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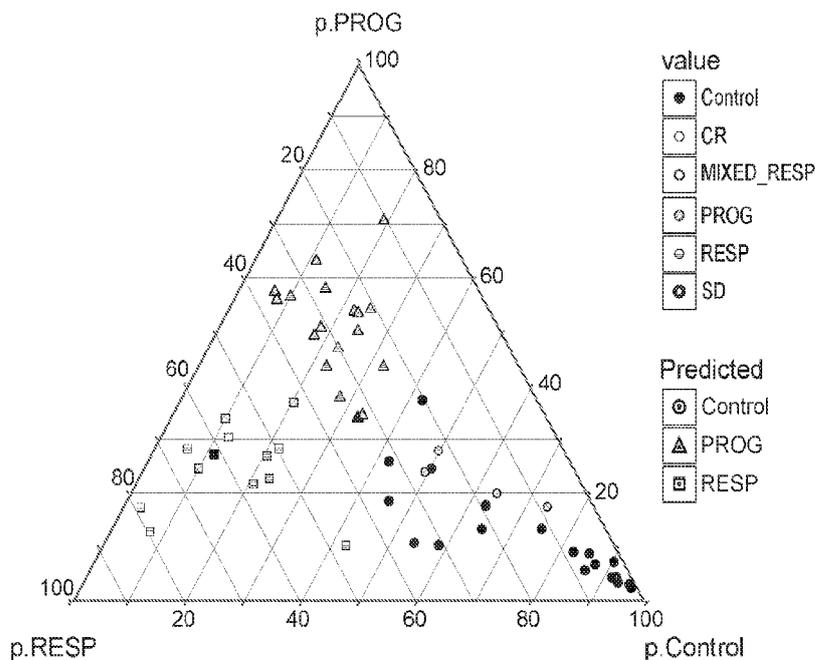


FIG. 1

(57) Abstract: The present invention is directed to methods for detecting a melanoma, methods for determining whether a melanoma is stable or progressive, methods for evaluating the extent of surgery resection in a subject having a melanoma, and methods for determining a response by a subject having a melanoma to a therapy.

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METHODS FOR MELANOMA DETECTION

Cross-Reference to Related Applications

[0001] This application claims the benefit of and priority to U.S. Provisional Application No. 62/511,058, filed on May 25, 2017, the contents of which are hereby incorporated by reference.

Incorporation by Reference of Sequence Listing

[0002] The contents of the text file named "LBIO-001_001WO SEQ LISTING.txt", which was created on May 12, 2018 and is 265 kB in size, are hereby incorporated by reference in their entireties.

Field of the Invention

[0003] The present invention relates to melanoma detection.

Background of the Invention

[0004] Melanoma is a common (~24-35/100,000 incidence – US), highly aggressive, skin cancer with an incidence that continues to rise. The most common, cutaneous melanomas, are associated with UV exposure and immune dysregulation. As a group, melanoma is known to carry the highest mutational burden (>10mutations/Mb). Major mutations include BRAF (~50%), N-Ras (~20%) and NF-1 (~5%), which together, comprise 75% of all mutations. Melanomas are addicted to MAPK pathway activation, regardless of whether tumors exhibit mutations in genes coding for proteins in this pathway. This provides the rationale for targeted therapy e.g., BRAF v600E agents, in this tumor. Other gain-of-function and loss-of-function mutations e.g., in RASopathy genes and amplification of cyclin D1/cdk4 and/or mutation/loss of the tumor suppressor PTEN, also characterize the tumor. This makes melanoma one of the most aggressive and therapy-resistant cancers.

[0005] Five-year survival rates range from 95-100% for stage I, 65-93% for stage II, to 41-71% and 9-28% for stage III and IV, respectively. Surgery, immunotherapy and targeted therapies provide the basis for management, with chemotherapy and radiation as adjuncts. Surgery, however, has a critical role in melanoma care (diagnosis, cure and palliation). Sentinel lymph node biopsy has become widespread, as it provides prognostic information. Melanoma, however, lacks a clinically useful non-invasive e.g., blood-based biomarker of disease activity to help guide patient management by providing predictive or prognostic information.

[0006] Blood-based factors include lactate dehydrogenase (LDH), detecting mutations in circulating tumor (ct) DNA, measurements of circulating tumor cells (CTCs) and circulating mRNA. LDH is typically used to identify aggressive tumor behavior and predict recurrence but its metrics are very low e.g., 30-50% accurate. It is also non-specific for melanoma. Mutations in target genes, like BRAF, can be detected in the blood in ctDNA but its utility as an indicator of therapeutic efficacy is limited e.g., 45-70% accurate. CTCs do not appear to be an accurate marker in melanomas and there is no consensus as to their clinical utility. Circulating microRNAs (miRNA) have been detected but there is no evidence yet for clinical usefulness.

[0007] For melanomas, there are no multi-mRNA circulating biomarkers that function as a diagnostic or as a prognostic for disease recurrence. A biomarker that can be used to monitor the efficacy of surgery or drug therapy in melanomas is currently lacking.

Summary of the Invention

[0008] Among other things, disclosed herein is a 28-gene expression tool for melanoma. It has high sensitivity and specificity (>95%) for the detection of melanoma and can differentiate aggressive untreated disease from stable, treated disease.

[0009] One aspect of the present disclosure relates to a method for detecting a melanoma in a subject in need thereof, comprising: (1) determining the expression level of at least 29 biomarkers from a test sample from the subject by contacting the test sample with a plurality of agents specific to detect the expression of the at least 29 biomarkers, wherein the at least 29 biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1,

FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, YY2, and at least one housekeeping gene; (2) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of the at least one housekeeping gene, thereby obtaining a normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (3) inputting each normalized expression level into an algorithm to generate a score; (4) comparing the score with a first predetermined cutoff value; and (5) producing a report, wherein the report identifies the presence of a melanoma in the subject when the score is equal to or greater than the first predetermined cutoff value or identifies the absence of a melanoma in the subject when the score is below the first predetermined cutoff value, wherein the first predetermined cutoff value is 20 on a scale of 0 to 100.

[0010] In some embodiments, the method further comprises treating the subject identified as having a melanoma with surgery or drug therapy.

[0011] Another aspect of the present disclosure relates to a method for determining whether a melanoma in a subject is stable or progressive, comprising: (1) determining the expression level of at least 29 biomarkers from a test sample from the subject by contacting the test sample with a plurality of agents specific to detect the expression of the at least 29 biomarkers, wherein the at least 29 biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, YY2, and at least one housekeeping gene; (2) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5,

SPATS2L, TDRD7, TXK, and YY2 to the expression level of the at least one housekeeping gene, thereby obtaining a normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (3) inputting each normalized expression level into an algorithm to generate a score; (4) comparing the score with a second predetermined cutoff value; and (5) producing a report, wherein the report identifies that the melanoma is progressive when the normalized expression level is equal to or greater than the second predetermined cutoff value or identifies that the melanoma is stable when the normalized expression level is below the second predetermined cutoff value, wherein the second predetermined cutoff value is 50 on a scale of 0 to 100.

[0012] Another aspect of the present disclosure relates to a method for evaluating the extent of surgical resection in a subject having a melanoma, comprising: (1) determining the expression level of at least 29 biomarkers from a test sample from the subject after the surgical resection by contacting the test sample with a plurality of agents specific to detect the expression of the at least 29 biomarkers, wherein the at least 29 biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, YY2, and at least one housekeeping gene; (2) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of the at least one housekeeping gene, thereby obtaining a normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (3) inputting each normalized expression level into an algorithm to generate a score; (4) comparing the score with a third predetermined cutoff value; and (5) producing a report, wherein

the report identifies that the surgical resection does not remove the entire melanoma when the normalized expression level is equal to or greater than the third predetermined cutoff value or identifies that the surgical resection removes the entire melanoma when the normalized expression level is below the third predetermined cutoff value, wherein the third predetermined cutoff value is 20 on a scale of 0 to 100.

