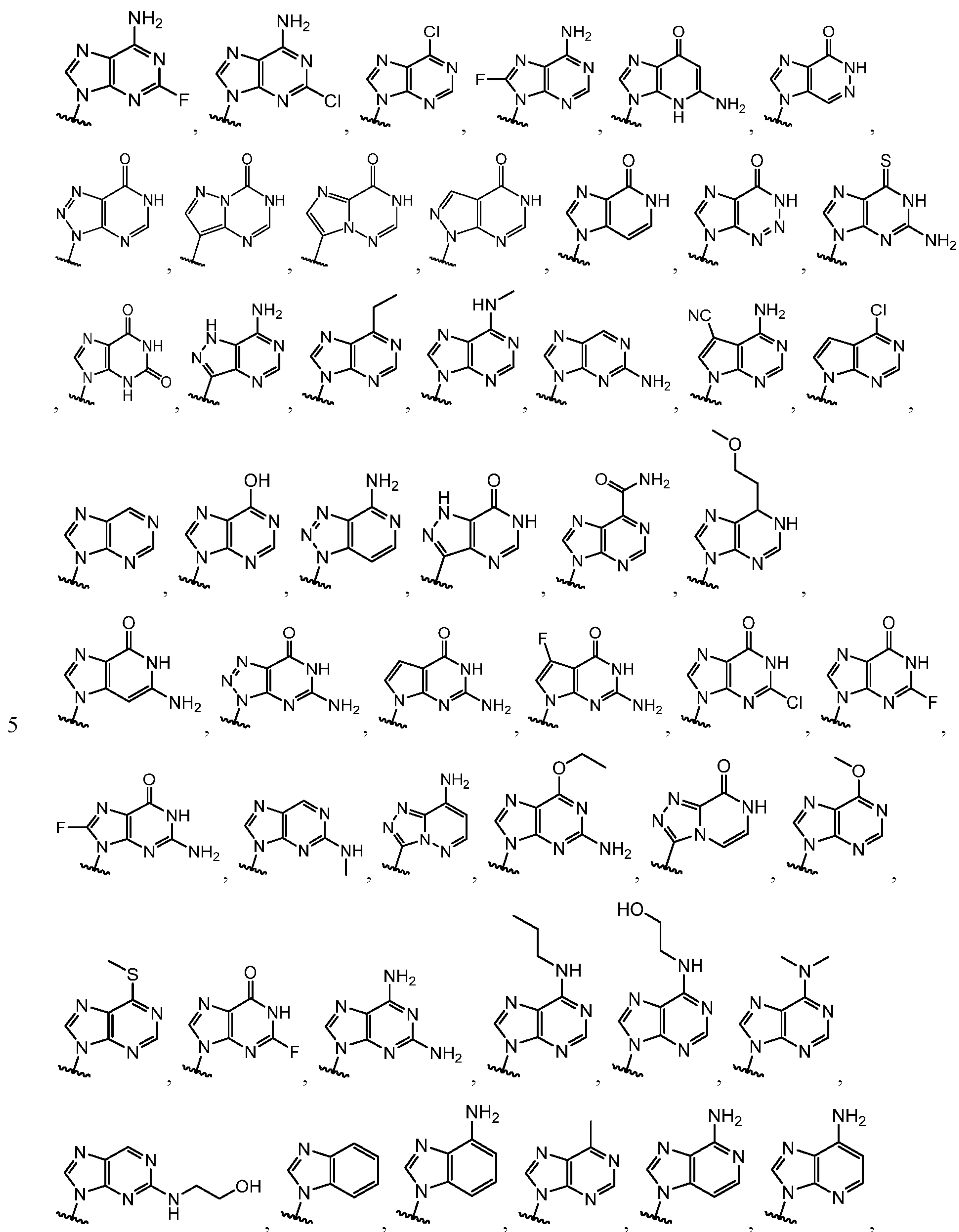
10 In specific aspects of these embodiments, the PD-1 antagonist is selected from nivolumab and pembrolizumab. In a more specific aspect, the PD-1 antagonist is nivolumab. In a further specific aspect, the PD-1 antagonist is pembrolizumab.

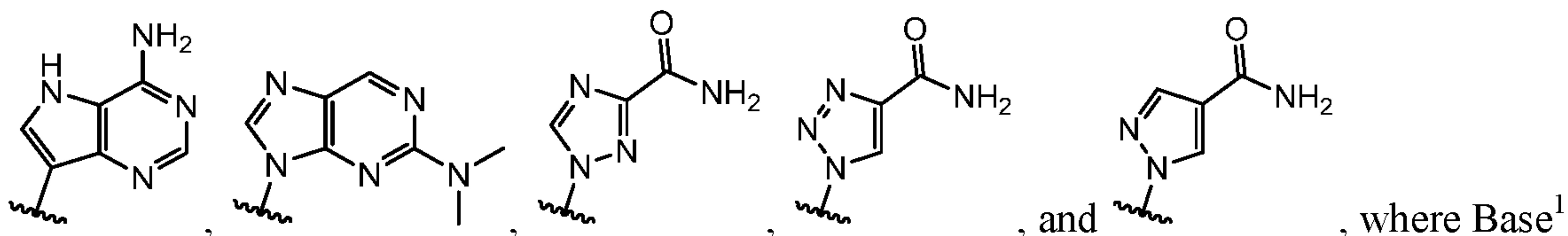
[0228] In embodiments of the methods disclosed herein, the cyclic dinucleotide STING agonist is selected from compounds of formula (I'):



or a pharmaceutically acceptable salt, hydrate, solvate, or prodrug thereof, wherein Base¹ and

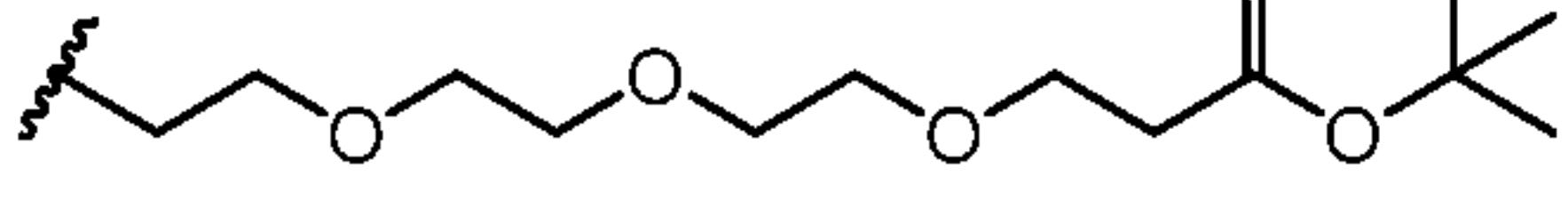
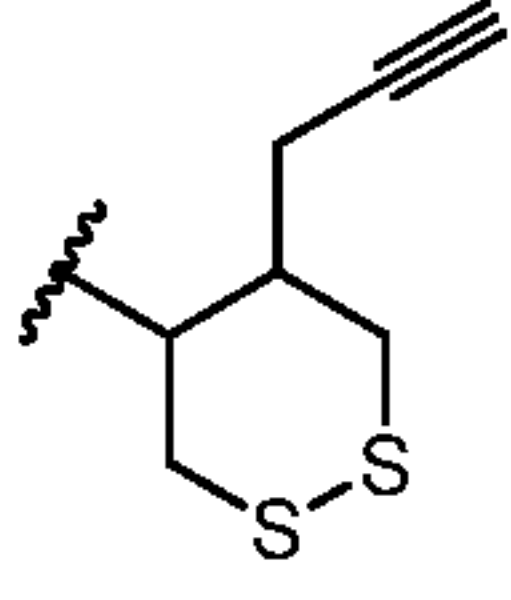






and Base² each may be independently substituted by 0-3 substituents R¹⁰, where each R¹⁰ is independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂; Y and Y^a are each independently selected from the group consisting of -O- and -S-; X^a and X^{a1} are each independently selected from the group consisting of O, and S; X^b and X^{b1} are each independently selected from the group consisting of O, and S; X^c and X^{c1} are each independently selected from the group consisting of OR⁹, SR⁹, and NR⁹R⁹; X^d and X^{d1} are each independently selected from the group consisting of O and S; R¹ and R^{1a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl, where said R¹ and R^{1a} C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R² and R^{2a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl, where said R² and R^{2a} C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R³ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl, where said R³ C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁴ and R^{4a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁₋₆ alkyl, C₂₋₆ alkenyl, C₂₋₆ alkynyl, C₁₋₆ haloalkyl, C₂₋₆ haloalkenyl, C₂₋₆ haloalkynyl, -O-C₁₋₆ alkyl, -O-C₂₋₆ alkenyl, and -O-C₂₋₆ alkynyl, where said R⁴ and

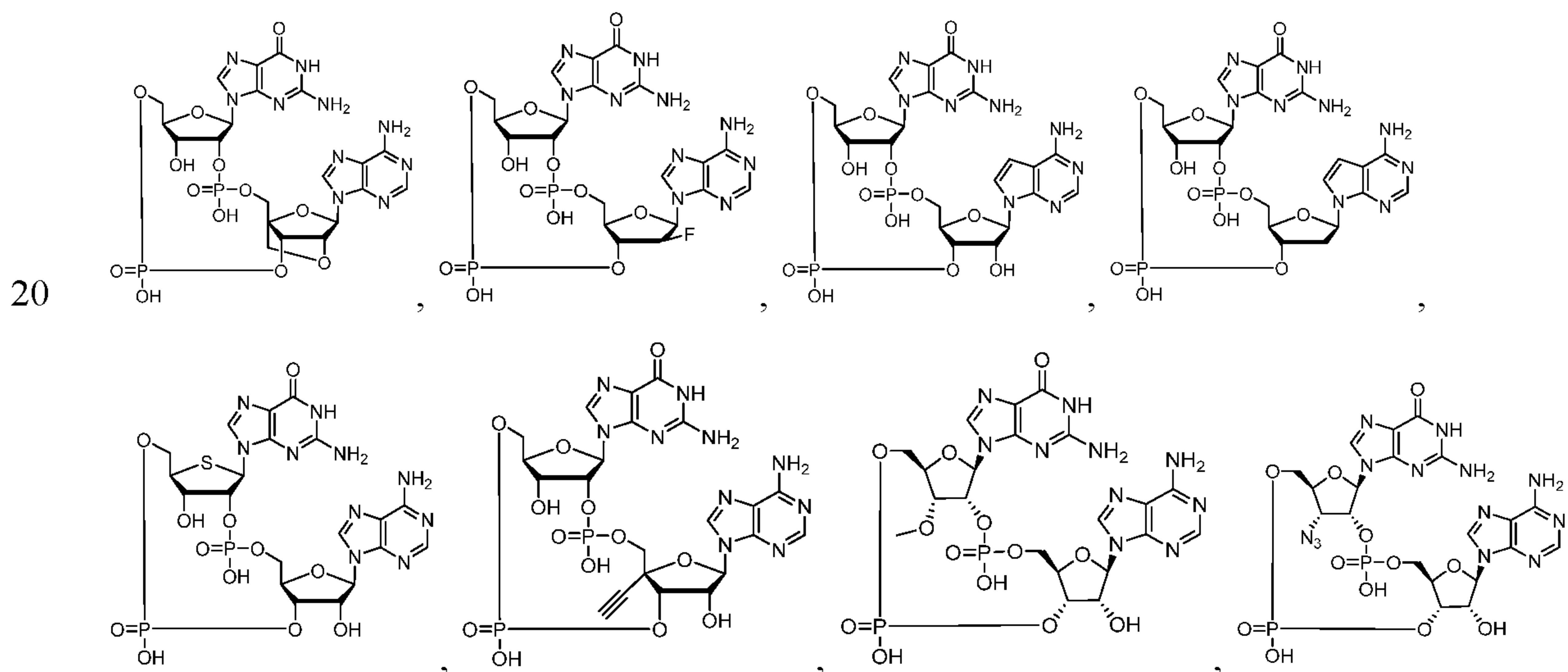
R^{4a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁵ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, NH₂, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁵ C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, NR⁹R⁹, and N₃; R⁶ and R^{6a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁶ and R^{6a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁷ and R^{7a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁷ and R^{7a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; R⁸ and R^{8a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁸ and R^{8a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃; each R⁹ is independently selected

from the group consisting of H, C₁-C₂₀ alkyl, , and ,

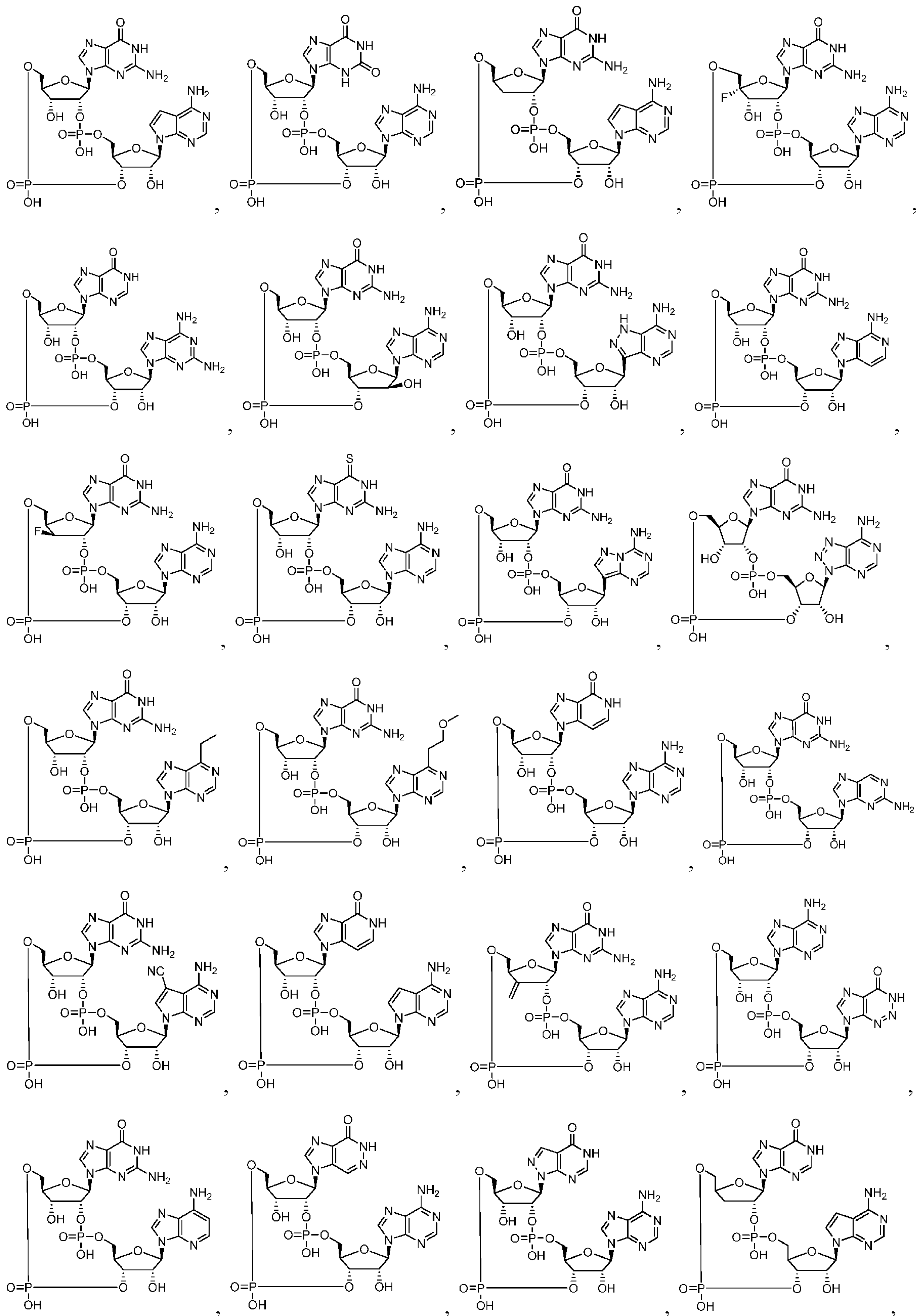
where each R⁹ C₁-C₂₀ alkyl is optionally substituted by 0 to 3 substituents independently selected from the group consisting of OH, -O-C₁-C₂₀ alkyl, -S-C(O)C₁-C₆ alkyl, and -C(O)OC₁-C₆ alkyl; optionally R^{1a} and R³ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene,

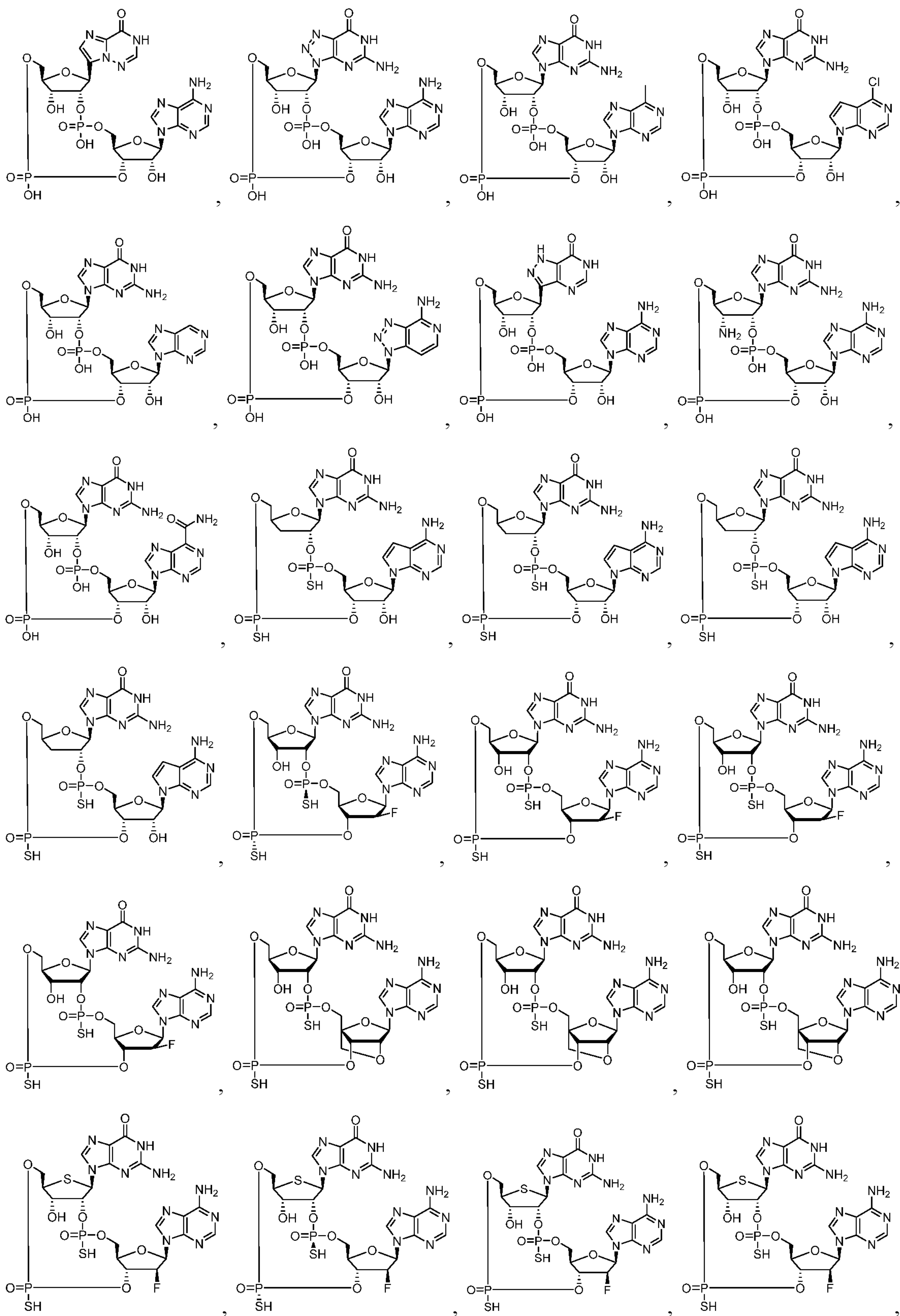
-O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R^{1a} and R³ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position; optionally R^{2a} and R³ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R^{2a} and R³ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position; optionally R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R³ and R^{6a} are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R³ position; optionally R⁴ and R⁵ are connected to form are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, C₂-C₆ alkynylene, -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R⁴ and R⁵ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R⁵ position; optionally R⁵ and R⁶ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, such that where R⁵ and R⁶ are connected to form -O-C₁-C₆ alkylene, -O-C₂-C₆ alkenylene, or -O-C₂-C₆ alkynylene, said O is bound at the R⁵ position; optionally R⁷ and R⁸ are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, or C₂-C₆ alkynylene; and optionally R^{7a} and R^{8a} are connected to form C₁-C₆ alkylene, C₂-C₆ alkenylene, or C₂-C₆ alkynylene.