[0013] In some embodiments, the report further identifies that the risk of melanoma recurrence is high when the normalized expression level is equal to or greater than the third predetermined cutoff value or identifies that the risk of melanoma recurrence is low when the normalized expression level is below the third predetermined cutoff value.

[0014] Yet another aspect of the present disclosure relates to a method for determining a response by a subject having a melanoma to a therapy, comprising: (1) determining a first expression level of at least 28 biomarkers from a first test sample from the subject at a first time point by contacting the first test sample with a plurality of agents specific to detect the expression of the at least 28 biomarkers, wherein the 28 biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (2) determining a second expression level of the at least 28 biomarkers from a second test sample from the subject at a second time point by contacting the second test sample with a plurality of agents specific to detect the expression of the at least 28 biomarkers, wherein the second time point is after the first time point and after the administration of the therapy to the subject; (3) comparing the first expression level with the second expression level; and (4) producing a report, wherein the report identifies that the subject is responsive to the therapy when the second expression level is significantly decreased as compared to the first expression level.

[0015] In some embodiments, the first time point is prior to the administration of the therapy to the subject. In some embodiments, the first time point is after the administration of the therapy to the subject. In some embodiments, the therapy comprises an immunotherapy or a targeted therapy (e.g., a BRAF inhibitor).

[0016] In some embodiments, the at least one housekeeping gene is selected from the group consisting of ALG9, SEPN, YWHAQ, VPS37A, PRRC2B, DOPEY2, NDUFB11, ND4, MRPL19, PSMC4, SF3A1, PUM1, ACTB, GAPD, GUSB, RPLP0, TFRC, MORF4L1, 18S, PPIA, PGK1, RPL13A, B2M, YWHAZ, SDHA, HPRT1, TOX4, and TPT1.

[0017] In some embodiments, the at least one housekeeping gene comprises TOX4 and TPT1.

[0018] In some embodiments, the normalized expression level is obtained by: (1) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of TOX4, thereby obtaining a first normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (2) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of TPT1, thereby obtaining a second normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; and (3) averaging the first normalized expression level and the second normalized expression level to obtain the normalized expression level.

[0019] In some embodiments, the method can have a specificity, sensitivity, and/or accuracy of at least 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99%. In some embodiments, the method has a sensitivity of greater than 90%. In some embodiments, the method has a specificity of greater than 90%.

[0020] In some embodiments, the biomarker is RNA, cDNA, or protein. When the biomarker is RNA, the RNA can be reverse transcribed to produce cDNA, and the produced cDNA expression level is detected. In some embodiments, the expression level of the biomarker is detected by forming a complex between the biomarker and a labeled probe or primer. When the biomarker is RNA or cDNA, the RNA or cDNA can be detected by forming a complex between the RNA or cDNA and a labeled nucleic acid probe or primer. When the biomarker is protein, the protein can be detected by forming a complex between the protein and a labeled antibody. In some embodiments, the label is a fluorescent label.

[0021] In some embodiments, the test sample is blood, serum, plasma, or neoplastic tissue.

[0022] In some embodiments, the first predetermined cutoff value can be derived from a plurality of reference samples obtained from subjects free of a neoplastic disease. The second predetermined cutoff value can be derived from a plurality of reference samples obtained from subjects whose melanomas are being adequately controlled by therapies like immune therapy. The third predetermined cutoff value can be derived from a plurality of reference samples obtained from subjects whose melanomas have been completely removed by surgery and they are considered “disease free.” In some embodiments, each reference sample can be blood, serum, plasma, or non-neoplastic tissue.

[0023] In some embodiments, the subject in need thereof is a subject diagnosed with a melanoma, a subject having at least one melanoma symptom, or a subject having a predisposition or familial history for developing a melanoma. In some embodiments, the subject is a human.

[0024] In some embodiments, the algorithm is XGB, RF, glmnet, cforest, CART, treebag, knn, nnet, SVM-radial, SVM-linear, NB, NNET, or mlp.

Brief Description of the Drawings

[0025] FIG. 1 is a graph showing visualization of the melanoma score as a system of three contributors to the clinical picture - Control, Response, and Progression. Samples towards each of the corner represent pure representations of each clinical group. Samples in the middle are in the area of both algorithmic and clinical uncertainty.

[0026] FIG. 2 is a graph showing the metrics for the test in the test set ranged from 87-100%.

[0027] FIGs. 3A-3C are a set of graphs showing the evaluation of the circulating melanoma gene test (Melanomx) in test set 2. Values were significantly higher in melanoma samples than in controls (FIG. 3A). Patients who were responding to therapy had values similar to controls. Receiver operator curve analysis of test set 2 identifying the AUC for differentiating melanoma from controls was >0.95 (FIG. 3B). The metrics for the test ranged from 78-92% (FIG. 3C).

[0028] FIGs. 4A-4B are a set of graphs showing the effect of surgery on the Melanomx. Levels were significantly decreased by surgery ($p<0.0001$) (FIG. 4A). Values in the NED (no evidence of disease after surgery) group were significantly lower than in those with residual disease after surgery ($p=0.0007$) (FIG. 4B).

[0029] FIG. 5 is a graph showing the effect of therapy on the Melanomx score. Levels were significantly decreased by immunotherapy (ipilimumab) or a BRAF inhibitor (Vemurafenib).

[0030] FIGs. 6A-6B are a set of graphs showing Melanomx score in 3 different melanoma cell lines. FIG. 6A identifies the cell lines demonstrate elevated expression – Melanomx score ranging from 40 (A375) to 95 (Hs294). FIG. 6B identifies that spiking these cells into blood from a subject that does not have a melanoma, resulted in detectable gene expression and scores. A minimum of 1 cell/ml of blood could be consistently identified.

[0031] FIGs. 7A-7B are a set of graphs showing expression in tumor tissue and its correlation with blood samples collected at the same time. In FIG. 7A, the Melanomx score ranged 40-97 in melanoma tumor tissue. In contrast, normal epithelium exhibited values <20 . In FIG. 7B, gene expression in tumor tissue is compared to matched blood samples. This is highly concordant (correlation ~ 0.80).

Detailed Description of the Invention

[0032] The details of the invention are set forth in the accompanying description below. Although methods and materials similar or equivalent to those described herein can be used in the practice or testing of the present invention, illustrative methods and materials are now

described. Other features, objects, and advantages of the invention will be apparent from the description and from the claims. In the specification and the appended claims, the singular forms also include the plural unless the context clearly dictates otherwise. Unless defined otherwise, 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 invention belongs. All patents and publications cited in this specification are incorporated herein by reference in their entireties.

[0033] Early signs of melanoma include changes to the shape or color of existing moles or, in the case of nodular melanoma, the appearance of a new lump anywhere on the skin. At later stages, the mole may itch, ulcerate or bleed. Visual inspection is the most common diagnostic technique. Melanoma can be divided into the following types: lentigo maligna, lentigo maligna melanoma, superficial spreading melanoma, acral lentiginous melanoma, mucosal melanoma, nodular melanoma, polypoid melanoma, and desmoplastic melanoma.

[0034] Measurements of circulating melanoma transcripts -- the Melanomx -- identify melanomas and decreases in the Melanomx score correlate with the efficacy of therapeutic interventions such as immunotherapy and targeted therapy. Targeted gene expression profile of RNA can be isolated from the peripheral blood of patients with melanoma. This expression profile is evaluated in an algorithm and converted to an output (prediction). It can identify active disease, provide an assessment of treatment responses, or predict risk of relapse, in conjunction with standard clinical assessment and imaging.

[0035] In one aspect, the present disclosure provides a method for detecting a melanoma in a subject in need thereof, including: (1) determining the expression level of at least 29 biomarkers from a test sample from the subject by contacting the test sample with a plurality of agents specific to detect the expression of the at least 29 biomarkers, wherein the at least 29 biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOHI2CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, YY2, and at least one housekeeping gene; (2) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1,

LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of the at least one housekeeping gene, thereby obtaining a normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (3) inputting each normalized expression level into an algorithm to generate a score; (4) comparing the score with a first predetermined cutoff value; and (5) producing a report, wherein the report identifies the presence of a melanoma in the subject when the score is equal to or greater than the first predetermined cutoff value or identifies the absence of a melanoma in the subject when the score is below the first predetermined cutoff value, wherein the first predetermined cutoff value is 20 on a scale of 0 to 100.

[0036] In some embodiments, the at least one housekeeping gene is selected from the group consisting of ALG9, SEPN, YWHAQ, VPS37A, PRRC2B, DOPEY2, NDUFB11, ND4, MRPL19, PSMC4, SF3A1, PUM1, ACTB, GAPD, GUSB, RPLP0, TFRC, MORF4L1, 18S, PPIA, PGK1, RPL13A, B2M, YWHAZ, SDHA, HPRT1, TOX4, and TPT1.

[0037] In some embodiments, the at least one housekeeping gene comprises TOX4 and TPT1. In some embodiments, the normalized expression level is obtained by: (1) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of TOX4, thereby obtaining a first normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (2) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2,

SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of TPT1, thereby obtaining a second normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; and (3) averaging the first normalized expression level and the second normalized expression level to obtain the normalized expression level.

[0038] Among the provided methods are those that are able to classify or detect a melanoma. In some embodiments, the provided methods can identify or classify a melanoma in a human blood sample. In some examples, the methods can provide such information with a specificity, sensitivity, and/or accuracy of at least 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99%.

[0039] The agents can be any agents for detection of the biomarkers, and typically are isolated polynucleotides or isolated polypeptides or proteins, such as antibodies, for example, those that specifically hybridize to or bind to the at least 29 biomarkers.

[0040] The biomarker can be RNA, cDNA, or protein. When the biomarker is RNA, the RNA can be reverse transcribed to produce cDNA (such as by RT-PCR), and the produced cDNA expression level is detected. The expression level of the biomarker can be detected by forming a complex between the biomarker and a labeled probe or primer. When the biomarker is RNA or cDNA, the RNA or cDNA detected by forming a complex between the RNA or cDNA and a labeled nucleic acid probe or primer. The complex between the RNA or cDNA and the labeled nucleic acid probe or primer can be a hybridization complex.

[0041] When the biomarker is protein, the protein can be detected by forming a complex between the protein and a labeled antibody. The label can be any label, for example a fluorescent label, chemiluminescence label, radioactive label, etc. The protein level can be measured by methods including, but not limited to, immunoprecipitation, ELISA, Western blot analysis, or immunohistochemistry using an agent, e.g., an antibody, that specifically detects the protein encoded by the gene.

[0042] In some embodiments, the methods are performed by contacting the test sample with one of the provided agents, more typically with a plurality of the provided agents, for example, a set of polynucleotides that specifically bind to the at least 29 biomarkers. In some embodiments, the set of polynucleotides includes DNA, RNA, cDNA, PNA, genomic DNA, or synthetic oligonucleotides. In some embodiments, the methods include the step of isolating RNA from the test sample prior to detection, such as by RT-PCR, e.g., QPCR. Thus, in some embodiments, detection of the melanoma biomarkers, such as expression levels thereof, includes detecting the presence, absence, or amount of RNA. In one example, the RNA is detected by PCR or by hybridization.

[0043] In some embodiments, the polynucleotides include sense and antisense primers, such as a pair of primers that is specific to each of the at least 29 biomarkers. In one aspect of this embodiment, the detection of the at least 29 biomarkers is carried out by PCR, typically quantitative or real-time PCR. For example, in one aspect, detection is carried out by producing cDNA from the test sample by reverse transcription; then amplifying the cDNA using the pairs of sense and antisense primers that specifically hybridize to the panel of at least 28 biomarkers, and detecting products of the amplification.

[0044] The test sample can be any biological fluid obtained from the subject. Preferably, the test sample is blood, serum, plasma or neoplastic tissue.

[0045] The first predetermined cutoff value can be derived from a plurality of reference samples obtained from subjects free of a neoplastic disease. Each reference sample can be any biological fluid obtained from a subject not having, showing symptoms of or diagnosed with a neoplastic disease. In some embodiments, the reference sample is blood, serum, plasma, or non-neoplastic tissue.

[0046] The subject in need thereof can be a subject diagnosed with a melanoma, a subject having at least one melanoma symptom or a subject having a predisposition or familial history for developing a melanoma. The subject can be any mammal. Preferably, the subject is human. The terms "subject" and "patient" are used interchangeably herein.

[0047] The score is the Melanomx score, which has a scale of 0 to 100. The Melanomx score is the product of a classifier built from predictive classification algorithms, e.g., XGB, RF, glmnet, cforest, CART, treebag, knn, nnet, SVM-radial, SVM-linear, NB, NNET, or mlp. The algorithm analyzes the data (i.e., expression levels) and then assigns a score.

[0048] The method can further include treating the subject identified as having a melanoma with surgery, drug therapy, radiation therapy, or a combination thereof. The drug therapy can be an immunotherapy, a targeted therapy, a chemotherapy, or a combination thereof. In some embodiments, the drug therapy includes an immunotherapy. Examples of immunotherapies for treating a melanoma include, but are not limited to, Imlygic (T-VEC), Yervoy in combination with Opdivo, Opdivo (nivolumab), Keytruda (pembrolizumab), Yervoy (ipilimumab), Interleukin-2 (IL-2), and Interferon alpha 2-b. In some embodiments, the drug therapy includes a targeted therapy such as a BRAF inhibitor. Examples of targeted therapies for treating a melanoma include, but are not limited to, Zelboraf in combination with Cotellic (cobimetinib), Tafenlar in combination with Mekinist, Tafenlar (dabrafenib), Mekinist (trametinib), and Zelboraf (vemurafenib). In some embodiments, the drug therapy includes a chemotherapy. In some embodiments, the chemotherapy includes dacarbazine.

[0049] The present disclosure also provides a method for determining whether a melanoma in a subject is stable or progressive, including: (1) determining the expression level of at least 29 biomarkers from a test sample from the subject by contacting the test sample with a plurality of agents specific to detect the expression of the at least 29 biomarkers, wherein the at least 29 biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, YY2, and at least one housekeeping gene; (2) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of the at least one housekeeping gene, thereby obtaining a normalized expression level of each of ATL1,

ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (3) inputting each normalized expression level into an algorithm to generate a score; (4) comparing the score with a second predetermined cutoff value; and (5) producing a report, wherein the report identifies that the melanoma is progressive when the normalized expression level is equal to or greater than the second predetermined cutoff value or identifies that the melanoma is stable when the normalized expression level is below the second predetermined cutoff value, wherein the second predetermined cutoff value is 50 on a scale of 0 to 100.

[0050] The second predetermined cutoff value can be derived from a plurality of reference samples obtained from subjects whose melanomas are being adequately controlled by therapies like immune therapy.

[0051] Surgical resection is a procedure that removes melanoma tissues from the subject in need thereof. The present disclosure also provides a method for evaluating the extent of surgical resection in a subject having a melanoma, including: (1) determining the expression level of at least 29 biomarkers from a test sample from the subject after the surgical resection by contacting the test sample with a plurality of agents specific to detect the expression of the at least 29 biomarkers, wherein the at least 29 biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, YY2, and at least one housekeeping gene; (2) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of the at least one housekeeping gene, thereby obtaining a normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L,

TDRD7, TXK, and YY2; (3) inputting each normalized expression level into an algorithm to generate a score; (4) comparing the score with a third predetermined cutoff value; and (5) producing a report, wherein the report identifies that the surgical resection does not remove the entire melanoma when the normalized expression level is equal to or greater than the third predetermined cutoff value or identifies that the surgical resection removes the entire melanoma when the normalized expression level is below the third predetermined cutoff value, wherein the third predetermined cutoff value is 20 on a scale of 0 to 100.

[0052] The third predetermined cutoff value can be derived from a plurality of reference samples obtained from subjects whose melanoma disease has been completely removed by surgery and they are considered “disease free.”