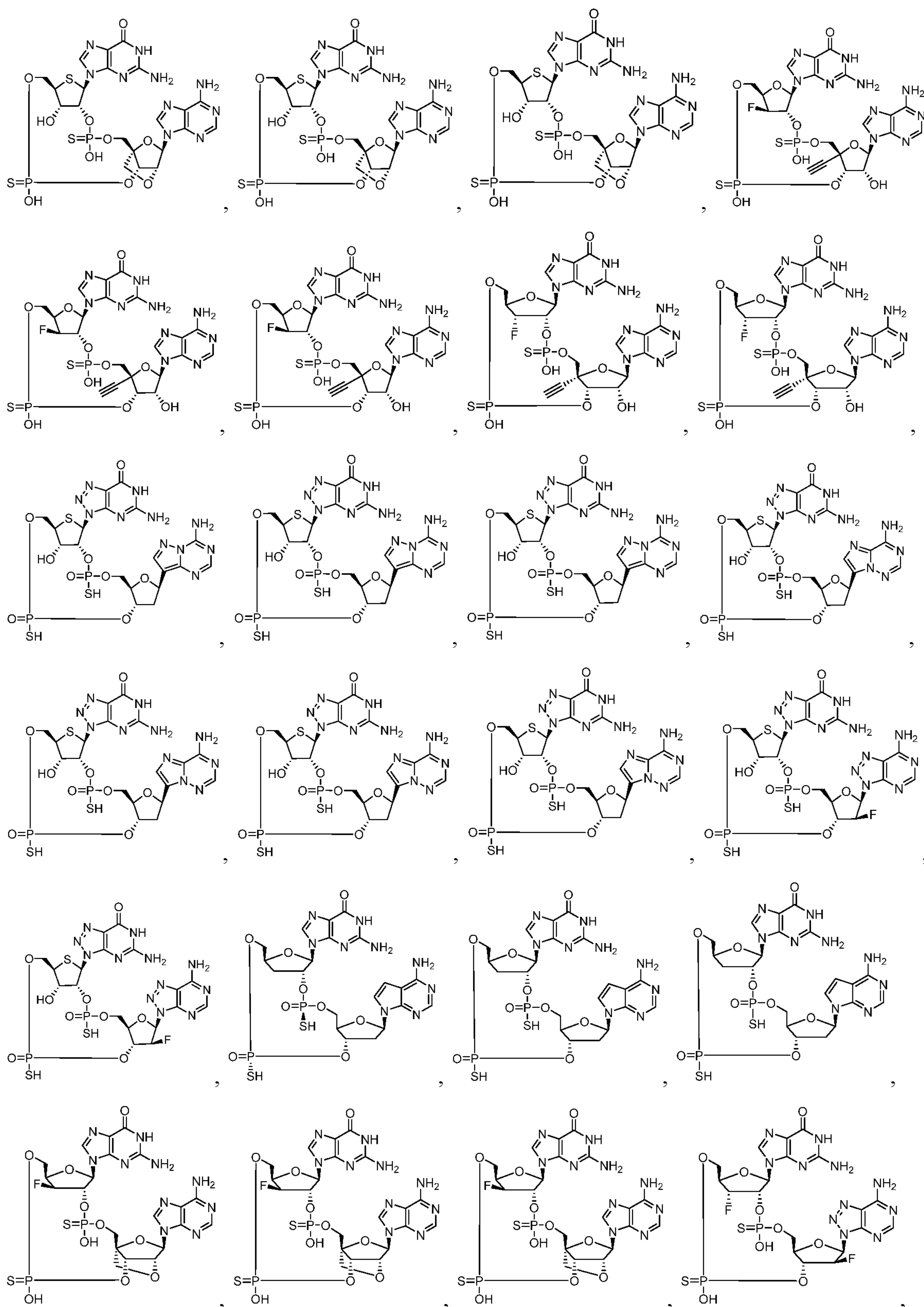
[0229] In instances of these embodiments, the cyclic dinucleotide STING agonist is selected from the group consisting of:

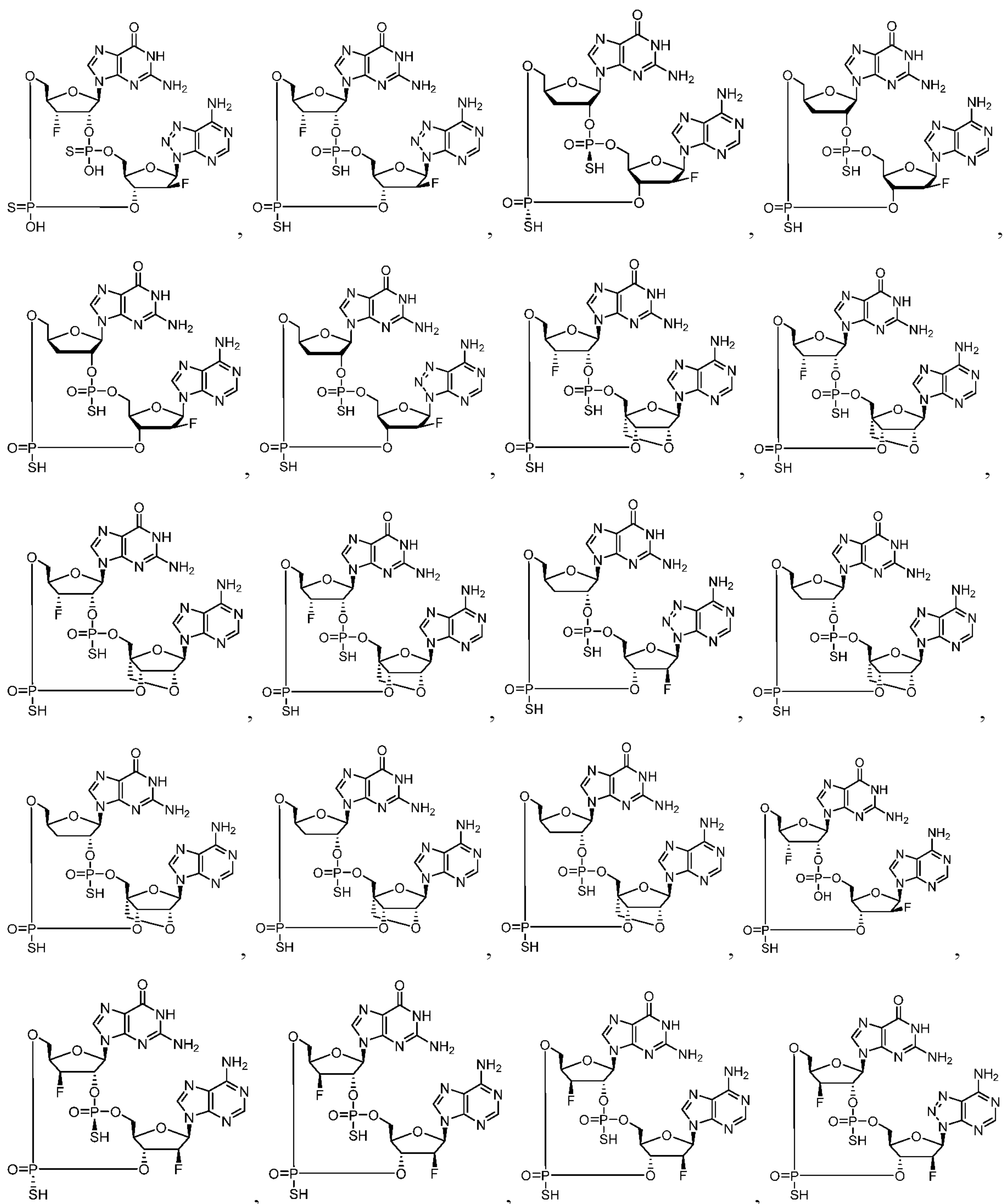




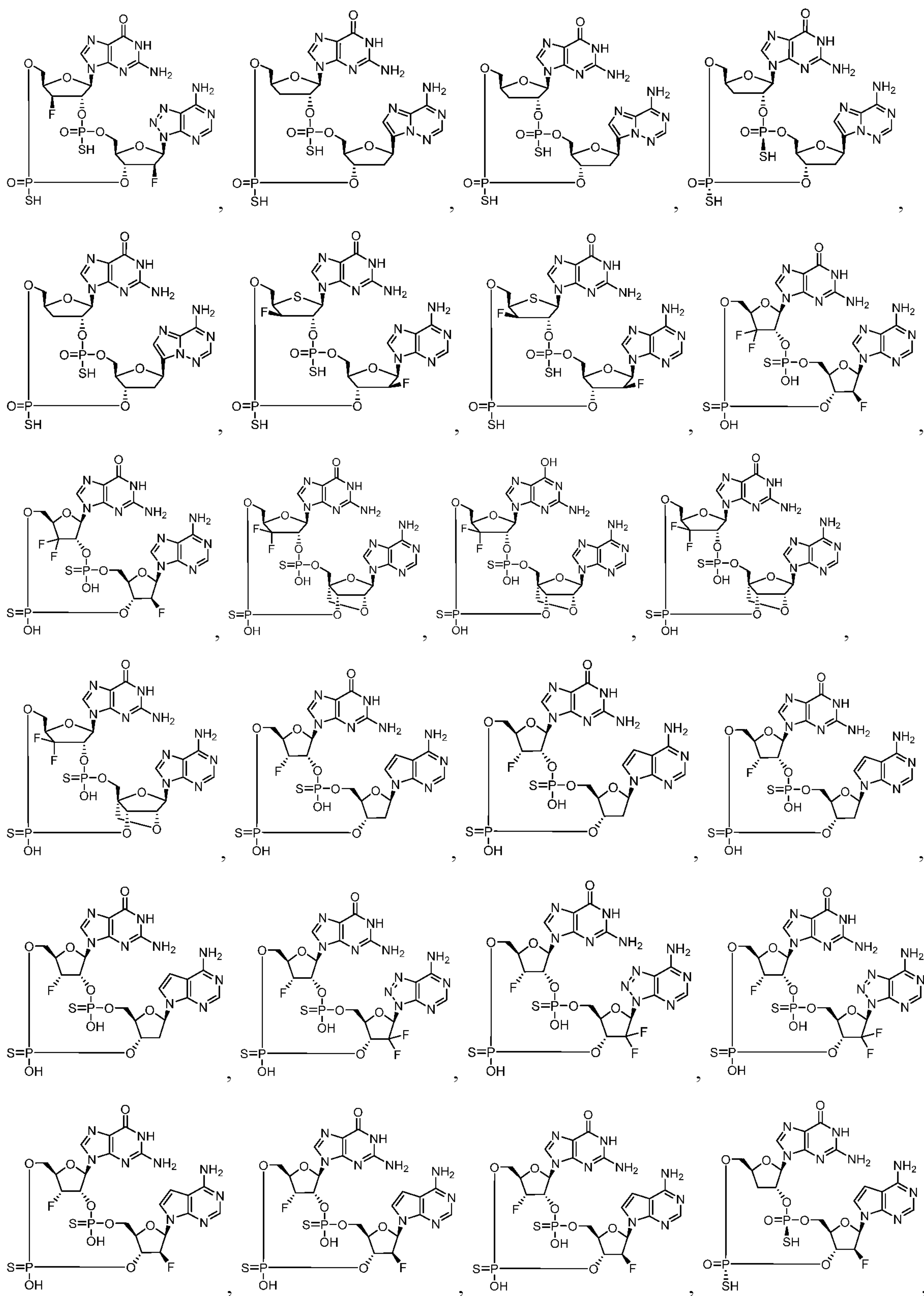


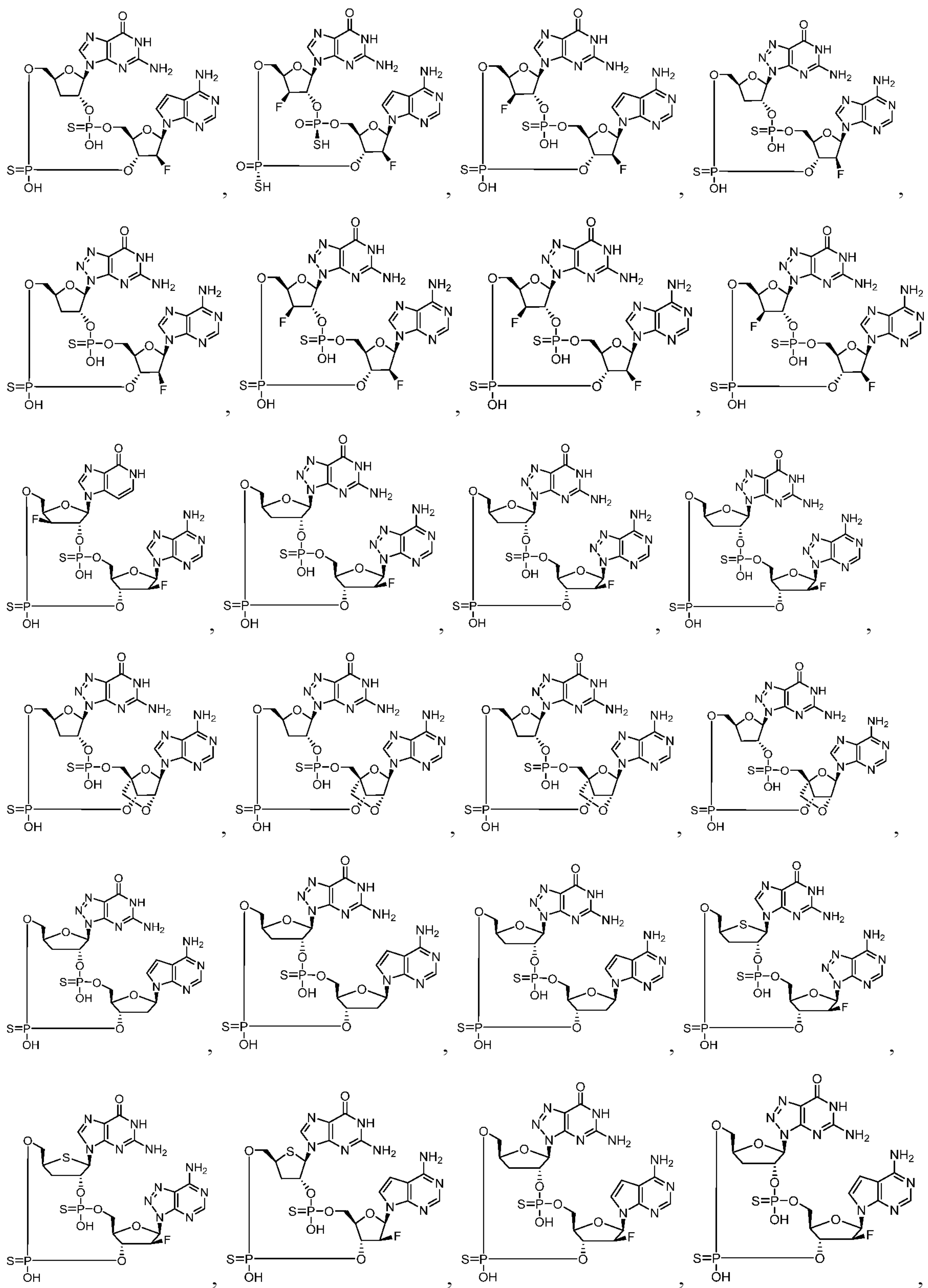


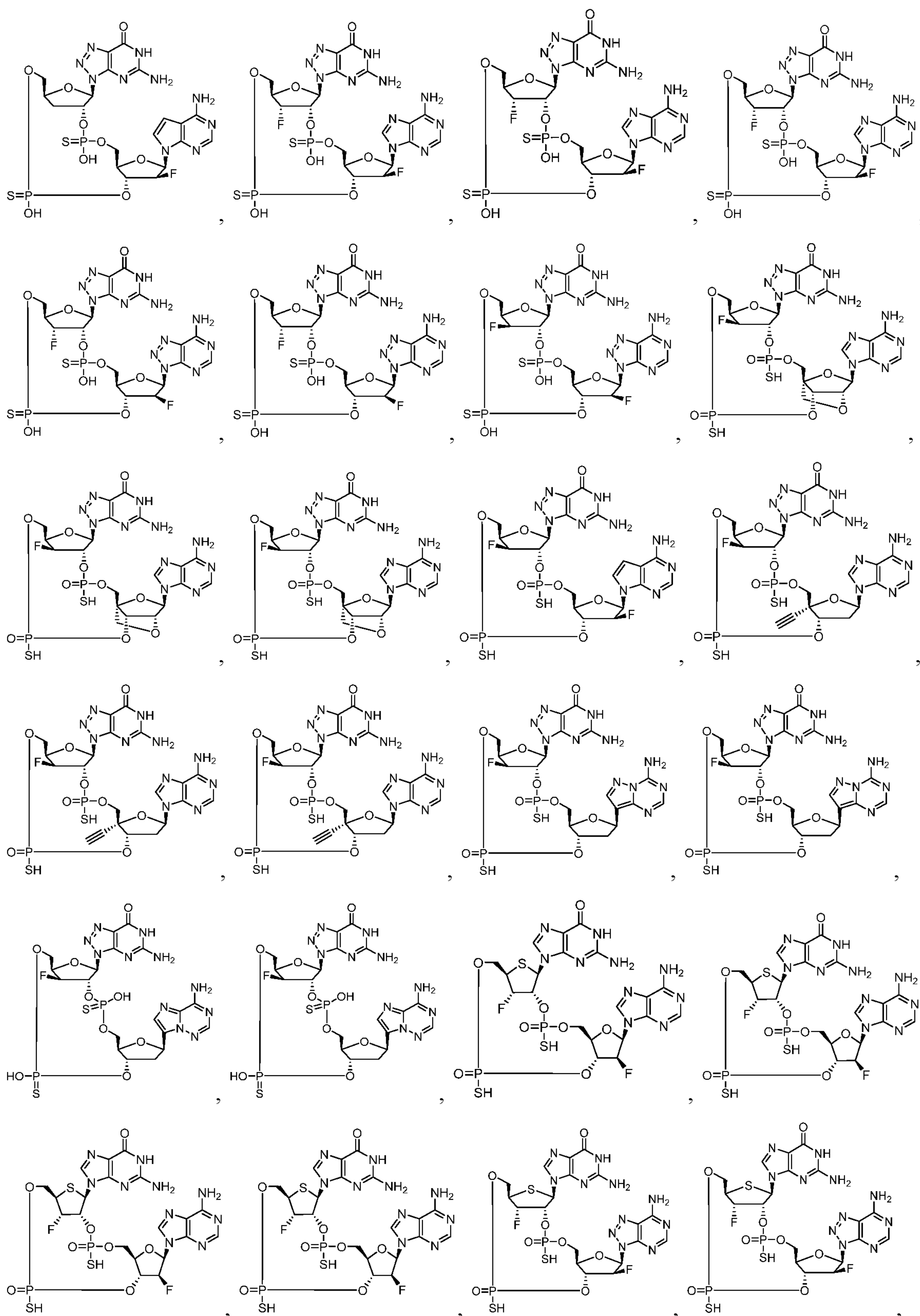


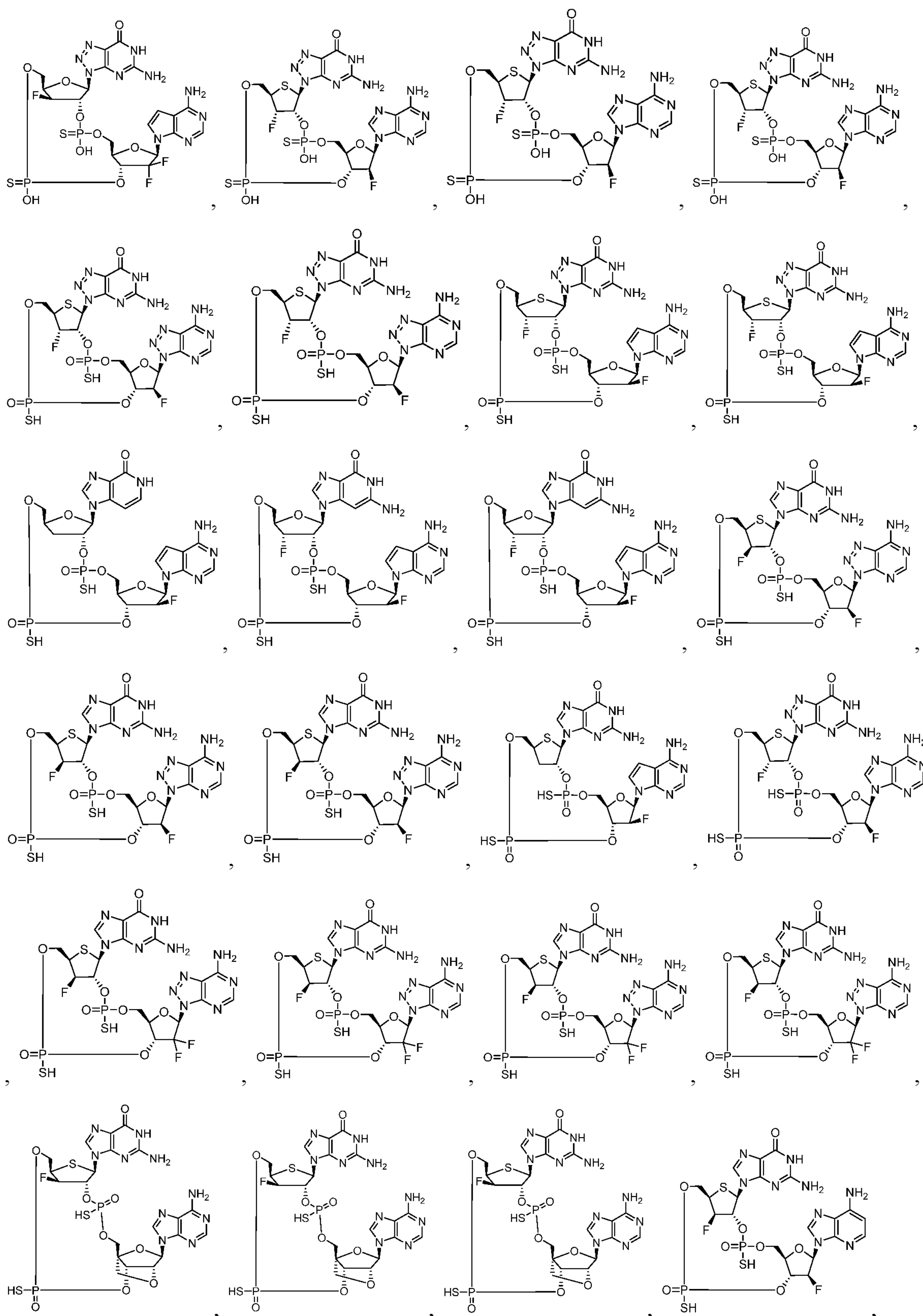


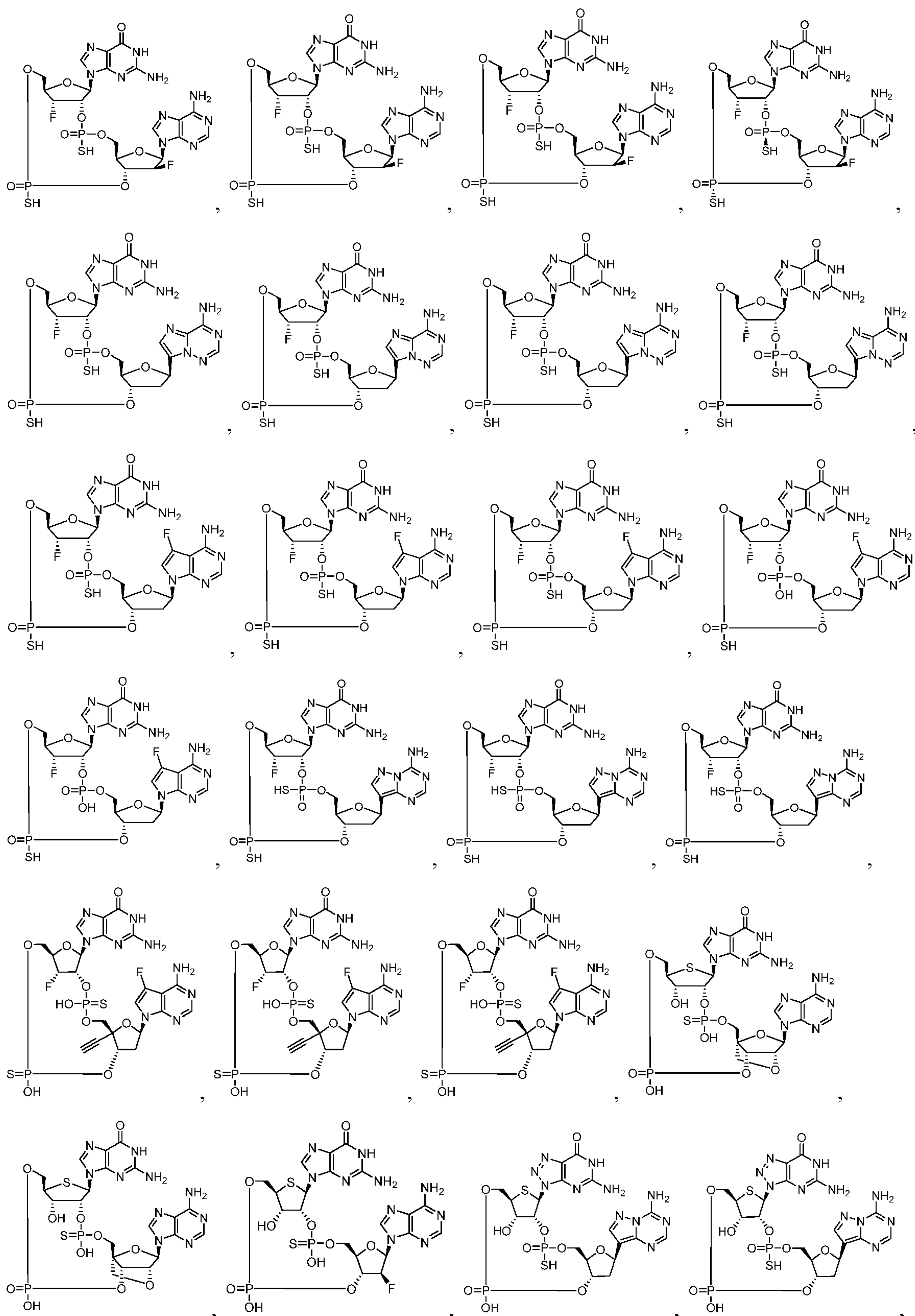
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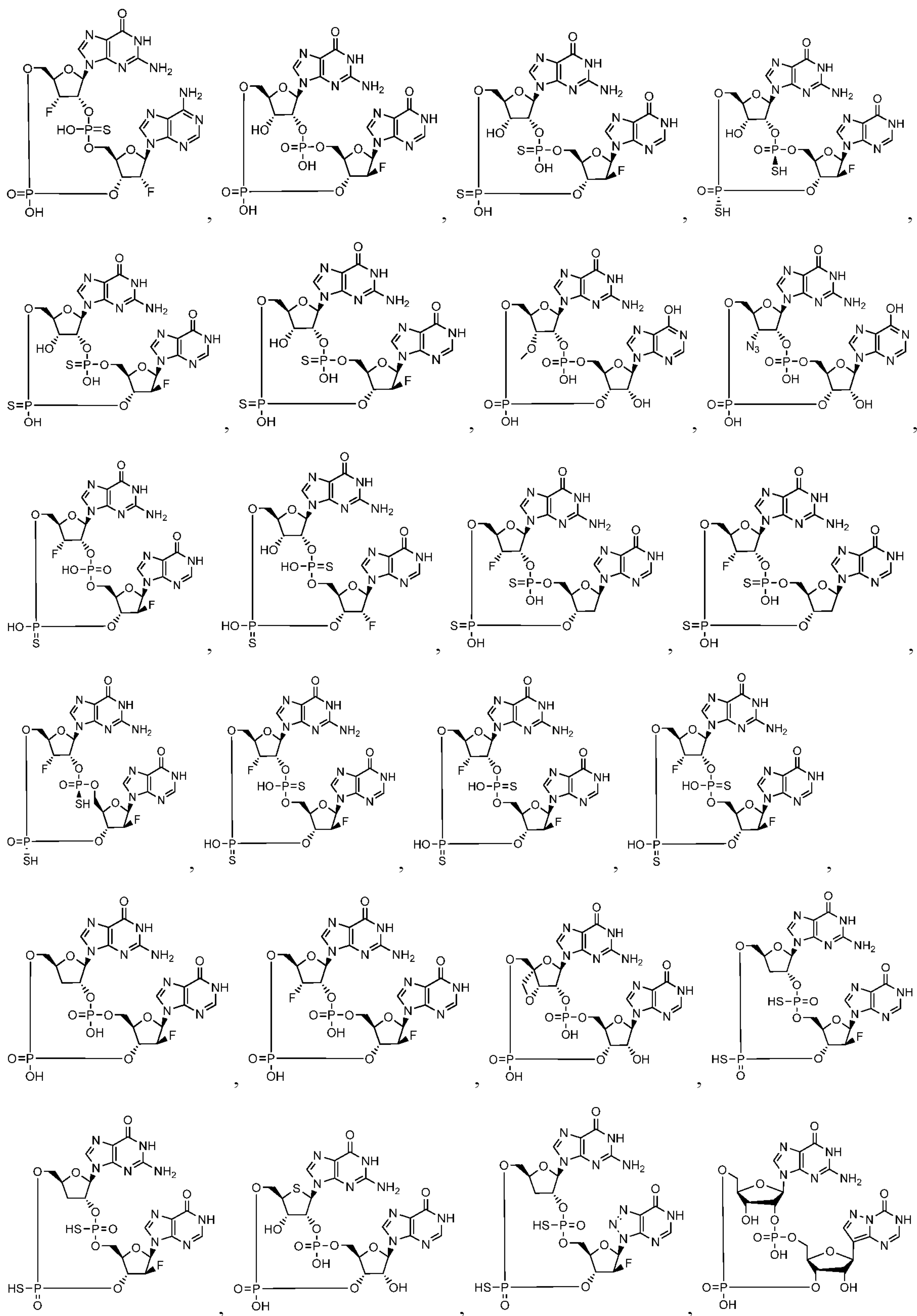


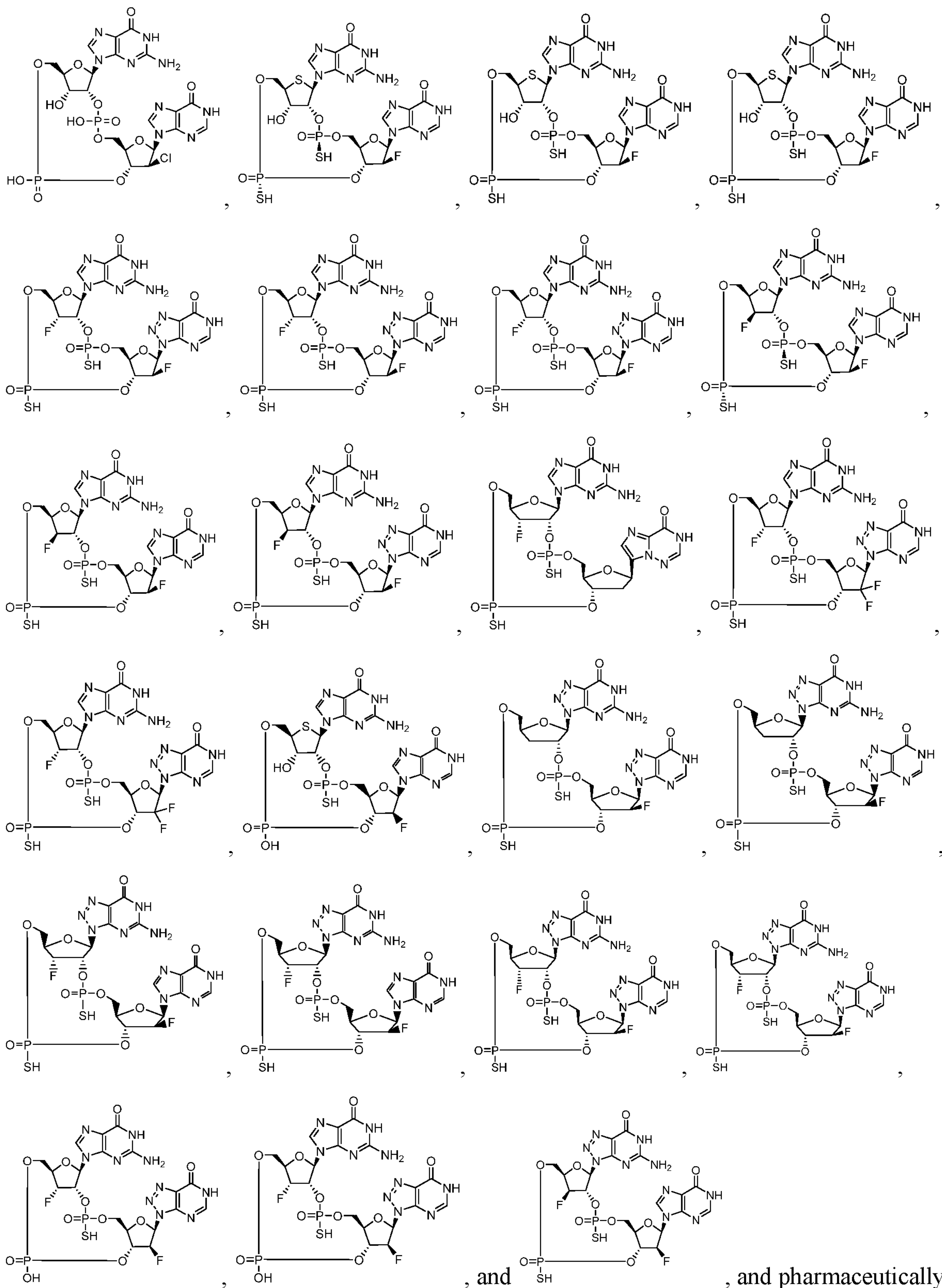












acceptable salts thereof.

[0230] In embodiments of the methods disclosed herein, the PD-1 antagonist is administered by intravenous infusion, and the cyclic dinucleotide STING agonist is orally, by intravenous infusion, by intertumoral injection or by subcutaneous injection.

[0231] In embodiments of the methods disclosed herein, the PD-1 antagonist is administered prior to administration of the cyclic dinucleotide STING agonist. In alternative embodiments of the methods disclosed herein, the cyclic dinucleotide STING agonist is administered prior to administration of the PD-1 antagonist.

[0232] In embodiments of the methods disclosed herein, the PD-1 antagonist is administered at a dose of 200mg; and the cyclic dinucleotide STING agonist is administered at a dose of from 10µg to 3000µg. In aspects of such embodiments, the cyclic dinucleotide STING agonist is administered at a dose of from 10µg to 270µg.

[0233] Additional embodiments of the disclosure include the pharmaceutical compositions, combinations, uses and methods set forth in above, wherein it is to be understood that each embodiment may be combined with one or more other embodiments, to the extent that such a combination is consistent with the description of the embodiments. It is further to be understood that the embodiments provided above are understood to include all embodiments, including such embodiments as result from combinations of embodiments.

GENERAL METHODS

[0234] Standard methods in molecular biology are described Sambrook, Fritsch and Maniatis (1982 & 1989 2nd Edition, 2001 3rd Edition) *Molecular Cloning, A Laboratory Manual*, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY; Sambrook and Russell (2001) *Molecular Cloning, 3rd ed.*, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY; Wu (1993) *Recombinant DNA*, Vol. 217, Academic Press, San Diego, CA). Standard methods also appear in Ausbel, *et al.* (2001) *Current Protocols in Molecular Biology, Vols. 1-4*, John Wiley and Sons, Inc. New York, NY, which describes cloning in bacterial cells and DNA mutagenesis (Vol. 1), cloning in mammalian cells and yeast (Vol. 2), glycoconjugates and protein expression (Vol. 3), and bioinformatics (Vol. 4).

[0235] Methods for protein purification including immunoprecipitation, chromatography, electrophoresis, centrifugation, and crystallization are described (Coligan, *et al.* (2000) *Current Protocols in Protein Science, Vol. 1*, John Wiley and Sons, Inc., New York). Chemical analysis, chemical modification, post-translational modification, production of fusion proteins, glycosylation of proteins are described (*see, e.g.,* Coligan, *et al.* (2000) *Current Protocols in*

Protein Science, Vol. 2, John Wiley and Sons, Inc., New York; Ausubel, *et al.* (2001) *Current Protocols in Molecular Biology*, Vol. 3, John Wiley and Sons, Inc., NY, NY, pp. 16.0.5-16.22.17; Sigma-Aldrich, Co. (2001) *Products for Life Science Research*, St. Louis, MO; pp. 45-89; Amersham Pharmacia Biotech (2001) *BioDirectory*, Piscataway, N.J., pp. 384-391).

- 5 Production, purification, and fragmentation of polyclonal and monoclonal antibodies are described (Coligan, *et al.* (2001) *Current Protocols in Immunology*, Vol. 1, John Wiley and Sons, Inc., New York; Harlow and Lane (1999) *Using Antibodies*, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY; Harlow and Lane, *supra*). Standard techniques for characterizing ligand/receptor interactions are available (*see, e.g.,* Coligan, *et al.* (2001) *Current*
10 *Protocols in Immunology*, Vol. 4, John Wiley, Inc., New York).

[0236] Monoclonal, polyclonal, and humanized antibodies can be prepared (*see, e.g.,* Sheperd and Dean (eds.) (2000) *Monoclonal Antibodies*, Oxford Univ. Press, New York, NY; Kontermann and Dubel (eds.) (2001) *Antibody Engineering*, Springer-Verlag, New York; Harlow and Lane (1988) *Antibodies A Laboratory Manual*, Cold Spring Harbor Laboratory
15 Press, Cold Spring Harbor, NY, pp. 139-243; Carpenter, *et al.* (2000) *J. Immunol.* 165:6205; He, *et al.* (1998) *J. Immunol.* 160:1029; Tang *et al.* (1999) *J. Biol. Chem.* 274:27371-27378; Baca *et al.* (1997) *J. Biol. Chem.* 272:10678-10684; Chothia *et al.* (1989) *Nature* 342:877-883; Foote and Winter (1992) *J. Mol. Biol.* 224:487-499; U.S. Pat. No. 6,329,511).