[0053] When it is determined that the surgical resection does not remove the entire melanoma, the subject is at risk of melanoma recurrence. Accordingly, in some embodiments, the report further identifies that the risk of melanoma recurrence is high when the normalized expression level is equal to or greater than the third predetermined cutoff value or identifies that the risk of melanoma recurrence is low when the normalized expression level is below the third predetermined cutoff value.

[0054] The present disclosure also provides a method for determining a response by a subject having a melanoma to a therapy, comprising: (1) determining a first expression level of at least 28 biomarkers from a first test sample from the subject at a first time point by contacting the first test sample with a plurality of agents specific to detect the expression of the at least 28 biomarkers, wherein the 28 biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (2) determining a second expression level of the at least 28 biomarkers from a second test sample from the subject at a second time point by contacting the second test sample with a plurality of agents specific to detect the expression of the at least 28 biomarkers, wherein the second time point is after the first time point and after the administration of the therapy to the subject; (3) comparing the first expression level with the second expression level; and (4) determining that the subject is

responsive to the therapy when the second expression level is significantly decreased as compared to the first expression level.

[0055] In some embodiments, the methods can predict treatment responsiveness to, or determine whether a patient has become clinically stable following, or is responsive or non-responsive to, a melanoma treatment, such as a surgical intervention or drug therapy (for example, an immunotherapy or targeted therapy). In some cases, the methods can do so with a specificity, sensitivity, and/or accuracy of at least 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99%. In some cases, it can differentiate between treated and untreated melanoma with a specificity, sensitivity, and/or accuracy of at least 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99%.

[0056] In some embodiments, the first and second test samples can be of the same type. In some embodiments, the first and second test samples can be of different types.

[0057] In some embodiments, the therapy can be a drug therapy. The drug therapy can be an immunotherapy, a targeted therapy, a chemotherapy, or a combination thereof. In some embodiments, the therapy can be a radiation therapy.

[0058] In some embodiments, the first time point is prior to the administration of the therapy to the subject. In some embodiments, the first time point is after the administration of the therapy to the subject. The second time point can be a few days, a few weeks, or a few months after the first time point. For example, the second time point can be at least 1 day, at least 7 days, at least 14 days, at least 30 days, at least 60 days, or at least 90 days after the first time point.

[0059] In some embodiments, the second expression level is significantly decreased as compared to the first expression level when the second expression level is at least 10% less than the first expression level. In some embodiments, the second expression level is significantly decreased as compared to the first expression level when the second expression level is at least 20% less than the first expression level. In some embodiments, the second expression level is significantly decreased as compared to the first expression level when the second expression level is at least 30% less than the first expression level. In some embodiments, the second expression level is significantly decreased as compared to the first expression level when the

second expression level is at least 40% less than the first expression level. In some embodiments, the second expression level is significantly decreased as compared to the first expression level when the second expression level is at least 50% less than the first expression level. In some embodiments, the second expression level is significantly decreased as compared to the first expression level when the second expression level is at least 60% less than the first expression level. In some embodiments, the second expression level is significantly decreased as compared to the first expression level when the second expression level is at least 70% less than the first expression level. In some embodiments, the second expression level is significantly decreased as compared to the first expression level when the second expression level is at least 80% less than the first expression level. In some embodiments, the second expression level is significantly decreased as compared to the first expression level when the second expression level is at least 90% less than the first expression level.

[0060] In some embodiments, the method further comprises determining a third expression level of the at least 28 biomarkers from a third test sample from the subject at a third time point by contacting the third test sample with a plurality of agents specific to detect the expression of the at least 28 biomarkers, wherein the third time point is after the second time point. The method can further comprise creating a plot showing the trend of the expression level change.

[0061] The present disclosure also provides an assay comprising: (1) determining the expression level of biomarkers consisting essentially of the following 30 biomarkers from a test sample from a patient diagnosed of a melanoma or a subject suspected of having a melanoma: ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, YY2, TOX4, and TPT1, wherein the expression level is measured by contacting the test sample with a plurality of agents specific to detect the expression of the 30 biomarkers; (2) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of

TOX4, thereby obtaining a first normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (3) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of TPT1, thereby obtaining a second normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (4) averaging the first normalized expression level and the second normalized expression level to obtain a normalized expression level for each biomarker; (5) inputting each normalized expression level into an algorithm to generate a score; and (6) comparing the score with a first predetermined cutoff value.

[0062] The present disclosure also provides an assay comprising: (1) determining the expression level of biomarkers consisting of the following 30 biomarkers from a test sample from a patient diagnosed of a melanoma or a subject suspected of having a melanoma: ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, YY2, TOX4, and TPT1, wherein the expression level is measured by contacting the test sample with a plurality of agents specific to detect the expression of the 30 biomarkers; (2) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of TOX4, thereby obtaining a first normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR,

CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (3) normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of TPT1, thereby obtaining a second normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2; (4) averaging the first normalized expression level and the second normalized expression level to obtain a normalized expression level for each biomarker; (5) inputting each normalized expression level into an algorithm to generate a score; and (6) comparing the score with a first predetermined cutoff value.

[0063] The sequence information of the melanoma biomarkers and housekeepers is shown in Table 1.

[0064] Table 1. Melanoma Biomarker/Housekeeper Sequence Information

Gene Name	RefSeq Accession	Sequence	SEQ ID NO:
ATL1	NM_001127713.1	AGTTTGAGGC TGCAC TGCAC ACCTGCACAA CCTGCCTTCT ACTTAGTTCT TCTGAGACAT TTCTGAAAGT CTGAATTCCT AGGACTGCTC AATGACCTTT GTCCCTGTTG GGCACACGCA GTGTCTCATC GCTGGTATTG CACCTTTAAT GAGACCAGGA GTTCCGCAAA AGTAAAACAA AGGGGTCACA GACTGGTTCT GGTTCACCA CTTCCTCAGT ATGTGCTCTG GGACAAAACA GCATATTTGG CAACATCTCG GTGTCCTTAT CTGCAGCTTG AAGAGGCGCC ACAGCAACAT CCTCAGAGTC TGAGCGAACT GCGCCCAGCG CGGGCACGGA GCCTCCCACC GCCAGCAACC TGCGGCCCCG GAGAAGGCAG CGAGCGCAGT GACAGCGCCT CACCGCCACC AGCTCCTGGA CCACCATGGC CAAGAACCGC AGGGACAGAA ACAGTTGGGG TGGATTTTCG GAAAAGACAT	1

	<p>ATGAATGGAG CTCAGAAGAG GAGGAGCCAG TGAAAAAGGC AGGACCAGTC CAAGTCCTCA TTGTCAAAGA TGACCATTCC TTTGAGTTAG ATGAAACTGC ATTAATCGG ATCCTTCTCT CGGAGGCTGT CAGAGACAAG GAGGTTGTTG CTGTATCTGT TGCTGGAGCA TTTAGAAAAG GAAAATCATT CCTGATGGAC TTCATGTTGA GATACATGTA CAACCAGGAA TCAGTTGATT GGGTTGGAGA CTACAATGAA CCATTGACTG GTTTTTCATG GAGAGGTGGA TCTGAGCGAG AGACCACAGG AATTCAGATA TGGAGTGAAA TCTTCCTTAT CAATAAACCT GATGGTAAAA AGGTTGCAGT GTTATTGATG GATACTCAGG GAACCTTTGA TAGTCAGTCA ACTTTGAGAG ATTCAGCCAC AGTATTTGCC CTTAGCACAA TGATCAGCTC AATACAGGTA TATAACTTAT CCCAAAATGT CCAGGAGGAT GATCTTCAGC ACCTCCAGCT TTTCACTGAG TATGGCAGAC TGGCAATGGA GGAAACATTC CTGAAGCCAT TTCAGAGTCT GATATTTCTT GTTCGAGACT GGAGTTTCCC ATACGAATTT TCATATGGAG CCGATGGTGG TGCCAAATTC TTGGAAAAAC GCCTCAAGGT CTCAGGGAAC CAGCATGAAG AACTACAGAA CGTCAGAAAA CACATCCATT CCTGTTTCAC CAACATTTCC TGTTTTCTGC TACCTCATCC TGGCTTAAAA GTAGCTACCA ATCCAAACTT TGATGGAAAA TTGAAAGAAA TAGATGATGA ATTCATCAA AACTTGAAAA TACTGATTCC TTGGCTACTT AGTCCCAGAGA GCCTAGATAT TAAAGAGATC AATGGGAATA AAATCACCTG CCGGGGTCTG GTGGAGTACT TCAAGGCTTA TATAAAGATC TATCAAGGTG AAGAATTACC ACATCCCAA TCCATGTTAC AGGCCACAGC AGAAGCTAAC AATTTAGCAG CCGTGGCAAC TGCCAAGGAC ACATACAACA AAAAAATGGA AGAGATTTGT GGTGGTGACA AACCATTTCT GGCCCCAAT GACTTGCAGA CCAAACACCT GCAACTTAAG GAAGAATCTG TGAAGCTATT CCGAGGGGTG AAGAAGATGG GTGGGGAAGA ATTTAGCCGG CGTTACCTGC AGCAGTTGGA GAGTGAAATA GATGAACTTT ACATCCAATA TATCAAGCAC AATGATAGCA AAAATATCTT CCATGCAGCT CGTACCCAG CCACACTGTT TGTAGTCATC TTTATCACAT ATGTGATTGC TGGTGTGACT GGATTCATTG GTTTGGACAT CATAGCTAGC CTATGCAATA TGATAATGGG ACTGACCCTT ATCACCTGT GCACTTGGGC ATATATCCGG TACTCTGGAG AATACCGAGA GCTGGGAGCT GTAATAGACC AGGTGGCTGC AGCTCTGTGG GACCAGGCTT TGTACAAGCT TTACAGTGCA GCAGCAACCC ACAGACATCT GTATCATCAA GCTTCCCTA CACCAAAGTC GGAATCTACT GAACAATCAG AAAAGAAAAA AATGTAATGC AAATTTAAG AAATACAGGT GCATGACCAA TTGTCAATTA AATATTCAGT TTTATGTCTC CATGCAAACA</p>	
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		<p>TTCAAAGTGC TTCCATCAGA ACGGAGTAAA ATACTAAACA CCTCTGAAGA CTGCAAACCTG GATTAGTTCT TTTACTTCAG TGTTTAATAA GCAGATGTAT GTATGCATGG TTATACTATT TTGTTAACAT GTACAATTTT CTGATTTTTT TTCAAAAATG CTGTTATAAA GTATTTGTCT ATTTATGATA ACAGTACACG TGTTCTGCTT GAATTTACTA AATTTCTACTA CTGGGTTATA ATTAATCAT GTGATATTCC ACGTTTGGAT ATGCTCATT AATTTCTACA GAAAAAATTT TAAATTATTT CACATTAGCC ATTTGTAAA ACACAGCATC ATAACTCAGC AGGCTGGATT TAATCTGTAT CATCTTATAT ATATCACAAT CTTATTTTTA AGCACATTTT AGAGTTCCTT AGTTGCTTTA TCAAAAACCA GATATTGCTT TTACATGGTT TAATAGAATA TAAACCTCTT GATAAAAAAT GCACAAAAAA TCACTTTGTA TATGTGAGTT TCACTGCATT GTATATTTTT TCATTTGGTA CACAAAGAAT GTATICTTCA TAGGTTTATT CTTTTAATAT GTGAACTATT ATTAAGTTT ACTCTGGTTC CTAAGATTAA AAACAAATGC TTACTGAATT TGAAAAAAA A</p>	
<p>ATP6V 0D</p>	<p>NM_0 04691. 4</p>	<p>ACTTGACAGC CCGCTGAGGA CGCAGCGTCA GCTGACCTGG GGAGTCGCGA TTCGTGCCGG CCGGTCCTGG TTCTCCGGTC CCGCCGCTCC CGCAGCAGCC ATGTCGTTCT TCCCGGAGCT TTACTIONAAC GTGGACAATG GCTACTTGGG GGGACTGGTG CGCGGCCTGA AGGCCGGGGT GCTCAGCCAG GCCGACTACC TCAACCTGGT GCAGTGCAG ACGCTAGAGG ACTTGAAACT GCATCTGCAG AGCACTGATT ATGGTAACTT CCTGGCCAAC GAGGCATCAC CTCTGACGGT GTCAGTCATC GATGACCGGC TCAAGGAGAA GATGGTGGTG GAGTTCCGCC ACATGAGGAA CCATGCCTAT GAGCCACTCG CCAGCTTCCT AGACTTCATT ACTTACAGTT ACATGATCGA CAACGTGATC CTGCTCATCA CAGGCACGCT GCACCAGCGC TCCATCGCTG AGCTCGTGCC CAAGTGCCAC CCACTAGGCA GCTTCGAGCA GATGGAGGCC GTGAACATTG CTCAGACACC TGCTGAGCTC TACAATGCCA TTCTGGTGGA CACGCCTCTT GCGGCTTTTT TCCAGGACTG CATTTCAGAG CAGGACCTTG ACGAGATGAA CATCGAGATC ATCCGCAACA CCCTTACAA GGCCTACCTG GAGTCCTTCT ACAAGTTCTG CACCCTACTG GCGGGGACTA CGGCTGATGC CATGTGCCCC ATCCTGGAGT TTGAAGCAGA CCGCCGCGCC TTCATCATCA CCATCAATTC TTTCGGCACA GAGCTGTCCA AAGAGGACCG TGCCAAGCTC TTTCCACACT GTGGGCGGCT CTACCCTGAG GGCCTGGCGC AGCTGGCTCG GGCTGACGAC TATGAACAGG TCAAGAACGT GGCCGATTAC TACCCGGAGT ACAAGCTGCT CTTGAGGGT GCAGGTAGCA ACCCTGGAGA CAAGACGCTG GAGGACCGAT TCTTTGAGCA CGAGGTAAAG CTGAACAAGT TGGCCTTCTT GAACCAGTTC CACTTTGGTG TCTTCTATGC CTTGCTGAAG CTCAAGGAGC AGGAGTGTGC CAACATCGTG TGGATCGCTG AATGTATCGC CCAGCGCCAC</p>	<p>2</p>