[0237] An alternative to humanization is to use human antibody libraries displayed on
20 phage or human antibody libraries in transgenic mice (Vaughan *et al.* (1996) *Nature Biotechnol.* 14:309-314; Barbas (1995) *Nature Medicine* 1:837-839; Mendez *et al.* (1997) *Nature Genetics* 15:146-156; Hoogenboom and Chames (2000) *Immunol. Today* 21:371-377; Barbas *et al.* (2001) *Phage Display: A Laboratory Manual*, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York; Kay *et al.* (1996) *Phage Display of Peptides and Proteins: A Laboratory*
25 *Manual*, Academic Press, San Diego, CA; de Bruin *et al.* (1999) *Nature Biotechnol.* 17:397-399).

[0238] Purification of antigen is not necessary for the generation of antibodies. Animals can be immunized with cells bearing the antigen of interest. Splenocytes can then be isolated from the immunized animals, and the splenocytes can fused with a myeloma cell line to produce
30 a hybridoma (*see, e.g.,* Meyaard *et al.* (1997) *Immunity* 7:283-290; Wright *et al.* (2000) *Immunity* 13:233-242; Preston *et al., supra*; Kaithamana *et al.* (1999) *J. Immunol.* 163:5157-5164).

[0239] Methods for flow cytometry, including fluorescence activated cell sorting (FACS), are available (*see, e.g., Owens, et al. (1994) Flow Cytometry Principles for Clinical Laboratory Practice*, John Wiley and Sons, Hoboken, NJ; Givan (2001) *Flow Cytometry*, 2nd ed.; Wiley-Liss, Hoboken, NJ; Shapiro (2003) *Practical Flow Cytometry*, John Wiley and Sons, Hoboken, NJ). Fluorescent reagents suitable for modifying nucleic acids, including nucleic acid primers and probes, polypeptides, and antibodies, for use, e.g., as diagnostic reagents, are available (Molecular Probes (2003) *Catalogue*, Molecular Probes, Inc., Eugene, OR; Sigma-Aldrich (2003) *Catalogue*, St. Louis, MO).

[0240] Standard methods of histology of the immune system are described (*see, e.g., Muller-Harmelink (ed.) (1986) Human Thymus: Histopathology and Pathology*, Springer Verlag, New York, NY; Hiatt, *et al.* (2000) *Color Atlas of Histology*, Lippincott, Williams, and Wilkins, Phila, PA; Louis, *et al.* (2002) *Basic Histology: Text and Atlas*, McGraw-Hill, New York, NY).

[0241] Software packages and databases for determining, e.g., antigenic fragments, leader sequences, protein folding, functional domains, glycosylation sites, and sequence alignments, are available (*see, e.g., GenBank*, Vector NTI® Suite (Informax, Inc., Bethesda, MD); GCG Wisconsin Package (Accelrys, Inc., San Diego, CA); DeCypher® (TimeLogic Corp., Crystal Bay, Nevada); Menne, *et al.* (2000) *Bioinformatics* 16: 741-742; Menne, *et al.* (2000) *Bioinformatics Applications Note* 16:741-742; Wren, *et al.* (2002) *Comput. Methods Programs Biomed.* 68:177-181; von Heijne (1983) *Eur. J. Biochem.* 133:17-21; von Heijne (1986) *Nucleic Acids Res.* 14:4683-4690).

Advanced MC38 Mouse Syngenic Tumor Model

[0242] Synergistic tumor models are recognized to be appropriate models to evaluate anti-tumor efficiency of agents that target specific molecules, pathways, or cell types and to provide mechanistic rationale that targeting similar specific molecules, pathways, or cell types in human tumors will lead to favorable clinical outcomes. The mouse syngeneic MC38 tumor model is a mouse colon adenocarcinoma cell line that was established by carcinogenic induction of tumors in the C57BL/6 background. This cell line is considered immunogenic and is responsive to immune modulation. It is generally injected subcutaneously (SC) to evaluate tumor growth and response to treatment. Specifically, each animal is inoculated in the right lower flank with a SC dose of 1×10^6 MC38 colon adenocarcinoma cells in 100 μ L of serum-free Dulbecco's modified Eagle's medium. Tumor progression is monitored by measuring tumor volume using Vernier calipers. *See T.H. Corbett et al., Tumor Induction Relationships in*

Development of Transplantable Cancers of the Colon in Mice for Chemotherapy Assays, with a Note on Carcinogen Structure, 35(9) Cancer Res. 2434-2439 (September 1, 1975).

Anti-Mouse PD-1 Antibody

[0243] In the Example below, the anti-tumor effects of selected CDN STING agonists in combination with an anti-mouse PD1 antibody are evaluated in mouse syngeneic tumor models. Anti-tumor activity (tumor growth inhibition, tumor regression) is observed on treatment of mouse syngeneic tumors with the combination. Both mouse and human tumor infiltrating T cells express high levels of PD-1, associated with what is referred to as an “exhausted phenotype” (See Y. Jiang *et al.*, “T-cell exhaustion in the tumor microenvironment”, *Cell Death and Disease* **2015**, 6, e1792). Induction of anti-tumor efficacy in mouse syngeneic tumor models following treatment with anti-mouse PD-1 antibodies provides a mechanistic rationale that treatment of cancer patients with anti-human PD-1 antibodies will induce anti-tumor efficacy (See S. Hu-Lieskovan *et al.*, “Improved antitumor activity of immunotherapy with BRAF and MEK inhibitors in BRAF(V600E) melanoma”, *Sci. Transl. Med.* 2015 Mar 18; 7(279):279ra41; C.D. Pham *et al.*, “Differential immune microenvironments and response to immune checkpoint blockade among molecular subtypes of murine medulloblastoma”, *Clin. Cancer Res.* 2016 Feb 1; 22(3):582-595; S. Budhu *et al.*, “The importance of animal models in tumor immunity and immunotherapy”, *Curr. Opin. Genet. Dev.* **2014**, 24, 46–51). Suitable anti-mouse PD-1 antibodies that may be used include muDX400 (Merck), *InVivoMAb* and *InVivoPlusMAb* anti-mouse PD-1 clone J43 (commercially available from BioXCell as catalog number BE0033-2), *InVivoMAb* anti-mouse PD-1 clone 29F.1A12 (commercially available from BioXCell as catalog number BE0273), and *InVivoMAb* and *InVivoPlusMAb* anti-mouse PD-1 clone RMP1-14 (commercially available from BioXCell as catalog number BE0146).

EXAMPLES

Example 1: Anti-Tumor Efficacy of a CDN STING Agonist in Combination with an Anti-PD-1 Antibody in Advanced MC38 Mouse Syngenic Tumor Model

[0244] To assess the combination anti-tumor efficacy of a CDN STING agonist and anti-mouse PD-1 antibody muDX400 in the advanced MC38 mouse syngeneic tumor model, a cohort of 8-12 week old female C57Bl/6 mice are implanted with 1×10^6 MC38 cells. When the tumors reach a median size of approximately 350 mm^3 , the animals are randomized into 6 treatment groups of 10 mice per group:

Treatment Group A: PBS and mIgG1 (5mg/kg)

Treatment Group B: PBS and anti-PD-1 antibody muDX400 (5mg/kg)

Treatment Group C: CDN STING agonist (5μg) and mIgG1 (5mg/kg)

Treatment Group D: CDN STING agonist (5μg) and anti-PD-1 antibody muDX400

5 (5mg/kg)

[0245] CDN STING agonist are administered intratumorally on every 3 to 7 days for up to 30 days. Antibodies are administered intraperitoneally every 5 days for 5 doses. The study period will be 30 days post initiation of the dosing regimens.

[0246] Tumors on animals in Treatment Group A are anticipated to progress rapidly.

10 The remaining groups are observed for tumor regression and number of CRs. It is anticipated that CDN STING agonist in combination with anti PD-1 muDX400 treatment (Treatment Group D) will demonstrate superior efficacy to single agent treatment groups.

[0247] When the foregoing experiment was conducted with selected combinations as described herein, the combination treatment (Treatment Group D) resulted in significant anti-

15 tumor efficacy compared to Treatment Group A.

Example 2: Clinical Study Evaluating a CDN STING Agonist in Combination with an Anti-PD-1 Antibody in Treatment of Patients with Advanced/Metastatic Solid Tumors or Lymphomas

20 [0248] A Phase I clinical study will be conducted to evaluate, in part, the effects of a combination therapy, consisting of administration of a pembrolizumab intravenous infusion and of a CDN STING agonist as described above intratumoral injection, on advanced or metastatic solid tumors or lymphomas. The study is a non-randomized, 2-arm, multi-site, open-label trial of CDN STING agonist monotherapy and CDN STING agonist in combination with

25 pembrolizumab in subjects with advanced/metastatic solid tumors or lymphomas. CDN STING agonist will be administered intratumorally (IT).

[0249] Unless deemed medically unsafe by the Investigator, all subjects will be required to provide a sample of the tumor to be injected and a sample from a distant site prior to CDN STING agonist administration during screening, as well as on Cycle 3, Day 15. Subjects with

30 amenable lesions at both injected and non-injected sites may undergo an additional optional tumor biopsy on Cycle 6, Day 15 of both the injected lesion and the non-injected lesion. Subjects will undergo a 24-hour observation period following the first dose administration on

Cycle 1, Day 1. Each cycle within the trial is a 21-day cycle. Dosing in the first 3 cycles is once a week (Q1W) and dosing in cycles 4 and beyond is once every 3 weeks (Q3W).

[0250] Dose escalation will proceed based on emerging safety and tolerability data of CDN STING agonist as monotherapy and as combination therapy with pembrolizumab. For each dose level, an assessment will be made of the safety and tolerability data in order to define the next dose level to be tested. Both treatment arms will start with an accelerated titration design (ATD) followed by the modified toxicity probability interval (mTPI) method to identify a maximum tolerated dose (MTD) or maximum administered dose (MAD) of CDN STING agonist alone (Arm 1) or CDN STING agonist in combination with pembrolizumab (Arm 2). Starting with a dose of 10 μ g of CDN STING agonist in single patient cohorts (Arm 1, Part A), the trial will proceed in an ATD up to a dose that meets at least 1 of the following 3 criteria: 1) The 270 μ g cohort is completed, 2) $\geq$ Grade 2 non-disease-related toxicity at any dose level, or 3) Elevation of systemic TNF- α in blood above baseline levels by $\geq$ 3 fold increase for a given subject at any time during the first cycle of CDN STING agonist. Upon completion of the ADT phase by reaching at least one of the above triggering criteria, the monotherapy arm (Arm 1) of the study will proceed to a dose escalation and confirmation phase (Part B), using an mTPI design. In addition, Arm 2 (Part C), the combination therapy arm, will initiate once 2 dose levels within Arm 1 have been cleared by dose-limiting toxicity (DLT) evaluation.

[0251] Starting with a dose that is at least 2 dose levels behind CDN STING agonist monotherapy, CDN STING agonist combination therapy with pembrolizumab (Arm 2 Part C) will begin in single patient cohorts. In Arm 2 Part C, CDN STING agonist combination arm with pembrolizumab, dose escalation will proceed in an ATD up to a dose level which meets at least 1 of the following 3 criteria: 1) The 270 μ g cohort in combination is completed, 2) $\geq$ Grade 2 non-disease-related toxicity at any dose level in combination, or 3) Elevation of systemic TNF- α in blood above baseline levels by $\geq$ 3 fold for a given subject at any time during the first cycle of CDN STING agonist in combination with pembrolizumab. Arm 2 will then proceed to mTPI (Arm 2, Part D) to determine the MTD/MAD of the combination of CDN STING agonist and pembrolizumab.

[0252] Intra-subject dose escalation of CDN STING agonist to the next dose level is permitted only in Arm 1, including Parts A and B. Intrasubject dose escalation will be at the discretion of the Investigator, provided that the subject remains on study after receiving 3 cycles of treatment without $\geq$ Grade 2 toxicity, and provided that the dose escalation has proceeded

beyond the next dose level. Intra-subject dose escalation is not permitted in Arm 2 (Parts C and D).

[0253] During CDN STING agonist dose escalation in both Arm 1 (Parts A and (b) and Arm 2 (Parts C and D), at least 7 days of observation will occur between each of the first 2 subjects at each dose level. Over-enrollment in ATD up to 3 subjects per cohort is permitted, provided that the first 2 subjects will receive CDN STING agonist treatment at least 7 days apart. Dose escalation of CDN STING agonist to determine the MTD/MAD will be guided by the mTPI design, targeting a DLT rate of 30%. Doses of CDN STING agonist used in combination with pembrolizumab will be at least 2 dose levels behind the monotherapy CDN STING agonist dose, and will not exceed the MTD for monotherapy. If an MTD for the monotherapy arm is established, then the dose of CDN STING agonist in combination may continue escalation up to that dose. For example, if the MTD for monotherapy (Arm 1, Part A) is 90µg, then the starting dose for combination therapy (Arm 2, Part C), if no DLTs occurred in monotherapy, may be 10µg, with a maximum dose escalation to 90µg. If the MTD for monotherapy (Arm 1, Part A) is ≤ 30µg, then the starting dose for combination therapy will be 10µg. In monotherapy (Arm 1, Part A), if the 270µg dose level is completed, then the starting dose in combination therapy (Arm 2, Part C) will be 90µg.

[0254] A fixed dose of intravenous pembrolizumab 200mg will be administered every 3 weeks in Arm 2. A minimum of 3 subjects are required at each dose level during mTPI in both Arm 1 and Arm 2. The mTPI phase will have up to 3 to 6 subjects per cohort, and based on the occurrence of DLTs, up to 14 subjects may enroll per dose level. Therefore, during mTPI, up to 14 subjects may be enrolled per dose level, depending on the occurrence of a dose-limiting toxicity (DLT). Subjects may continue on their assigned treatment for up to 35 cycles (approximately 2 years) from the start of treatment. Treatment may continue until one of the following occurs: disease progression, unacceptable adverse event(s), intercurrent illness that prevents further administration of treatment, Investigator decision to withdraw the subject, subject withdraws consent, pregnancy of the subject, noncompliance with trials treatment or procedure requirements, or administrative reasons requiring cessation of treatment.

[0255] Subjects who progress by either clinical or radiographic evaluation on monotherapy with CDN STING agonist (Arm 1), may cross over into the combination arm of CDN STING agonist and pembrolizumab (Arm 2), provided that they meet crossover eligibility

criteria. Subjects who cross over from Arm 1 to Arm 2 are eligible for up to 35 cycles of treatment within Arm 2. Subjects who cross over will enter Arm 2 at the start of Arm 2.

[0256] Treatment allocation to Arm 1 will be accomplished by non-random assignment through an interactive voice response system/integrated web response system (IVRS/IWRS).

- 5 When both treatment arms are open for enrollment, IVRS/IWRS will alternate subject assignment between Arm 1 and 2, starting with Arm 1. Establishment of the MTD/MAD in the combination therapy of CDN STING agonist and pembrolizumab (Arm 2) requires that at least half of the subjects in Arm 2 have had no prior exposure to CDN STING agonist (i.e. non-crossover subjects). New subjects who are CDN STING agonist-naïve (non-crossover subjects)
- 10 will be given priority for enrollment into Arm 2.

- [0257] The final number of subjects enrolled in the dose escalation and confirmation parts of the study will depend on the empirical safety data (DLT observations, in particular, at which dose the mTPI design is triggered and at which dose the preliminary recommended Phase 2 dose is identified). For example, in a scenario where CDN STING agonist monotherapy starts
- 15 at 10µg and continues to the highest dose, the sample size across Parts A and B may be approximately 40 subjects. For combination therapy of CDN STING agonist with pembrolizumab, in a scenario where Arm 2 starts at 10µg of CDN STING agonist with 200mg of pembrolizumab, and continues to the highest dose, the sample size across Parts C and D may be approximately 40 subjects. In this scenario, the total sample size across Parts A-D will be
- 20 approximately 80 subjects. An administrative analysis may be conducted to enable future trial planning at the Sponsor's discretion, and data will be examined on a continuous basis to allow for dose escalation and confirmation decisions.

[0258] The trial will be conducted in conformance with Good Clinical Practices.

- [0259] Adverse Experiences (AEs) will be evaluated according to criteria outlined in the
- 25 National Cancer Institute (NCI) Common Terminology Criteria for Adverse Events (CTCAE) v4.

- [0260] It will be appreciated that various of the above-discussed and other features and functions, or alternatives thereof, may be desirably combined into many other different systems
- 30 or applications. Also that various presently unforeseen or unanticipated alternatives, modifications, variations or improvements therein may be subsequently made by those skilled in the art which are also intended to be encompassed by the following claims.