		<p>CGCGCCAAAA TCGACAATA CATCCCTATC TTCTAGCGTC CTGGCCCAAG GCTCTCAATT GCACTCTTTG TGTGTGTGTG TGTGTGTGTG CGCGTGTGTG TGCGTGTGTG TGTATGTGGT CTGTGACAAG CCTGTGGCTC ACCTGCCTGT CCGGGGTGTA GTACGCTGTC CTAGCGGCTG CCCAGTTCTC CTGACCCTCT TAGAGACTGT TCTTAGGCCT GAAAAGGGGC TGGGCACCCC CCCCCACCA GGATGGACGA AGACCCCTC CAGAGCAAGG AGGCCCCCTC AGCCCTGTGG TTACAGCCGC TGATGTATCT AAGAAGCATG TCACTTTCAT GTTCCTCCCT AACTCCCTGA CCTGAGAACC CTGGGGCCTG GGGGCAGTTT GAGCCTCCTC TCCCTTCTGT GGGTCGCTCC CAGAGCCATG GCCCATGGGA AGGACAGAGT GTGTGTGTCC TTGGGGCCTG GGGGGATGTT GCTCCTCAGC TCCCTCCCTC AGCCCTGCC CTCTGAGACA ATAAACTGC CCTCTCTAAG GCCAACTGTC AAAAAAAAAA AAAAAAAAA</p>	
<p>C1ORF 21</p>	<p>NM_0 30806. 3</p>	<p>CCCTCCCTCG CTCGCTCCTC GCAAGCTCCC GCTCGCTCCC TGCCCACTCC CGGGGGGACG TTCCGTGCCG CGGCCGCCGC GGCCGCTGCT TCTTTCACAC TTAGTTGGG AGCTGCGCGC CGCGCTCAGT TACTGGAGAG CTGGCCGCGC GCCGCCGCT CCCGCACGCT TGCACGCGGG CCCGGCTTCG GGGTTTTGGG TTCTTACTCC AAGCGGCGGG GAGGAGGGGG AGCCCCGGAC AACTGTGGG GAGGAGGAGG AAGAAGAGGA GGAGGGAGGA AGAAAAAGA CGAGGAGGAC AGGGGCGGGG GGCGGGAGGC TTGCCACCTT CAGCCCCCCC GCGAACGCC AAGGTGCACA CATCTTGACC AACTCAGCAG CAAGGTGGAT TTTCTTTGTG TTTAAAGAAA AAAAATGTCC CTGTGTCTGT AGAGATGATT TGCAGTTCAG CCCGGCTGAA GCTGACCGAA TGAGACTATG GGCTGTGCCT CCGCCAAGCA TGTTGCCACT GTTCAAAATG AAGAGGAAGC CCAGAAAGGG AAAAACTACC AGAACGGAGA TGTGTTTGGC GATGAGTATA GGATCAAACC AGTGGAAGAG GTCAAATACA TGAAAAATGG GGCAGAAGAA GAGCAGAAAA TAGCAGCCAG GAACCAAGAA AACTTGGAAA AAAGTGCCAG CTCAAATGTA AGACTTAAAA CTAATAAAGA GGTTCCGGGA TTAGTTCATC AACCCAGAGC AACATGCAC ATCTCTGAAA GCCAACAAGA ATTCTTCAGA ATGCTGGATG AAAAAATTGA AAAGGGTCGG GATTACTGTT CGGAAGAAGA GGATATCACA TAGCACCAAT TTTACCACTC AAACCAGGAG CTACTACTGT GTAAATAGGT TACACCCAG TTGAAATCTT TGCAAAGGTC GGTTCATTG AGCGAACAGC ACTATAGCAA AAGAAGATCG TTCCATATTG TACGCCCCAT TAAATTACAG TGTTTCTTAA TGAACCTGCA AAGGAATATT GCTAAAAACA AACAAAAAAA ACTGTTATCG AACTTCTTT GTTGCTGCTA GTTAAACTT GTTGCAACTT TCACTTCTC TTGTGTCCAG GTATGCAGCA AAATTCTGCA ATTTACCTT AAAGATACTG TTGGTTTTAC</p>	<p>3</p>