WHAT IS CLAIMED IS:

1. A method of treating a cell-proliferation disorder, said method comprising administering to a subject in need thereof a combination therapy that comprises

- 5 a) a PD-1 antagonist; and
b) a cyclic dinucleotide STING agonist;

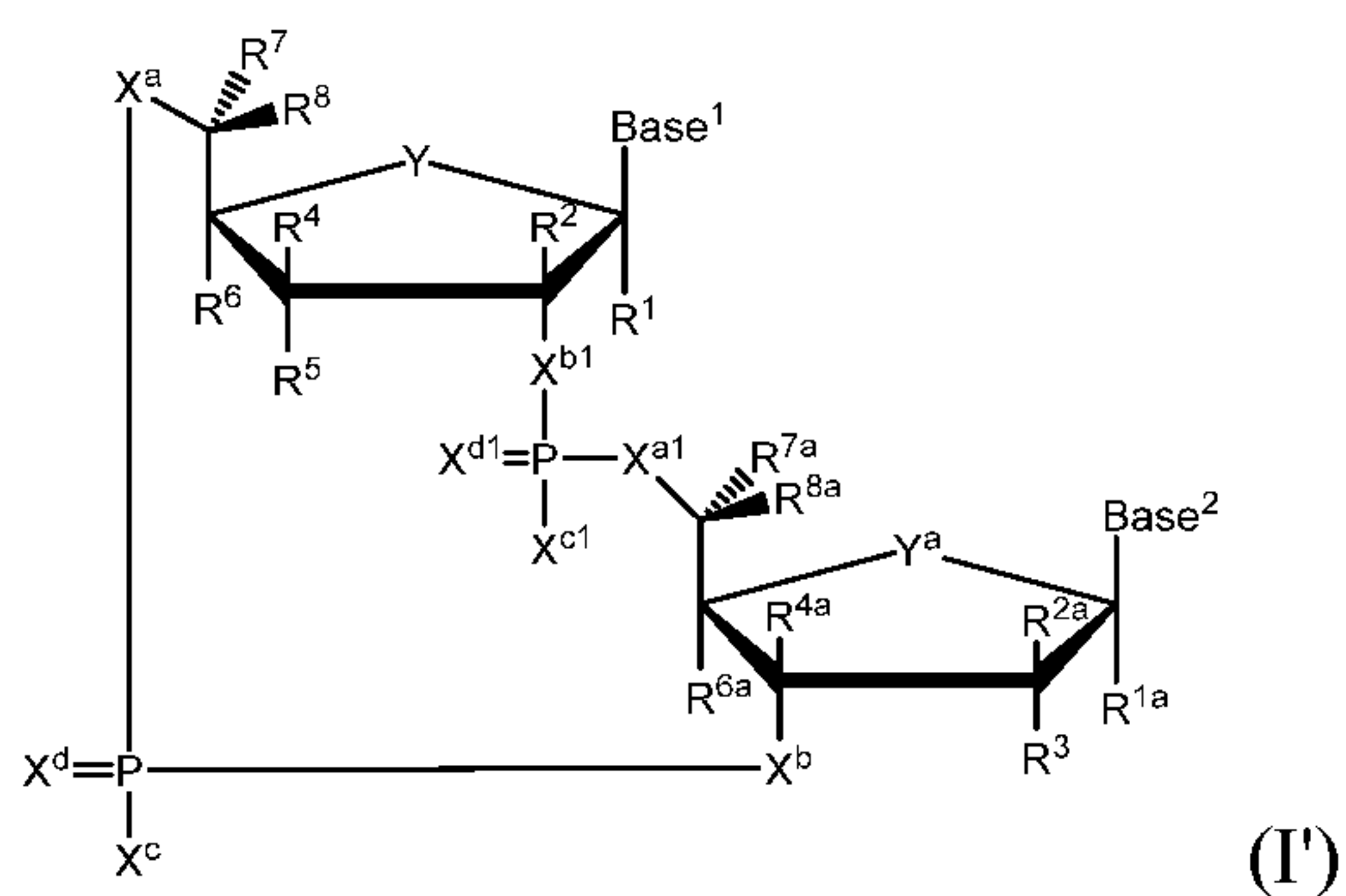
wherein

the PD-1 antagonist is administered once every 21 days; and

the cyclic dinucleotide STING agonist is administered once every 3 to 28 days;

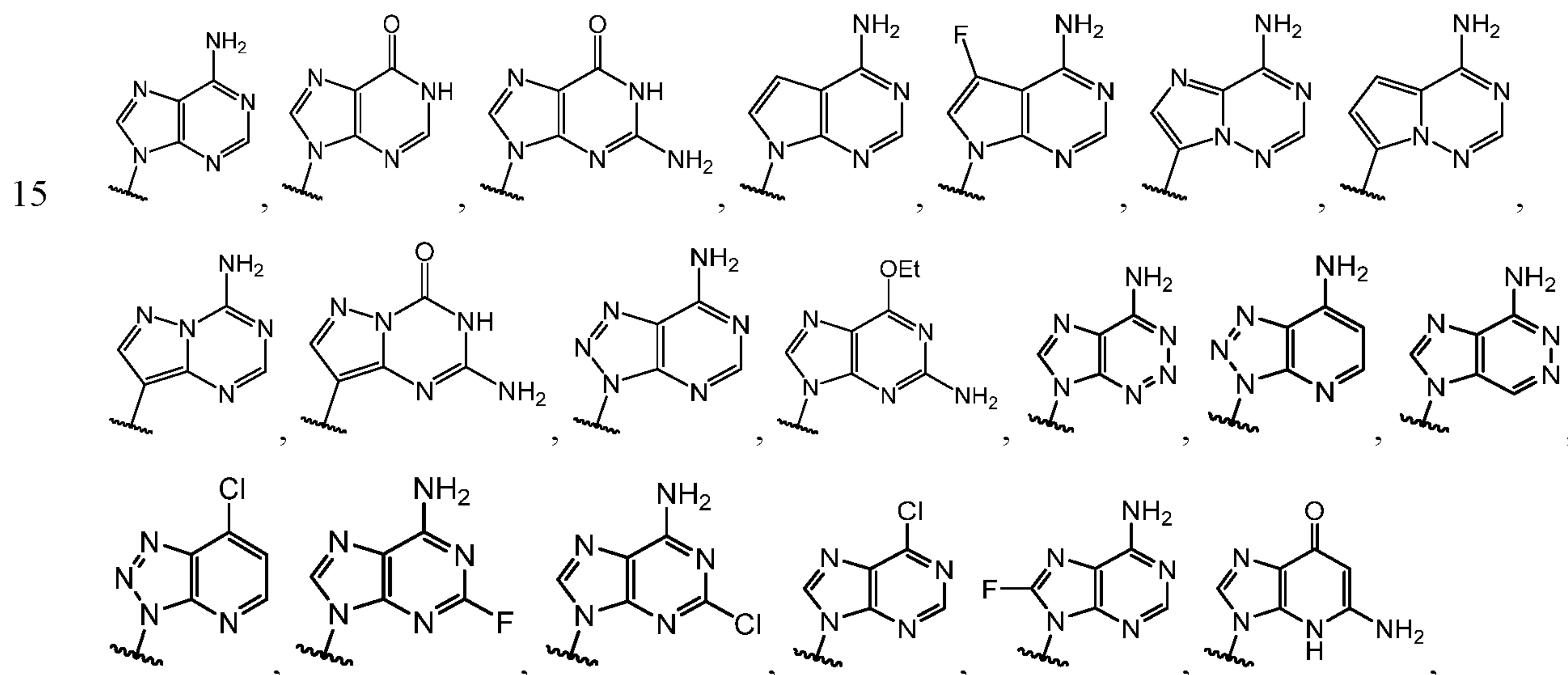
10 and

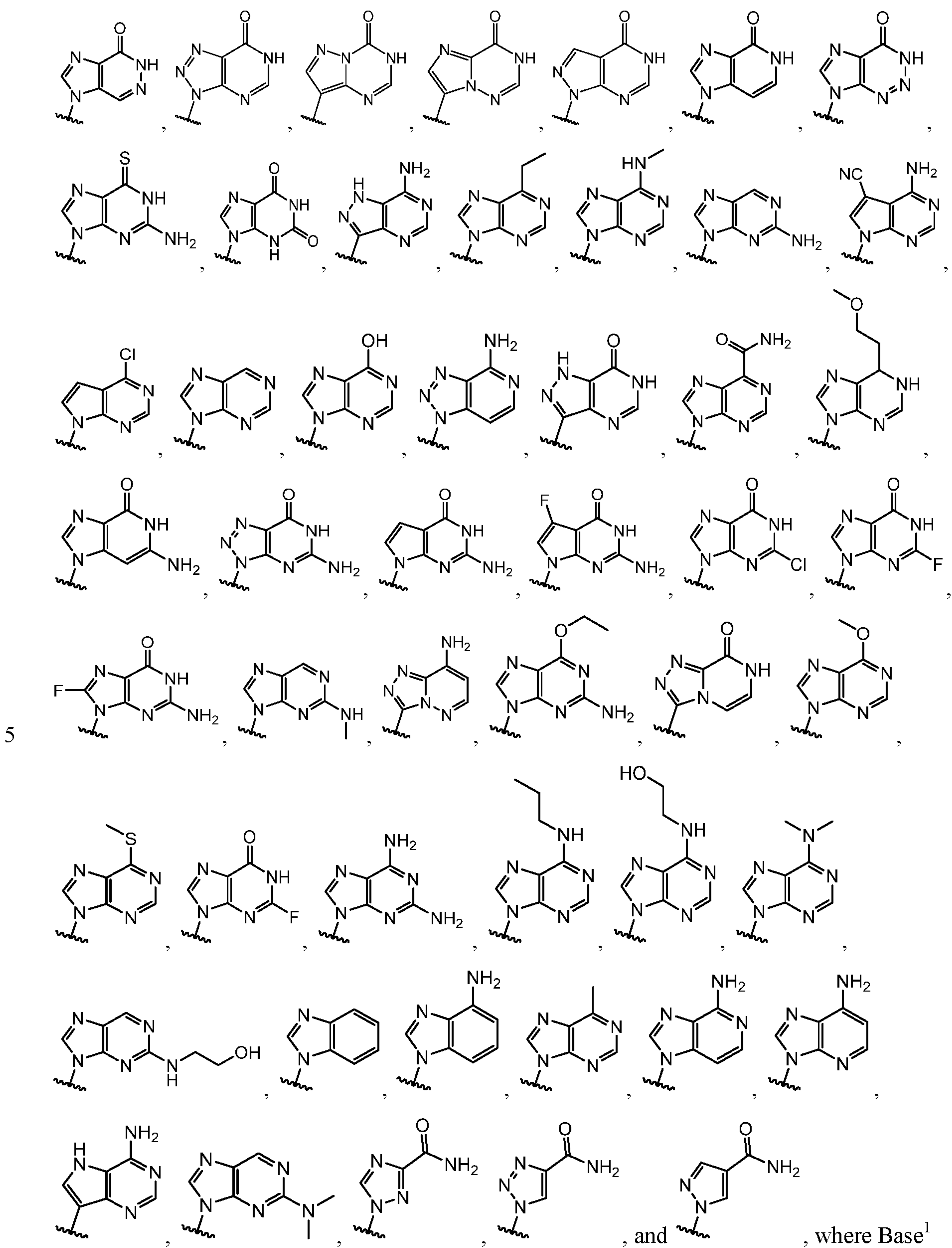
the cyclic dinucleotide STING agonist is selected from compounds of formula (I'):



or a pharmaceutically acceptable salt thereof, wherein

Base¹ and Base² are each independently selected from the group consisting of





independently selected from the group consisting of F, Cl, I, Br, OH, SH, NH₂, C₁₋₃ alkyl, C₃₋₆ cycloalkyl, O(C₁₋₃ alkyl), O(C₃₋₆ cycloalkyl), S(C₁₋₃ alkyl), S(C₃₋₆ cycloalkyl), NH(C₁₋₃ alkyl), NH(C₃₋₆ cycloalkyl), N(C₁₋₃ alkyl)₂, and N(C₃₋₆ cycloalkyl)₂;

Y and Y^a are each independently selected from the group consisting of -O- and
5 -S-;

X^a and X^{al} are each independently selected from the group consisting of O, and S;

X^b and X^{bl} are each independently selected from the group consisting of O, and S;

X^c and X^{cl} are each independently selected from the group consisting of OR⁹,
SR⁹, and NR⁹R⁹;

10 X^d and X^{dl} are each independently selected from the group consisting of O and S;

R¹ and R^{1a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C_{1-C6} alkyl, C_{2-C6} alkenyl, C_{2-C6} alkynyl, C_{1-C6} haloalkyl, C_{2-C6} haloalkenyl, C_{2-C6} haloalkynyl, -O-C_{1-C6} alkyl, -O-C_{2-C6} alkenyl, and -O-C_{2-C6} alkynyl, where said R¹ and R^{1a} C_{1-C6} alkyl, C_{2-C6} alkenyl, C_{2-C6} alkynyl, C_{1-C6} haloalkyl, C_{2-C6} haloalkenyl, C_{2-C6} haloalkynyl, -O-C_{1-C6} alkyl, -O-C_{2-C6} alkenyl, and -O-C_{2-C6} alkynyl are substituted by 0
15 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃;

R² and R^{2a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C_{1-C6} alkyl, C_{2-C6} alkenyl, C_{2-C6} alkynyl, C_{1-C6} haloalkyl, C_{2-C6} haloalkenyl, C_{2-C6} haloalkynyl, -O-C_{1-C6} alkyl, -O-C_{2-C6} alkenyl, and -O-C_{2-C6} alkynyl, where
20 said R² and R^{2a} C_{1-C6} alkyl, C_{2-C6} alkenyl, C_{2-C6} alkynyl, C_{1-C6} haloalkyl, C_{2-C6} haloalkenyl, C_{2-C6} haloalkynyl, -O-C_{1-C6} alkyl, -O-C_{2-C6} alkenyl, and -O-C_{2-C6} alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃;

R³ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C_{1-C6} alkyl, C_{2-C6} alkenyl, C_{2-C6} alkynyl, C_{1-C6} haloalkyl, C_{2-C6} haloalkenyl, C_{2-C6} haloalkynyl, -O-C_{1-C6} alkyl, -O-C_{2-C6} alkenyl, and -O-C_{2-C6} alkynyl, where said R³ C_{1-C6} alkyl, C_{2-C6} alkenyl, C_{2-C6} alkynyl, C_{1-C6} haloalkyl, C_{2-C6} haloalkenyl, C_{2-C6} haloalkynyl, -O-C_{1-C6} alkyl, -O-C_{2-C6} alkenyl, and -O-C_{2-C6} alkynyl are substituted by 0 to 3 substituents selected from the
25 group consisting of F, Cl, Br, I, OH, CN, and N₃;

R⁴ and R^{4a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C_{1-C6} alkyl, C_{2-C6} alkenyl, C_{2-C6} alkynyl, C_{1-C6} haloalkyl, C_{2-C6} haloalkenyl, C_{2-C6} haloalkynyl, -O-C_{1-C6} alkyl, -O-C_{2-C6} alkenyl, and -O-C_{2-C6} alkynyl, where
30 said R⁴ and R^{4a} C_{1-C6} alkyl, C_{2-C6} alkenyl, C_{2-C6} alkynyl, C_{1-C6} haloalkyl, C_{2-C6} haloalkenyl,

C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃;

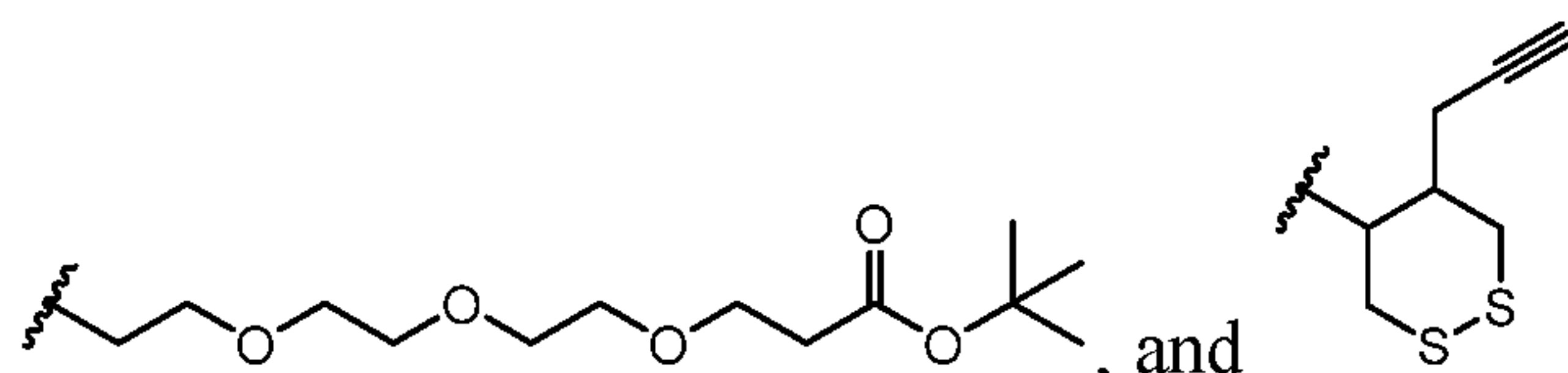
R⁵ is selected from the group consisting of H, F, Cl, Br, I, OH, CN, NH₂, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁵ C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, NR⁹R⁹, and N₃;

R⁶ and R^{6a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁶ and R^{6a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃;

R⁷ and R^{7a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁷ and R^{7a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃;

R⁸ and R^{8a} are each independently selected from the group consisting of H, F, Cl, Br, I, OH, CN, N₃, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl, where said R⁸ and R^{8a} C₁-C₆ alkyl, C₂-C₆ alkenyl, C₂-C₆ alkynyl, C₁-C₆ haloalkyl, C₂-C₆ haloalkenyl, C₂-C₆ haloalkynyl, -O-C₁-C₆ alkyl, -O-C₂-C₆ alkenyl, and -O-C₂-C₆ alkynyl are substituted by 0 to 3 substituents selected from the group consisting of F, Cl, Br, I, OH, CN, and N₃;

each R⁹ is independently selected from the group consisting of H, C₁-C₂₀ alkyl,



, where each R⁹ C₁-C₂₀ alkyl is optionally substituted by 0 to 3 substituents independently selected from the group consisting of OH, -O-C₁-C₂₀ alkyl, -S-C(O)C₁-C₆ alkyl, and -C(O)OC₁-C₆ alkyl;

optionally R^{1a} and R^3 are connected to form C_1-C_6 alkylene, C_2-C_6 alkenylene, C_2-C_6 alkynylene, $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, such that where R^{1a} and R^3 are connected to form $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, said O is bound at the R^3 position;

5 optionally R^{2a} and R^3 are connected to form C_1-C_6 alkylene, C_2-C_6 alkenylene, C_2-C_6 alkynylene, $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, such that where R^{2a} and R^3 are connected to form $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, said O is bound at the R^3 position;

10 optionally R^3 and R^{6a} are connected to form $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, such that where R^3 and R^{6a} are connected to form $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, said O is bound at the R^3 position;

optionally R^4 and R^5 are connected to form are connected to form C_1-C_6 alkylene, C_2-C_6 alkenylene, C_2-C_6 alkynylene, $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, such that where R^4 and R^5 are connected to form $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, said O is bound at the R^5 position;

15 optionally R^5 and R^6 are connected to form $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, such that where R^5 and R^6 are connected to form $-O-C_1-C_6$ alkylene, $-O-C_2-C_6$ alkenylene, or $-O-C_2-C_6$ alkynylene, said O is bound at the R^5 position;

20 optionally R^7 and R^8 are connected to form C_1-C_6 alkylene, C_2-C_6 alkenylene, or C_2-C_6 alkynylene; and

optionally R^{7a} and R^{8a} are connected to form C_1-C_6 alkylene, C_2-C_6 alkenylene, or C_2-C_6 alkynylene.

25 2. The method according to claim 1, wherein the cell-proliferation disorder is cancer.

3. The method according to claim 2, wherein the cancer occurs as one or more solid tumors or lymphomas.

30 4. The method according to claim 2, wherein the cancer is selected from the group consisting of advanced or metastatic solid tumors and lymphomas.

5. The method according to claim 2, wherein the cancer is selected from the group consisting of malignant melanoma, head and neck squamous cell carcinoma, breast adenocarcinoma, and lymphoma.

5 6. The method according to any one of claims 3 to 5, wherein the lymphoma is selected from the group consisting of diffuse large B-cell lymphoma, follicular lymphoma, mantle cell lymphoma, small lymphocytic lymphoma, mediastinal large B-cell lymphoma, splenic marginal zone B-cell lymphoma, extranodal marginal zone B-cell lymphoma of mucosa-associated lymphoid tissue (malt), nodal marginal zone B-cell lymphoma, lymphoplasmacytic
10 lymphoma, primary effusion lymphoma, Burkitt lymphoma, anaplastic large cell lymphoma (primary cutaneous type), anaplastic large cell lymphoma (systemic type), peripheral T-cell lymphoma, angioimmunoblastic T-cell lymphoma, adult T-cell lymphoma, nasal type extranodal NK/T-cell lymphoma, enteropathy-associated T-cell lymphoma, gamma/delta hepatosplenic T-cell lymphoma, subcutaneous panniculitis-like T-cell lymphoma, mycosis fungoides, and
15 Hodgkin lymphoma.