	<p>AGATGCTCTC CAACCTATTT TCTATAAGAT GAGGTAGTGG TGAACTCAGA TAACAAACTT CTCTTCTAAA CTGGTTCTGC TTCTAAGACA AGCATCTCCT GCCCTCTCTC CTCCTCCCC ATCTCTCGCA CGCAGTCTAG AGATGGACTG AGCCTTGCTT CTCACTGGCA GTGTTGAGCT TTGGAGATGG GATGGTTGCT ATGCCAAGCC TTGTTTCTCT GCTCAGAAGA AGTAGAGAAG CTATTATCAA TTAAGAGCAT GCTGTGATGT GACTCCTGGA AGTGACGTAG GAGTGAGTGG CAGGTTGTTT GATTAAATAG GTATCTTAAT CAAGAATTA GCTTGCAACA TTGGCTTTGC TCAGATGCAG ATGGAAGTGT GATCACAATC ATTTTGAATC CCTCTTCCTC ACTTTTTTTT CTAAGAAAAT AACATTTTA CTGTTTTTAT GGATCCTTGT CTTCTCCCAT TCATCCAGCT CAGTGTTTTA AGATGATCCT GGGTGCAGAA GTTGAGCCCT CCTTTGCATT GACTGATA ATTAGCCTAT AGGGCTCCCT ACCTTCCAT TAAGAATCTA CCAAGCATT GCAAGGCTGA AAGTGGTCTA AGAGGTGAGG GGACATCCTA TGACTTTTTA GGAAGGCCTG AAACCACCTT GTTACCTTTC ATTTTGTTAG CAAATAAACC ATCCTATTTT GTAACCTCTC CCCTTCAAAA TGCTACATGA GCCTTGCCAC TTCCTTTTTT TCTTACTTCC AGCACACTAG ACATAGCAAA AGTGTITGCC TACTCAAAAA CATAATACTT TTATGCTGAT GATGGTATTT GGAGATGTGA AAGCCAAAAG CCCCTGGCAG TGGTGGGGAA TGTTGACTGA GTGTTTCAGCA GAGTTTATTT TTCCATACTA TATGAAGAGA ATGATCTCTT CTCAAAGACA GAAGTGATAT TTTTAAACAA TATTGTCACA AGTAAATAGC AATCAAAAGG AGAAAATAAC TTTTGTATTT TTTTAAATGT TTTGATAGCT TTGACGAGGG TTCTCTTGT TACTTTCAGG GGAGGGCATC CTATTAATG CCACGCCAGC AGTCCGGGTC TGGGTTTGTCC CACAAAATC ACAGGAGCAC TGTATGTTC TCTCTTTTGG AGTTGTGACT TTGAAGGGCC TCAATATTAG CCACACTGCC GCCTGCAGAA GGTGGAGAGT TAAGATGTT TATGTCAATT TGCTCTTGCC GAAAAGATGA GCCTCGATTT TAAAATCTAT CCACATCCAA CTGATGGCAC CATTGATGTG CAAATAATGA GATTCCTAT CTCTTTTAG ACCTGGGACG GCAAAAGGGA AGGGAAGGAA ACTTAGCAGA GTGCTATTGA CTATAGATT ACATATTAGC AACAAAATCC CGTAATTCCT TTGGCCAACA GCAGCTATTT TGGGGAGCAG CTGTGGCTGT TACATAAATA GAGATGCAGC CAAATTTTA GGCCTTTTAT CCTGCTTCTA GCAGAAAAAT GCAGGGAGAG TCAAGTAGTC TAGGGTTTCA GGTTGCCTCC CCTCATATGG TTTTGGCCA AGTGACTAAA ACAGTTTTCC ACAACGTAA ACAACTGCT AAGCCCCACC TCAAACCTGT TCACTGGGGA CTTTGCTTAC CGTTCGTGG GTGACCTTTT</p>	
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	<p>CCGGGATTTTCTTGTTCTTAT CAAGCAAGAA TTAAGCACAT GCTAAACGTC TTCCATTTGA CTTCTCTACT CGGTGTCTCA GACAGTGTCT TCCCAGAAAA CCACCACCCT CTACCCAAAAG ATGAAACATG CTCATGTCAT TTTTCTCATG GTCACATTTA ACAGTTTTGA CATGTTATAC TTGCGCATAG ATCCAAGCGT TTCTTGGGAA CCTGACTTTT GAGTGTITAA TAAAGCCGGA AGTGGTGTG CCCTGAACCA GCAGATTTT ACCTGGGTTC TGGCTCCGGT GTTTAACACT GGATACATCT TTGATGTGCG AAAGTGAGTT CATCTTCAGA CACATTTGGT ACATCCAGAA ATAGATCCAA GAAATGGGGT GGTTGAGTGG GTCCGCACGA AATGCTTGAT TATGTCAGCA ACACCCAACA CTGTCTGTTT TCCATTTGTT GGTTTTAATC ATAAAATTGT CAAGTGATTC GTGTTTGTAC TTTATTTTTT TGTGCCTTCT GAAAGGATCT AAAACAAAAA TATTTTGCCT TTTTTTCCC ACGTGTATCT GAACATTAAG CAGATTGGCT CAGACACAAT GAAAAGGATA ATCCAATGTA CGTGCTGGTG CACTCTGCTA GTTGTATCT CTGTAGGGCT CAGGAAGCTG GAAGGAGGAA GGGAGGGTAA GTGGCCTGGT GAGTGGAGGT AGAAAAATGA TGAGAAATGA ACTGAGAGCA TTAAGCAGAG AGGGTTGATA GGCTGGCCGT GTCCGGGGTG AAATTGGAAA TCCAGCTGCC TAGTGGCCAG TGGGTGGGGC AAGACTGTCA ACGAGATTTA CAGCTGGCTT ACACATGCCT TATGTCCTCT GAGTTGTAGA GTTGTAAGG TTCAGCAGTG TGTGCACAGC TTTCTTTTGG TTGGCAGAGA TTCAGGATCA TGGAGTACTG CTCTCATAAT TGAAGACGTG TTTGTTATTG GCAGAGAACC TAAAAAAGG CCTTTACCTC AGCGATGCTT CCTAGCCCCA GGCTTGCAGA GAACACAGAG TGGTGTGTTG GTCTATTTAG GGACAAAGAA GGTATAAAGT CCAGAGATGA GAAACTGGG TCAGCCCTCA GAAACTGCAG CAGCCACGCA CACAGAAGCC TGCTGGAAGA CAGGTCTCTC TCCGTCCACA GTGCCCATCA TCTGAGCCTG GGCTGGGATG ACTCAACTTA GCAAAGACGG ACCCAGGAGG AGTGCTGGTT CTTCAGTCTT TGFACTGGCC CCATCCTCTC CTCACTGTAA TGTGAGGAAG CACCTCTGTG TCAGGGCTCA CCTGGGCATC CAAAGCGGCC ACGCCACAA TCCGACAGCC CCCAGGAGCA GGTCCAGGGA TGATGCAGCC CCCTTCTTGG TCCCATGTGA TGTCATCCTG CTTTGTATC TTCTTATAAC TTTATCCTGC TATAACTTTA TCCTCTTCCC AGCCTCATCC CTGTTTTTCT GTTAGGGCAA GACTCTTCCA TAAGCCTGCT AAAAAACAGA GATGATACCT CTTACAAACT TTACCTCATA GCCTGTGAAG CAGGTTGGCA TGTGGATTAC AAGTCCTGCT TTGACACTGG GCAAGAATTT AAGATTGTTT TATCTCTACT AGTCATAGAA AAGAAACATT GTTAAACATG TTGAGTTTTA AAGGAAGAAA</p>	
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	<p>TATTTTCAA TTCTTAATCC AAGAAAATAC TGAGTTGGAA TCTTAGACTT CGGGACTCTG ACACGTTCTT TATGAAAGGC AAAATAATTG GTATCTAAAG TTCTCTCCTT CCTGCCTCCC CTAAAGAAAA AGGATATTAG ATTGCACACT ATAATTTTAC ATAAGATCTG CCTTCCACAC TTCCCTGCTG GAAGGCATTC TCAGAGCTTT ATGTCTTCGT ACCTCTCAGA GATTGGACTT TTTCTTGTTT AAAACCCCAA CCAAAAAAAAA TAATAAGGCA TGATIGGTGG GGAGGGAATG TGTATTTAGG GGCATAATAA AAAGGTGGCT CGAAGCAGGA ACTTTGGCCT CATGGTGTCA TGGGTGGATG CCTGGACTCC AGTGTGCCTG TGAGGGGCTG GGTTAGGCAG TCGGCTGTCA CACTCACATG TGCCTGCAAT AAACCTTTTG GAATTTTCATG AACGAGGTCT ATGAATTGCC TTTTGCCAAT GAATGGATGT ATTTTCCAA GGGGGGAATA GTATCCTTGA CTTTGGCAGT CACCTTTTTG TATGTCTCTA GAAAGGGGTC AAAAACTAT GGTAAGATG AGGCTTATGA GTGAATACCT CTGGGACAAA CCTTAGGACT CACAAGCTAT GCCATGTTTT TCAGGAGACT CTTGTACCTT ATCTGGAATC TAATCTTGGG AGAAGAGGAA AAAGGAGCTA ATATTTGTCA TTTATACTCA CTCTGTGCCA GATACTGTGC TAGGCATTTT ATAATTGTTT TGTGTCATCC TCATGATAAT CCTGTGAAGT AGATCTATTA CCCCCGTGTT CTAATAATA GATCTTAAGT GTGGAATGCA TCTATCCAAC ATAAATGCC ATGTTGAAAG AAGGAAAGAT GTCATTCAAG TATTTTCAA ATTCTTTTTA TTATGACTAT GCCCTTCGCA AACTGTGAA AACAACCCCT GGGGGCATCT GCCTTCCAGA ATCTCTCTCT GGCTTCTCAC CAGCTTGGTT TCCTCATGGG GAGTGTTTTA TTTGGCCTCC CCTATCTGAG CTGCACACAC ACCAGGGGAG GCCACTGGCT AACAGTAGGA CTTCAGTGCC CTGAGGAAAA GGCTTTGGGA ATTTAGGCAC ACGTTTCTCT CCTTGGAATC CTTCAGCAT CTGGAAAAGG AACCTGGTAT TTCTGCAGTT AGATAACCAGA TGCAACCAGA ACAGGCTTTC GAGTGCTGTG ATTTCTTTCC TGGGTTCCTA AGTCTTGTTG TTTCATCACA AACTGTATCT TTTTAAGGTT AAAAGTCTTG ACCTTCATGG GGGTCTGGGA CAATCCGATC TCCAAGCATG GAGGAAAGGC AATGCCTGGA CCACTGACTT GCATTGAAAT CCTTTCTTGT GGGCTAGGGT TTGATGTCTC TTTTTCATCT TTGGACTGGG GATCTGCATC TTCCAGGTCC ATTTAAGCAC TGAAACTAGA TGCAAATCTC TTTCGAGACC TTACATGTTT TAGATAGTCA TGTAATGACT TGGATAGACA TTAAATAAC TTGTTCCAAG GTCGGAAGAG CCCAAGAACT CTCAGAGTT CCTCTTCTTG CTTCCTGAAT CACATCTCT AAAGATGACT CATGTCTCCA TAGCAACTGT TAAAGGTGCT GCCTAGACGG GACCCGCTCC CACCTTACT</p>	