7. The method according to claim 2, wherein the cell-proliferation disorder is a cancer that has metastasized.

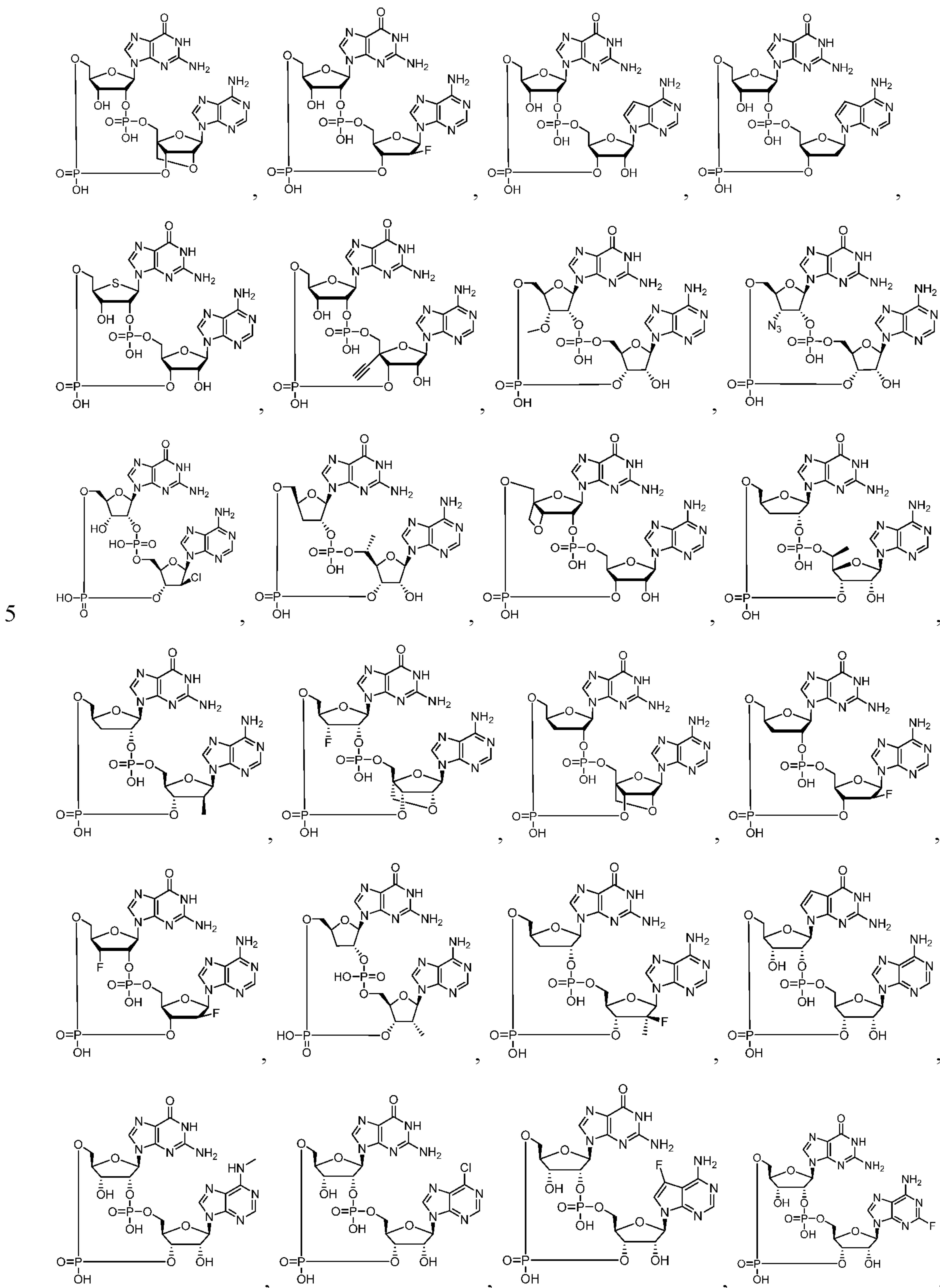
20 8. The method according to any one of claims 1 to 7, wherein the PD-1 antagonist is an anti-PD-1 monoclonal antibody.

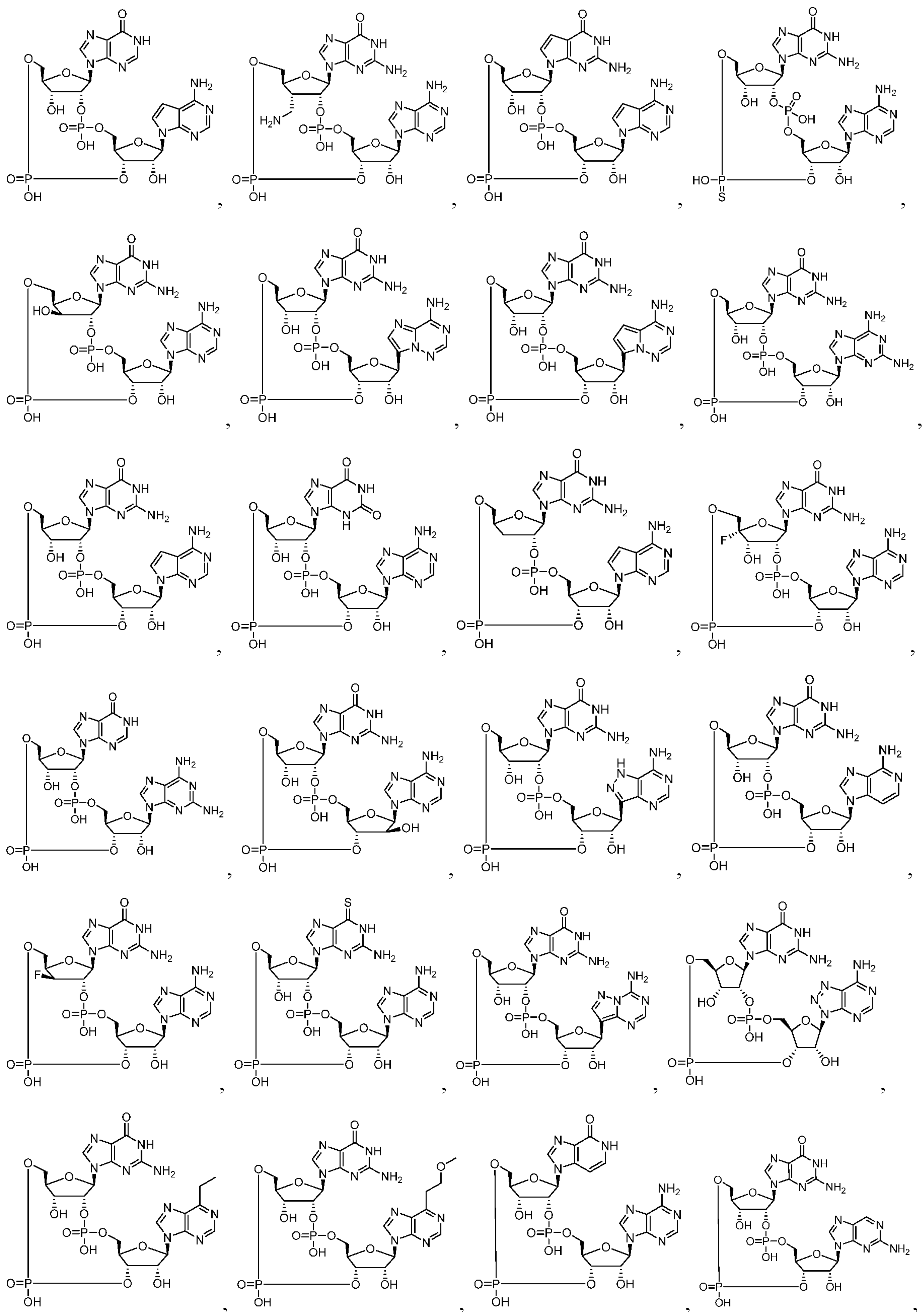
9. The method according to claim 8, wherein the PD-1 antagonist is selected from the group consisting of nivolumab, pembrolizumab, pidilizumab, and AMP-224.
25

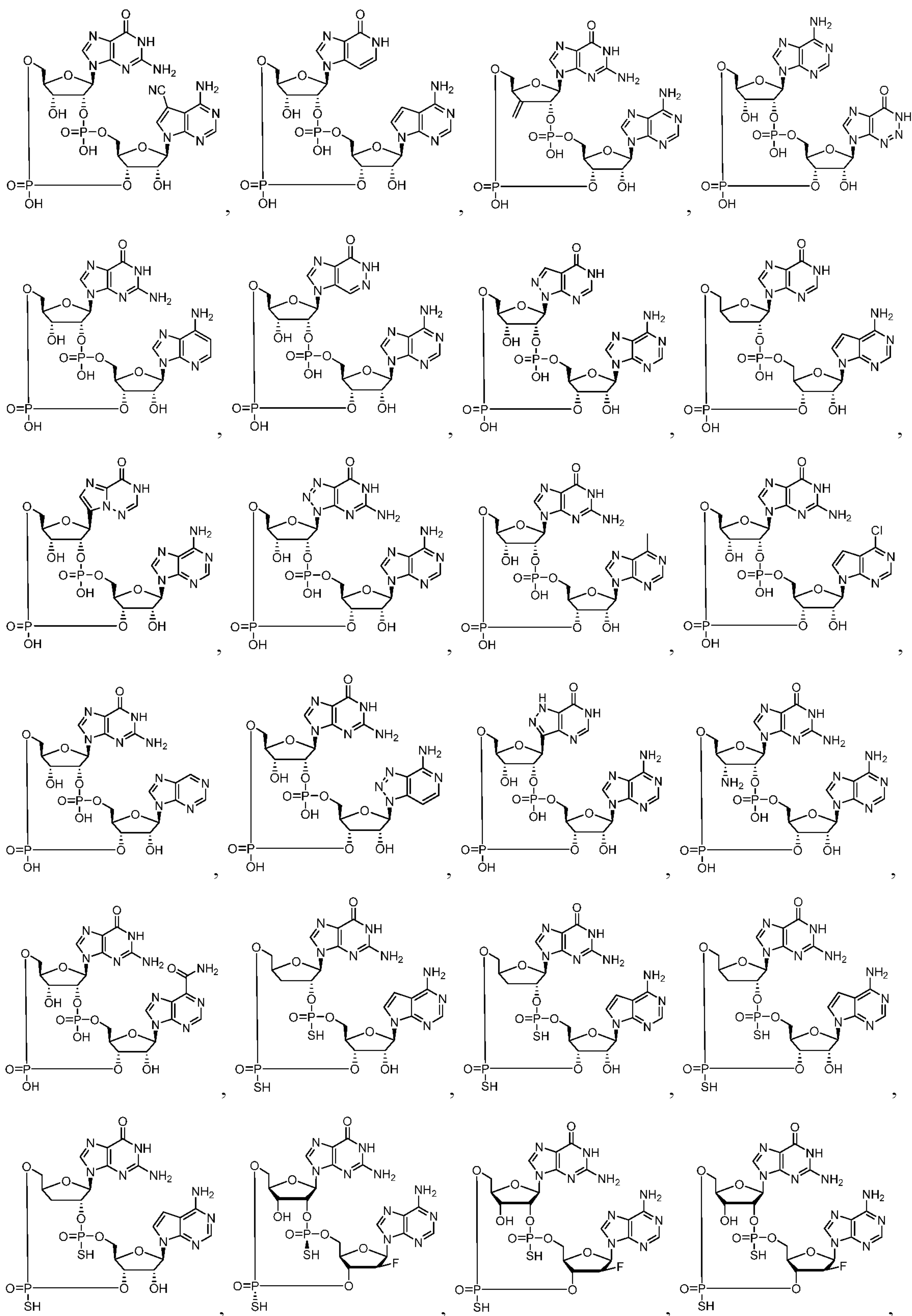
10. The method according to claim 9, wherein the PD-1 antagonist is nivolumab.

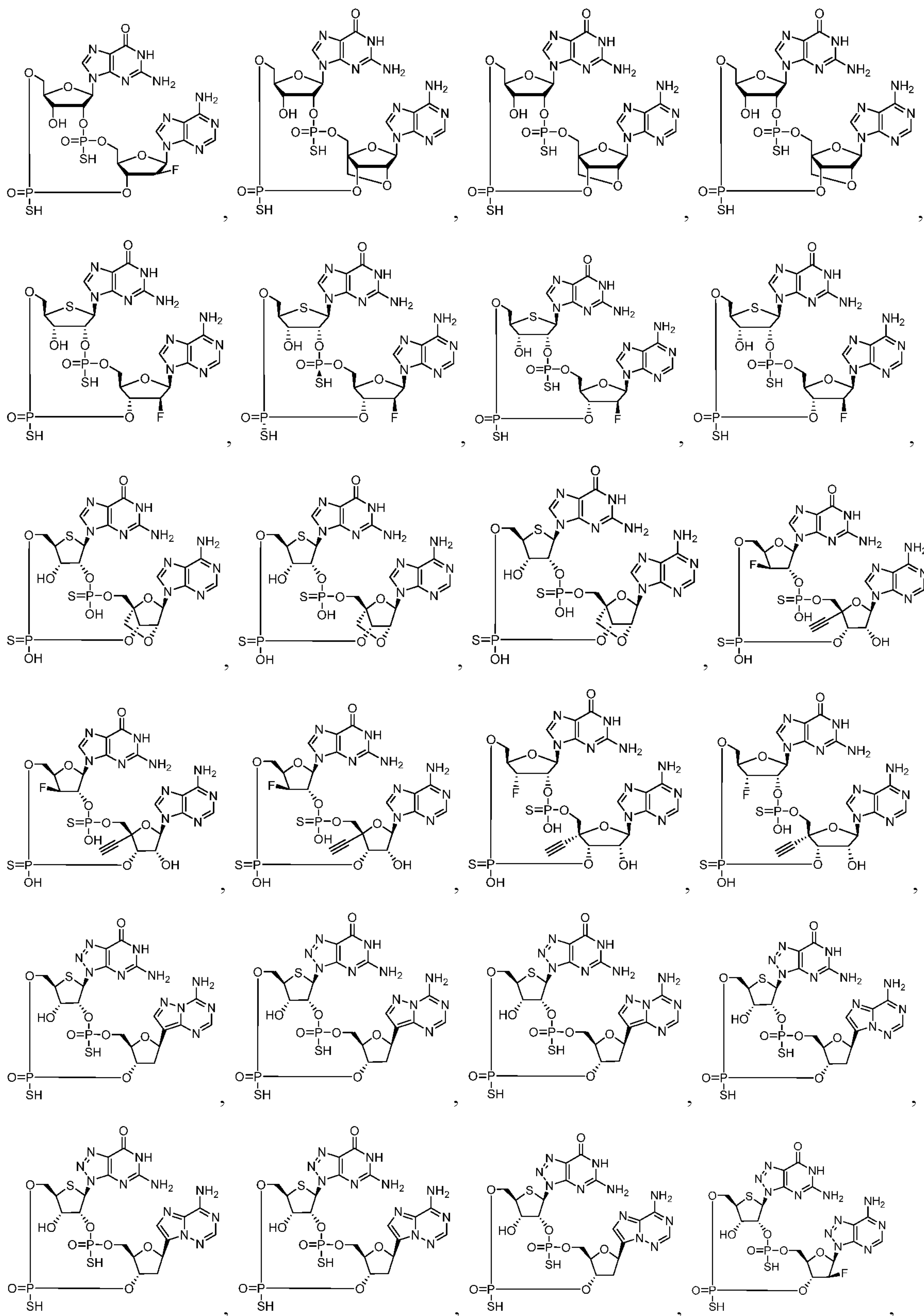
11. The method according to claim 9, wherein the PD-1 antagonist is
30 pembrolizumab.

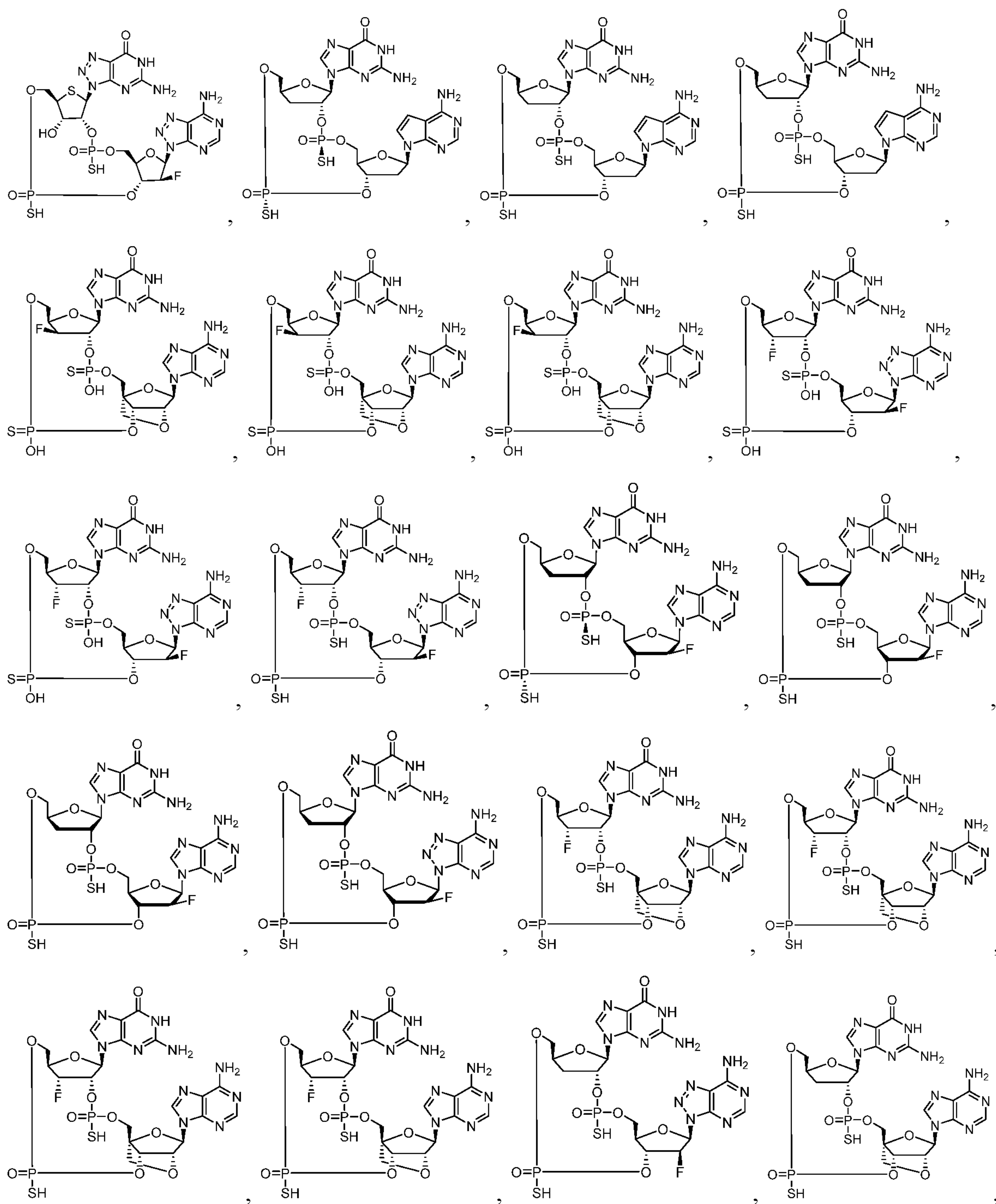
12. The method according to any one of claims 1 to 11, wherein the cyclic dinucleotide STING agonist is selected from the group consisting of:

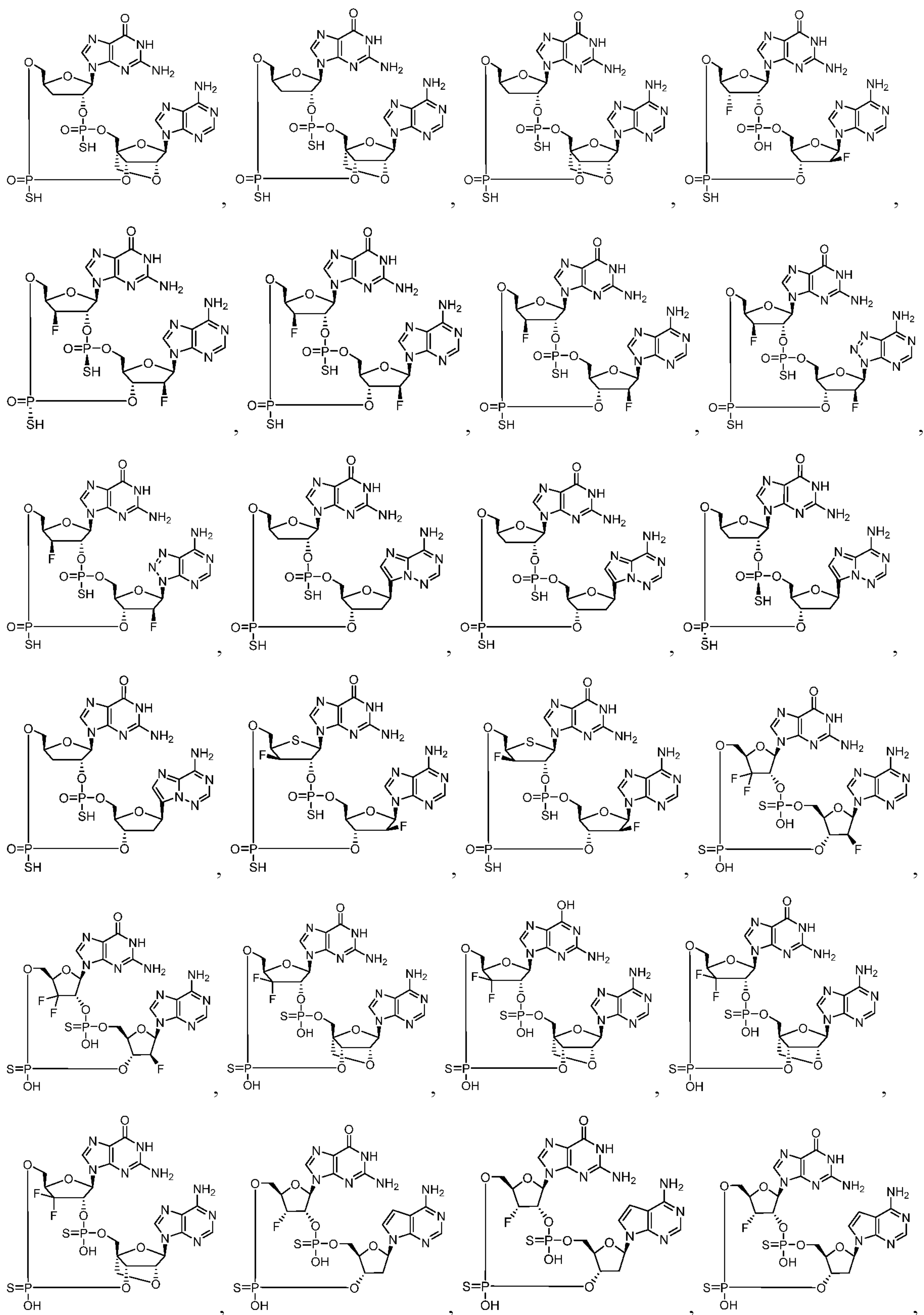




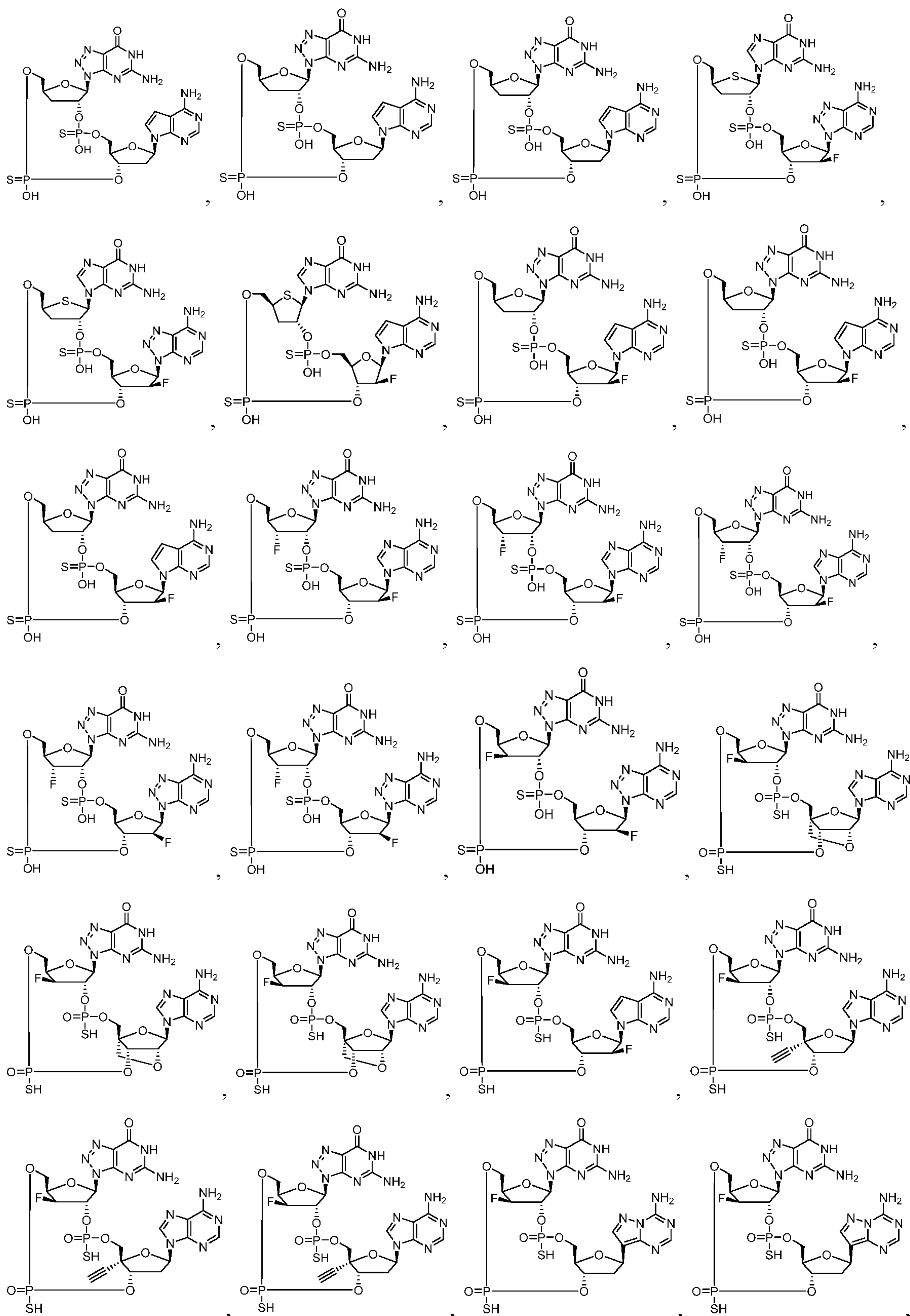


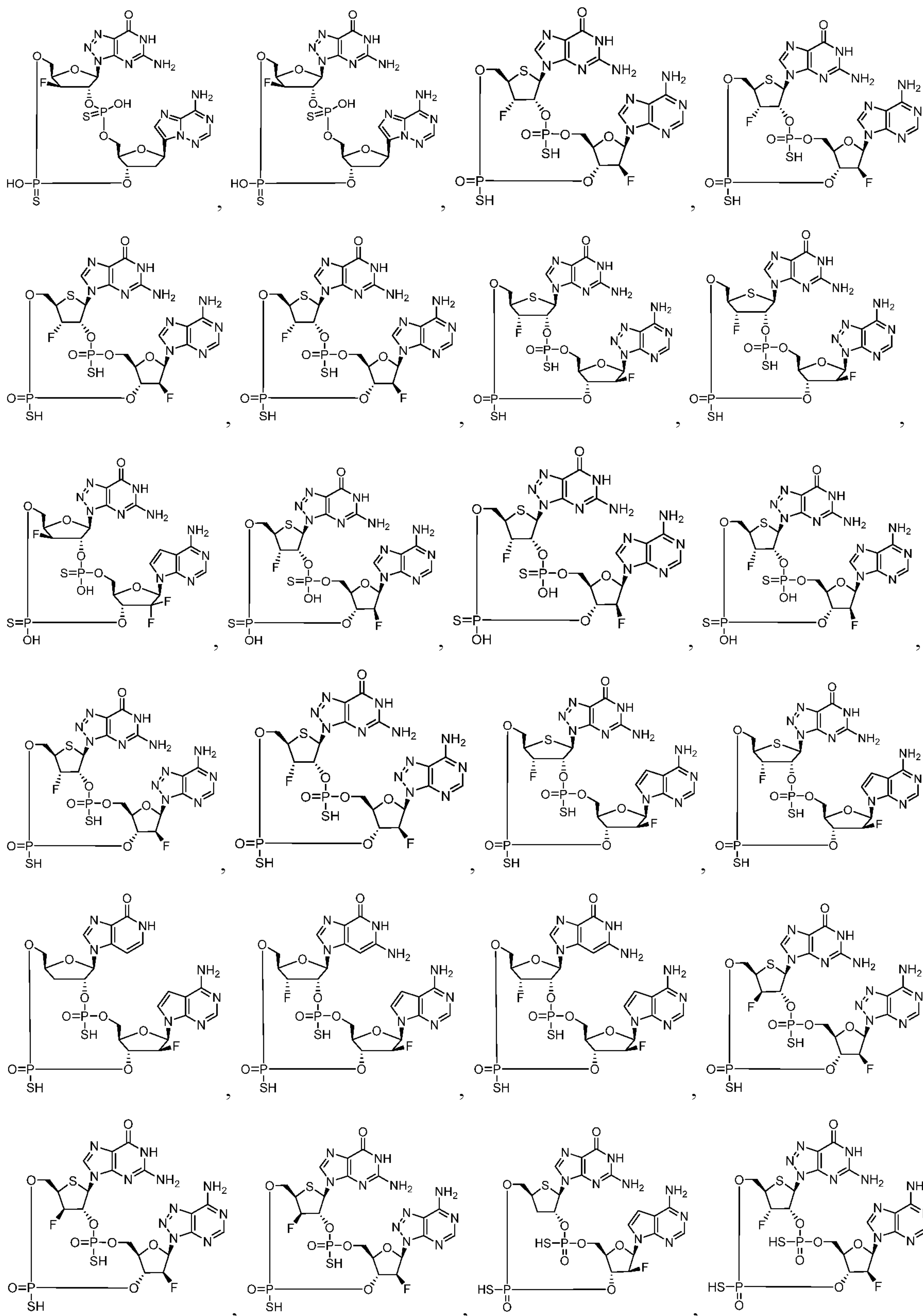


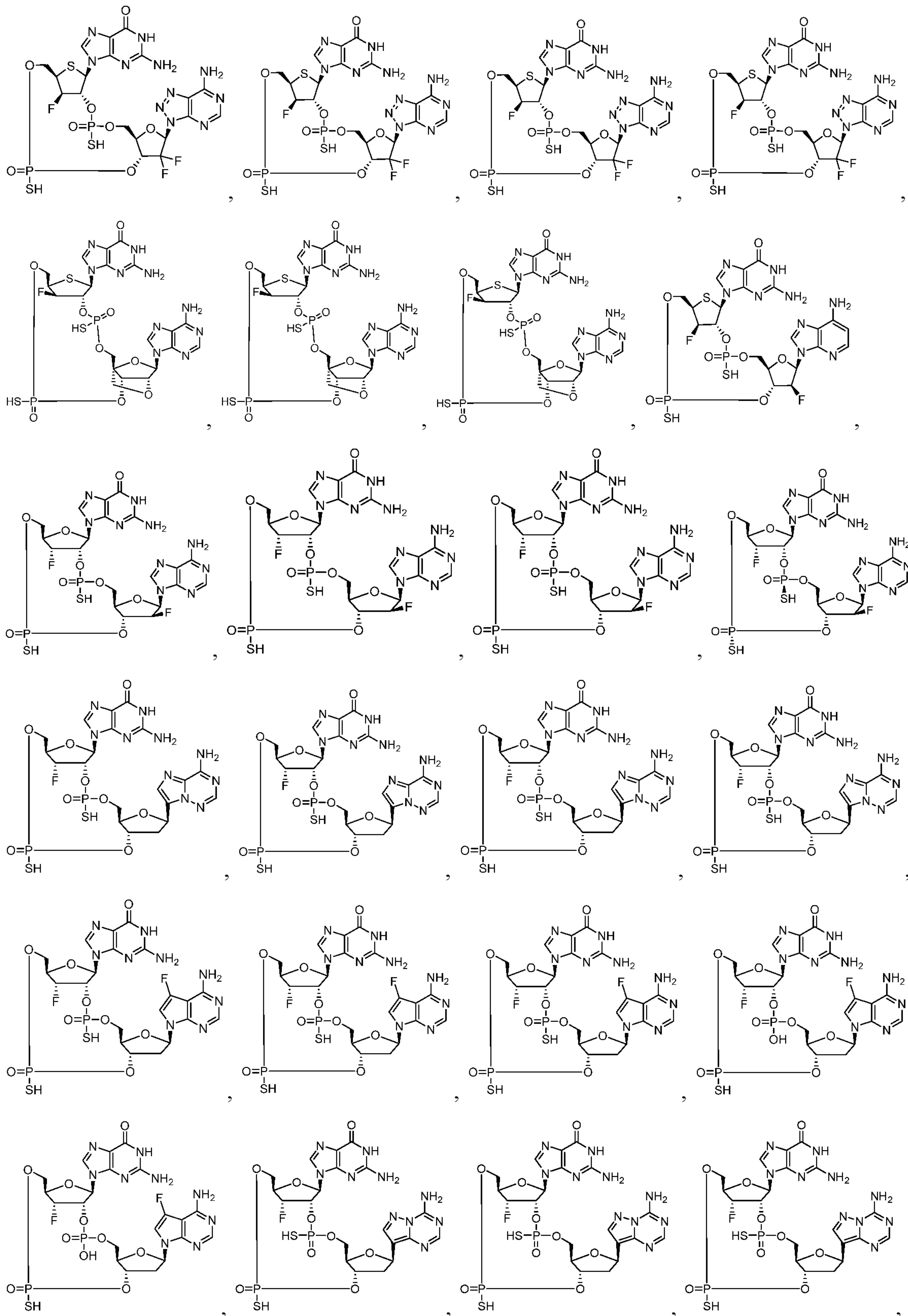


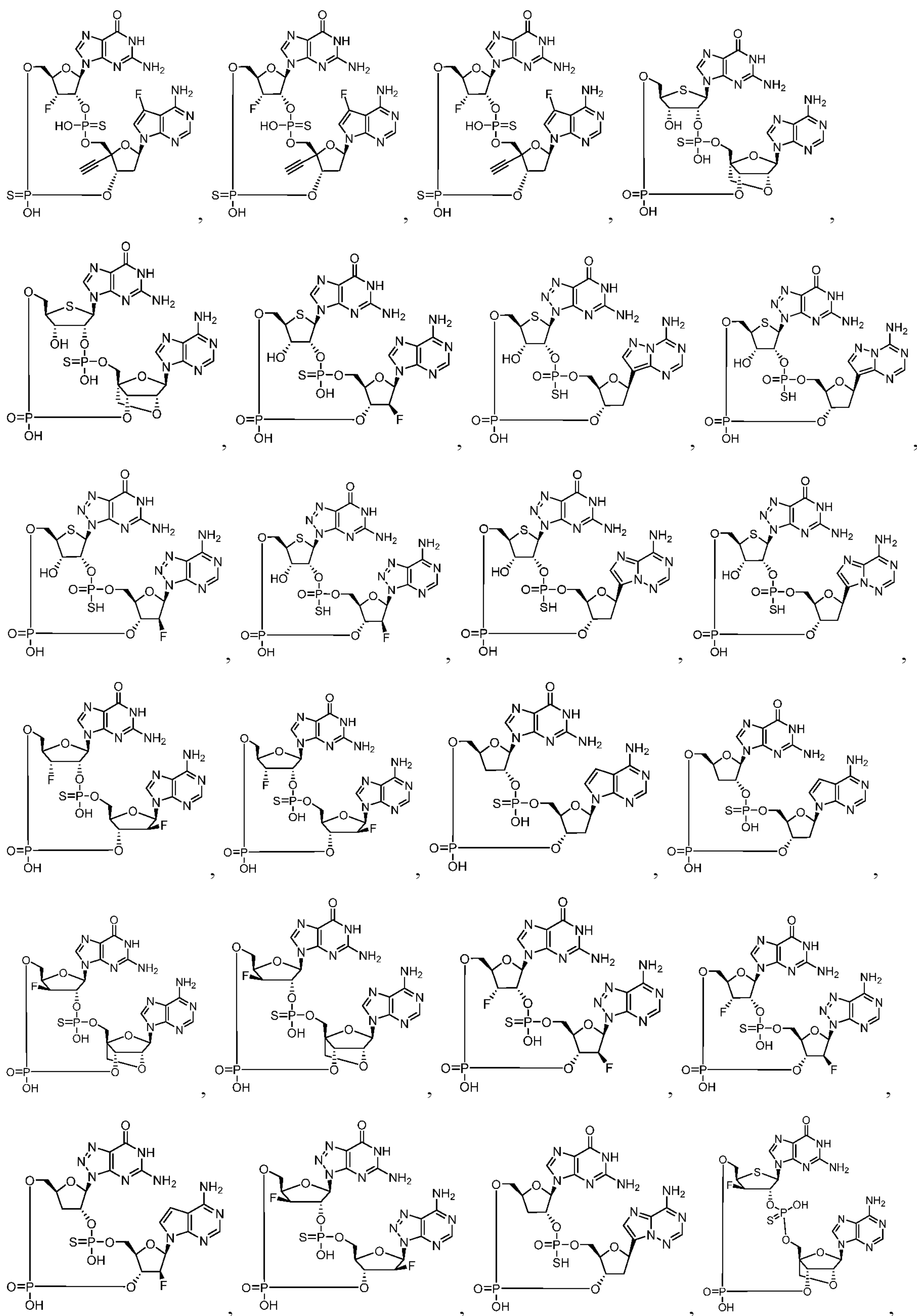




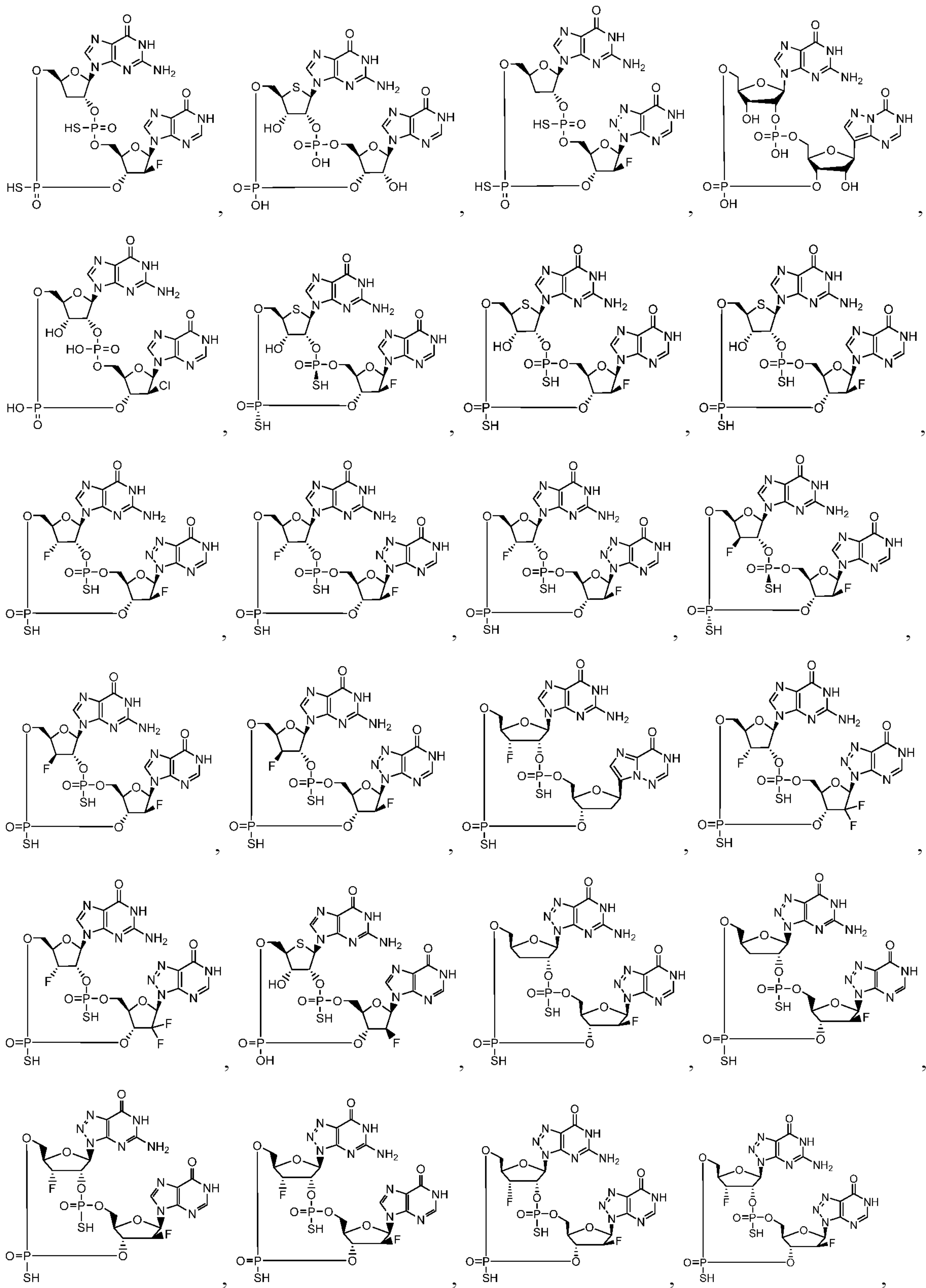


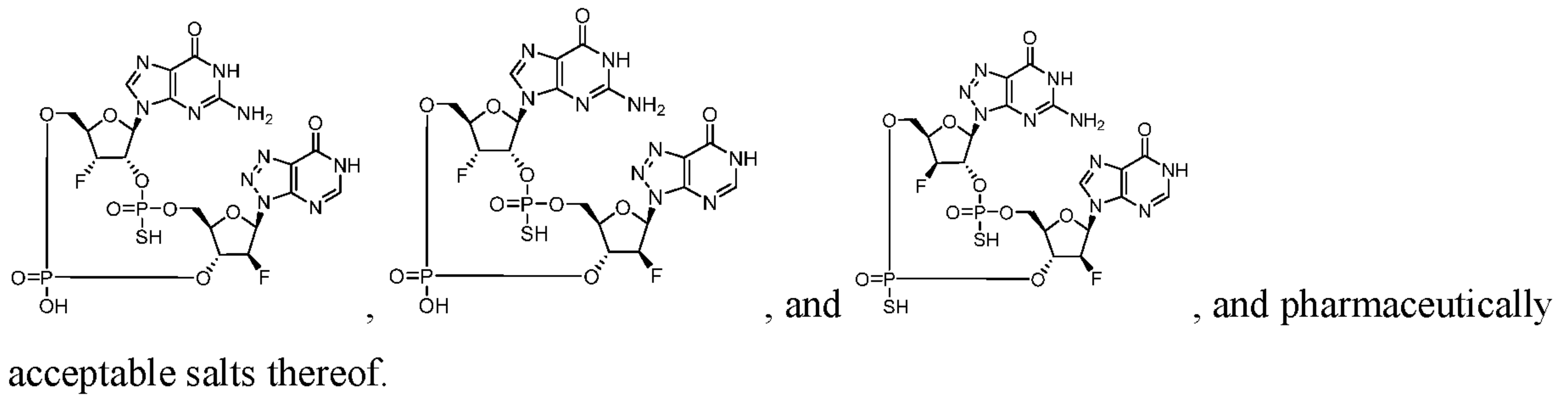












13. The method of any one of claims 1 to 12, wherein the PD-1 antagonist is administered by intravenous infusion, and the cyclic dinucleotide STING agonist is administered orally, by intravenous infusion, by intertumoral injection, or by subcutaneous injection.

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SEQ ID NO 1: Pembrolizumab, Variable Light chain, amino acid

Glu Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser
Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Lys Gly
Val Ser Thr Ser Gly Tyr Ser Tyr Leu His Trp Tyr Gln Gln
Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr Leu Ala Ser
Tyr Leu Glu Ser Gly Val Pro Ala Arg Phe Ser Gly Ser Gly
Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro
Glu Asp Phe Ala Val Tyr Tyr Cys Gln His Ser Arg Asp Leu
Pro Leu Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys

SEQ ID NO 2: Pembrolizumab, Variable Heavy chain, amino acid

Gln Val Gln Leu Val Gln Ser Gly Val Glu Val Lys Lys Pro
Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr
Phe Thr Asn Tyr Tyr Met Tyr Trp Val Arg Gln Ala Pro Gly
Gln Gly Leu Glu Trp Met Gly Gly Ile Asn Pro Ser Asn Gly
Gly Thr Asn Phe Asn Glu Lys Phe Lys Asn Arg Val Thr Leu
Thr Thr Asp Ser Ser Thr Thr Thr Ala Tyr Met Glu Leu Lys
Ser Leu Gln Phe Asp Asp Thr Ala Val Tyr Tyr Cys Ala Arg
Arg Asp Tyr Arg Phe Asp Met Gly Phe Asp Tyr Trp Gly Gln
Gly Thr Thr Val Thr Val Ser Ser

FIG.1

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SEQ ID NO 3: Pembrolizumab, Light chain, amino acid

Glu Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser
Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Lys Gly
Val Ser Thr Ser Gly Tyr Ser Tyr Leu His Trp Tyr Gln Gln
Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr Leu Ala Ser
Tyr Leu Glu Ser Gly Val Pro Ala Arg Phe Ser Gly Ser Gly
Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro
Glu Asp Phe Ala Val Tyr Tyr Cys Gln His Ser Arg Asp Leu
Pro Leu Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys Arg
Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu
Asn Asn Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val
Asp Asn Ala Leu Gln Ser Gly Asn Ser Gln Glu Ser Val Thr
Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu Ser Ser Thr
Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Val Tyr
Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr
Lys Ser Phe Asn Arg Gly Glu Cys

FIG.2

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SEQ ID NO 4: Pembrolizumab, Heavy chain, amino acid

Gln	Val	Gln	Leu	Val	Gln	Ser	Gly	Val	Glu	Val	Lys	Lys	Pro
Gly	Ala	Ser	Val	Lys	Val	Ser	Cys	Lys	Ala	Ser	Gly	Tyr	Thr
Phe	Thr	Asn	Tyr	Tyr	Met	Tyr	Trp	Val	Arg	Gln	Ala	Pro	Gly
Gln	Gly	Leu	Glu	Trp	Met	Gly	Gly	Ile	Asn	Pro	Ser	Asn	Gly
Gly	Thr	Asn	Phe	Asn	Glu	Lys	Phe	Lys	Asn	Arg	Val	Thr	Leu
Thr	Thr	Asp	Ser	Ser	Thr	Thr	Thr	Ala	Tyr	Met	Glu	Leu	Lys
Ser	Leu	Gln	Phe	Asp	Asp	Thr	Ala	Val	Tyr	Tyr	Cys	Ala	Arg
Arg	Asp	Tyr	Arg	Phe	Asp	Met	Gly	Phe	Asp	Tyr	Trp	Gly	Gln
Gly	Thr	Thr	Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro
Ser	Val	Phe	Pro	Leu	Ala	Pro	Cys	Ser	Arg	Ser	Thr	Ser	Glu
Ser	Thr	Ala	Ala	Leu	Gly	Cys	Leu	Val	Lys	Asp	Tyr	Phe	Pro
Glu	Pro	Val	Thr	Val	Ser	Trp	Asn	Ser	Gly	Ala	Leu	Thr	Ser
Gly	Val	His	Thr	Phe	Pro	Ala	Val	Leu	Gln	Ser	Ser	Gly	Leu
Tyr	Ser	Leu	Ser	Ser	Val	Val	Thr	Val	Pro	Ser	Ser	Ser	Leu
Gly	Thr	Lys	Thr	Tyr	Thr	Cys	Asn	Val	Asp	His	Lys	Pro	Ser
Asn	Thr	Lys	Val	Asp	Lys	Arg	Val	Glu	Ser	Lys	Tyr	Gly	Pro
Pro	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Phe	Leu	Gly	Gly	Pro
Ser	Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met
Ile	Ser	Arg	Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val
Ser	Gln	Glu	Asp	Pro	Glu	Val	Gln	Phe	Asn	Trp	Tyr	Val	Asp
Gly	Val	Glu	Val	His	Asn	Ala	Lys	Thr	Lys	Pro	Arg	Glu	Glu
Gln	Phe	Asn	Ser	Thr	Tyr	Arg	Val	Val	Ser	Val	Leu	Thr	Val
Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr	Lys	Cys	Lys
Val	Ser	Asn	Lys	Gly	Leu	Pro	Ser	Ser	Ile	Glu	Lys	Thr	Ile
Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr
Leu	Pro	Pro	Ser	Gln	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser
Leu	Thr	Cys	Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala
Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys
Thr	Thr	Pro	Pro	Val	Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe	Leu
Tyr	Ser	Arg	Leu	Thr	Val	Asp	Lys	Ser	Arg	Trp	Gln	Glu	Gly
Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	Leu	His	Asn
His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Leu	Gly	Lys	

FIG.3

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SEQ ID NO 5: Pembrolizumab, CDRL1

Arg Ala Ser Lys Gly Val Ser Thr Ser Gly Tyr Ser Tyr Leu His

SEQ ID NO 6: Pembrolizumab, CDRL2

Leu Ala Ser Tyr Leu Glu Ser

SEQ ID NO 7: Pembrolizumab, CDRL3

Gln His Ser Arg Asp Leu Pro Leu Thr

SEQ ID NO 8: Pembrolizumab, CDRH1

Asn Tyr Tyr Met Tyr

SEQ ID NO 9: Pembrolizumab, CDRH2

Gly Ile Asn Pro Ser Asn Gly Gly Thr Asn Phe Asn Glu Lys Phe Lys

SEQ ID NO 10: Pembrolizumab, CDRH3

Arg Asp Tyr Arg Phe Asp Met Gly Phe Asp Tyr

FIG.4

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SEQ ID NO 11: Nivolumab, Variable Light chain, amino acid

Glu Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser
Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser
Val Ser Ser Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln
Ala Pro Arg Leu Leu Ile Tyr Asp Ala Ser Asn Arg Ala Thr
Gly Ile Pro Ala Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp
Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro Glu Asp Phe Ala
Val Tyr Tyr Cys Gln Gln Ser Ser Asn Trp Pro Arg Thr Phe
Gly Gln Gly Thr Lys Val Glu Ile Lys

SEQ ID NO 12: Nivolumab, Variable Heavy chain, amino acid

Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln Pro
Gly Arg Ser Leu Arg Leu Asp Cys Lys Ala Ser Gly Ile Thr
Phe Ser Asn Ser Gly Met His Trp Val Arg Gln Ala Pro Gly
Lys Gly Leu Glu Trp Val Ala Val Ile Trp Tyr Asp Gly Ser
Lys Arg Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile
Ser Arg Asp Asn Ser Lys Asn Thr Leu Phe Leu Gln Met Asn
Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Thr
Asn Asp Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser
Ser

FIG.5

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SEQ ID NO 13: Nivolumab, Light chain, amino acid

Glu Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser
Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser
Val Ser Ser Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln
Ala Pro Arg Leu Leu Ile Tyr Asp Ala Ser Asn Arg Ala Thr
Gly Ile Pro Ala Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp
Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro Glu Asp Phe Ala
Val Tyr Tyr Cys Gln Gln Ser Ser Asn Trp Pro Arg Thr Phe
Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala Ala
Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys
Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr
Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu
Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser
Lys Asp Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser
Lys Ala Asp Tyr Glu Lys His Lys Val Tyr Ala Cys Glu Val
Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys Ser Phe Asn
Arg Gly Glu Cys

FIG.6

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SEQ ID NO 14: Nivolumab, Heavy chain, amino acid

Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln Pro
 Gly Arg Ser Leu Arg Leu Asp Cys Lys Ala Ser Gly Ile Thr
 Phe Ser Asn Ser Gly Met His Trp Val Arg Gln Ala Pro Gly
 Lys Gly Leu Glu Trp Val Ala Val Ile Trp Tyr Asp Gly Ser
 Lys Arg Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile
 Ser Arg Asp Asn Ser Lys Asn Thr Leu Phe Leu Gln Met Asn
 Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Thr
 Asn Asp Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser
 Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro
 Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu Gly Cys
 Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
 Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala
 Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val
 Thr Val Pro Ser Ser Ser Leu Gly Thr Lys Thr Tyr Thr Cys
 Asn Val Asp His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg
 Val Glu Ser Lys Tyr Gly Pro Pro Cys Pro Pro Cys Pro Ala
 Pro Glu Phe Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro
 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val
 Thr Cys Val Val Val Asp Val Ser Gln Glu Asp Pro Glu Val
 Gln Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala
 Lys Thr Lys Pro Arg Glu Glu Gln Phe Asn Ser Thr Tyr Arg
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn
 Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Gly Leu Pro
 Ser Ser Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro
 Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Gln Glu Glu
 Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly
 Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly
 Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Arg Leu Thr Val Asp
 Lys Ser Arg Trp Gln Glu Gly Asn Val Phe Ser Cys Ser Val
 Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu
 Ser Leu Ser Leu Gly Lys

FIG. 7

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SEQ ID NO 15: Nivolumab, CDRL1

Arg Ala Ser Gln Ser Val Ser Ser Tyr Leu Ala

SEQ ID NO 16: Nivolumab, CDRL2

Asp Ala Ser Asn Arg Ala Thr

SEQ ID NO 17: Nivolumab, CDRL3

Gln Gln Ser Ser Asn Trp Pro Arg Thr

SEQ ID NO 18: Nivolumab, CDRH1

Asn Ser Gly Met His

SEQ ID NO 19: Nivolumab, CDRH2

Val Ile Trp Tyr Asp Gly Ser Lys Arg Tyr Tyr Ala
Asp Ser Val Lys Gly

SEQ ID NO 20: Nivolumab, CDRH3

Asn Asp Asp Tyr

FIG.8

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SEQ ID NO 21: Human PD-L1

Met	Arg	Ile	Phe	Ala	Val	Phe	Ile	Phe	Met	Thr	Tyr	Trp	His
Leu	Leu	Asn	Ala	Phe	Thr	Val	Thr	Val	Pro	Lys	Asp	Leu	Tyr
Val	Val	Glu	Tyr	Gly	Ser	Asn	Met	Thr	Ile	Glu	Cys	Lys	Phe
Pro	Val	Glu	Lys	Gln	Leu	Asp	Leu	Ala	Ala	Leu	Ile	Val	Tyr
Trp	Glu	Met	Glu	Asp	Lys	Asn	Ile	Ile	Gln	Phe	Val	His	Gly
Glu	Glu	Asp	Leu	Lys	Val	Gln	His	Ser	Ser	Tyr	Arg	Gln	Arg
Ala	Arg	Leu	Leu	Lys	Asp	Gln	Leu	Ser	Leu	Gly	Asn	Ala	Ala
Leu	Gln	Ile	Thr	Asp	Val	Lys	Leu	Gln	Asp	Ala	Gly	Val	Tyr
Arg	Cys	Met	Ile	Ser	Tyr	Gly	Gly	Ala	Asp	Tyr	Lys	Arg	Ile
Thr	Val	Lys	Val	Asn	Ala	Pro	Tyr	Asn	Lys	Ile	Asn	Gln	Arg
Ile	Leu	Val	Val	Asp	Pro	Val	Thr	Ser	Glu	His	Glu	Leu	Thr
Cys	Gln	Ala	Glu	Gly	Tyr	Pro	Lys	Ala	Glu	Val	Ile	Trp	Thr
Ser	Ser	Asp	His	Gln	Val	Leu	Ser	Gly	Lys	Thr	Thr	Thr	Thr
Asn	Ser	Lys	Arg	Glu	Glu	Lys	Leu	Phe	Asn	Val	Thr	Ser	Thr
Leu	Arg	Ile	Asn	Thr	Thr	Thr	Asn	Glu	Ile	Phe	Tyr	Cys	Thr
Phe	Arg	Arg	Leu	Asp	Pro	Glu	Glu	Asn	His	Thr	Ala	Glu	Leu
Val	Ile	Pro	Glu	Leu	Pro	Leu	Ala	His	Pro	Pro	Asn	Glu	Arg
Thr	His	Leu	Val	Ile	Leu	Gly	Ala	Ile	Leu	Leu	Cys	Leu	Gly
Val	Ala	Leu	Thr	Phe	Ile	Phe	Arg	Leu	Arg	Lys	Gly	Arg	Met
Met	Asp	Val	Lys	Lys	Cys	Gly	Ile	Gln	Asp	Thr	Asn	Ser	Lys
Lys	Gln	Ser	Asp	Thr	His	Leu	Glu	Glu	Thr				

FIG.9

SEQ ID NO 1: Pembrolizumab, Variable Light chain, amino acid

Glu	Ile	Val	Leu	Thr	Gln	Ser	Pro	Ala	Thr	Leu	Ser	Leu	Ser
Pro	Gly	Glu	Arg	Ala	Thr	Leu	Ser	Cys	Arg	Ala	Ser	Lys	Gly
Val	Ser	Thr	Ser	Gly	Tyr	Ser	Tyr	Leu	His	Trp	Tyr	Gln	Gln
Lys	Pro	Gly	Gln	Ala	Pro	Arg	Leu	Leu	Ile	Tyr	Leu	Ala	Ser
Tyr	Leu	Glu	Ser	Gly	Val	Pro	Ala	Arg	Phe	Ser	Gly	Ser	Gly
Ser	Gly	Thr	Asp	Phe	Thr	Leu	Thr	Ile	Ser	Ser	Leu	Glu	Pro
Glu	Asp	Phe	Ala	Val	Tyr	Tyr	Cys	Gln	His	Ser	Arg	Asp	Leu
Pro	Leu	Thr	Phe	Gly	Gly	Gly	Thr	Lys	Val	Glu	Ile	Lys	

SEQ ID NO 2: Pembrolizumab, Variable Heavy chain, amino acid

Gln	Val	Gln	Leu	Val	Gln	Ser	Gly	Val	Glu	Val	Lys	Lys	Pro
Gly	Ala	Ser	Val	Lys	Val	Ser	Cys	Lys	Ala	Ser	Gly	Tyr	Thr
Phe	Thr	Asn	Tyr	Tyr	Met	Tyr	Trp	Val	Arg	Gln	Ala	Pro	Gly
Gln	Gly	Leu	Glu	Trp	Met	Gly	Gly	Ile	Asn	Pro	Ser	Asn	Gly
Gly	Thr	Asn	Phe	Asn	Glu	Lys	Phe	Lys	Asn	Arg	Val	Thr	Leu
Thr	Thr	Asp	Ser	Ser	Thr	Thr	Thr	Ala	Tyr	Met	Glu	Leu	Lys
Ser	Leu	Gln	Phe	Asp	Asp	Thr	Ala	Val	Tyr	Tyr	Cys	Ala	Arg
Arg	Asp	Tyr	Arg	Phe	Asp	Met	Gly	Phe	Asp	Tyr	Trp	Gly	Gln
Gly	Thr	Thr	Val	Thr	Val	Ser	Ser						

FIG.1