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	<p>TACTTTGCTC GAAGGAACCA AAACAATGTC CTTGTGTAAA GGGGCTGTCA TATCCAGTTT TCCTTTGAAA TCTGGCCCC AAGATCCTGC TTCTTICTAA CCTGGCAGCT GTCATGTCCC TTCAAGATGA TGTGGGAAAT GGCCCTTACA ACTTGGGAAC CACAGAAATT GCTGTATTTT GGGAAGATTC ACCTCTAAAC TGAAGGCTTC ATTCTGATAG TGTCTGCCCT CTCTACCCTG ATTTGCCCCT TCTTTGCTTC CTTTTTAGC CCAAGGCTTT GAATTTGATT GAGTAAGACT TAGAGGCAGT ATAAAGAACA CCATAAACTT AGGCAGAGGT CCCTTAGGGT CTCTAGAGTT GAAAATAATT CTACAGCCTT AGGGGGACCT CTTGGCATTG ACTCTAAAGG GAGAGAATAG CCCCTGTGTC CTGGCATTTC AGCTAGACC TTCAAGGACT GTTCTCTCTT GACAGGCAAG CAAGCAAAGA AAGTTTTGCA ATAGATTCA AGCCAGTTTT TCCATTCAA CCAAGATGCA AATTCATAAA ATTACTCTTT TCCTGGAATA GATCCAGGCA GCTGCCTTAT TAGAACTTTA GATTCGGATC TATTTTCTTA ACACACATAC ACATACACGC GCATACATAC ATATACAGAG AGATACGTGG AGAAAGGAAA TTTACTCTAT CATTGCAATA CTTCAAGAAA GAGCTGTATT TTGCCTTTCT GTAATCTCCA AGATAGTGTC TAGGAAAGTA ATAGTATAAC TATAGGGATA CCGAAACAGG AAAAACCAGC CATCACTCTT GAGAAAGTTT GAGTTCGACT CACATGGGAG AATCGAGGTC TGCTACTCGT CTTGCTTTGT GCCCATCTG TGCTGGATG CCCTACTACA TCTGCTTGAC TCGTCTGGGC TGCTAGCCGG GGTGTTGTGG CTGACATCCT TTCCTGGCCT TACACACATA ATAGACACAT CCCTAACGGC GTGTGCCTGG TCCAGCCACA TACAGCCACC ACATGTGTCA CACTGTCC CCCTCATCCA TGTGGACTTG ACTGGCATTI CAGCAGCTCC ACTGGGATGC TCTAACCCCA GTGTGTGGAG TTGGGGTCCC TTCATCTAGG TTGACCCAGG TATAGCATTI TTAGCATTGC CTTTCCAGTC TTGATGATTC ATTCATTGAA CTCATTTATT TCTGGAGCCC CTGGTACACT CCAGGCACTG CGCTAATTGC CAGCAAAGCA CAACTGAACT AAATCCACCT TCAAGGAACC TAGCCATAAC GAGGGAGGCA GCATGGAAGT ACCCTACAGG GGAAAGTCCT GAGTGCTGTG GGAGCATCTC ACCGTGGCAG CCAGCCCAGT TTTGGCAATC AGGGGCTTCC TGAAAGAGGT GACATCAAAG CCCGGATGTG TCAGAGGACT GAGGGAGAGT GTTACTAAAG GACTTTCAGG CTGAGAGGAT AGCACAGGAC TCAGCCCAA GGAGGGCCAG TGTGGACTGT CCAGGGCCAG CCTGCAGTAC AGAGGCTGGA GCTTGGACTT GTAGAGGGAG AGAGAAGAGC AAGGGACGTG GACGGGGCAG TGAGCCAGGC TAGCCACAGA GGGTTCCCGG GGCTTTGCTG GGGATTCAGG GAGCATAAAT AAGAGCTTTA GGTGGTGCTG TGTCCTCTGC</p>	
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	<p>AGCCCACTGC TGAGGTCTC CAGACAGGTA AGGTGTGGTC ACAATCAGGG CCGGGGTTTC CCTGCTCACT GCGGCAGTGC AGGGGTGCTT GCTGAGATGA TTCATCCCAG GGTGTCTCT GTCCCTTACC CAGCCCCAAC TCCTCTTCCT CTGCCAAAAG CTATTTGAAT TCAAGGACTT TAACCTGGGC CGGATCTGGT TTGGAGACAA AGGGGACAGC TCTGGGTCAG CATGACCTTC TTTAGAGCCA CTAAGGCGAA AAATACCGTT TGGGACCAGG CTGGCCTAGA CCCAGGGATG AGAATGCACC CTAATAATAA TATACGGGAA GCAGCAGAGG GCTTCCCTGT CTAGTGTGTG ATCCTAACTA AAGGCAGCTC TCTTGGACAG CCTTCCCCTG GATTAGGTCA CATAACCTG GTGGCCAAGC CTCTGCTGGG TCCCAAATAC ACACCCGAGT CCTGCCAAAG AAAGGAGATT TTTAAAAGC ACAGACAAAT TGTATGCAAG TGGAAAATAC CCATAGGCCT AGACAGCTGT GGAGGGAAGA CCTCGTGGGT ACCTGGAGGC TGCCAGAGCT GGGAGCTCTG CAGGTATGAG TCAGGGAAGG CTCAGAGACA AGCAGAATCT CTCTATGGAG ACAAC TTGCA GTGCC TTTTA GGTTTTCAA ATAACCTCGG AGTTCAGAGC ATGGGGTTT TTTCTCCCCT CCCACCCCC AGAAAATAA TTAGAAAAT GTTTAGGAGA AAGGAAAAGA ATTAGATGCA TCAGAATACC AGCTATAAGC CAACACTGTT TCCAGAACT CAAGAAAAG CTCAAACAGA AGACAGTTCC CCTGAGAGGC TGGAGGCGTT GGTGCTGAAG GCAATTTTCC TAGCTAAGGG GCACTGGGCC TTGCTGCACC TTGGGGCTGA CCTTTTTTGC AAAACACCCA CCCCTGCCCT CCTGGCATA TCAACAGCAA CGCCAGCTTT CTGGACCCTT GGAAAGATGT TAGCTCAAAC ACCCACTTTT TCCAGATCTT CCTCTTGCTC TTCCTGAGG AATTGTGAAT TCTGAGGCTA GCGATGCCA CTCGGATATT CCGCAGGCC AGGTGTTTAG ATTAGAATTT GTCCAGCGGT AATCCTGATG CTGGAAACCA ACAAACATTT GGCCTCATAT TCACCCATTT AAAA ACTAGA GCCCCTGGCA GGTCCCCTTA GGGCCATGTG TTCATGGAAT ATAAGCCAAG TTTGCCCTAG GCTTGTTTAT GGAATATAAG CCAAGTTTAC CTCTCCCAT TTTCTGCCCT GGCCCCTTC CCACTCACCT CCACCTCATT GCCAGGAAGG GATCAAAATG CCTCCATGCC AGTTGTTAAT GGCTACATAT TTGCCCTTCC CAAGGGTATT TGCATTTTAT TTAGGAACAT GGCCTTATAT TCAAGGAAAA TCTAGCATCA AGATTACGAG GCATCACCTC TCAATCAGGT CTGGGAGGTA TCTTGGGGCA TTGCTCTTCT GAACACCTGC AGAGGCTTCC TCAGGTGAGT GTGGGAGCCC GGAAGGGTGG CCTCCCTAAC CACTCTGCCT GCACATGAAT TCTCAAAGC AGTGGGCCCC CATCTGTTT AATTACACAT GCCTGTCAGC AAAACTTCG TGAGATGCAC TCTCTCTGTG TGTTTATTAA</p>	
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		<p>TTTATTTAAA GCATATATCC CTTTACTTTT GTACTACTAT ATTAGGCACA TTATAAAAAG TATACAGCAT AGAAACTTTA AATGAATAAG ACACAAAATA TTATAAACAG AGGTTCTGGC ATTTTCTCTC TGAACTCCTG AGGGGGACCT TGGGCACCTC CTGGTATGTG CACACCCAC TTTGAACACC CTTGCTTTAG TGCAAATCAG CACGCCTAAA TAGCATCCCA AACCACTGTG CGACATTTGG CCTGCAGAAT AAGAAAGTCC TAAGGCAGAA TCCCCAGGAA CTTCAAATCT GGGAGATGAG GAAAGAAAAC TACTCACAA ACAACATAAA TGTAATAAAT TCAAAACCCT AAAGGAGAGC TGTCCCTAGA CAGTAGTGGC TACAGGTGCT AAGCAAAAGT CAGGTACACT GGACAGAGCC ATGCGATTGA TTGTTTCTCT TCCCTCTCTG GGTCTCCAGA AAAGGACACT GGCAATATCC CCAGCTCTCT CCTGATTGGA ATTCTTTGAA CCCTCGAAGT GCTCCAGCAG TACTACCCCC CACGTAGCAG TGGGCCCATG TCGCTTGAAT ATTATTTTTG ATTTGACCAC CAAGATGTAA TGATGATTCT ATTCCATTTT GAAAAGGGTC TGAGAAAGTG TACAGGTCTA ATTATATATA CATATTTATA CATGTGTATT TTTGTTTATT GTTACATTTT GACATTGGAC TTTCTATTAA ATAATTTTAA AGAGTTGG</p>	
<p>CFLAR</p>	<p>NM_0 011271 83.2</p>	<p>GACCACGCAT GGAGAATTTT ACCACCCAGA GACACGCGAG TGGCCCTGTG CAAGTTTCAA CTGCGCGGGG GGCGGGGAAT TCCGCAGACA GGATTCTAGA ATTTATGTCT TGTGTGGTAA CATTTACAGCC GGTGGGTGGC GGGGATTAGG CGTGAAGCGG TTCAGCAGGC AGAGGTCTC GGACGCCCTC CGGCGAAGCC ACCTGTTGAT GCTTTTIGACT TTCTGTCCCT GTTCCTCGTC CCATCTGGAG CATTTCCAAT TCTGGTTTTG CGGAGCAGCA GGTCTGAGCT TGTCGGCGA GGGTGGGAGT TGGTCCCGGC GGAGATCCAG TGGGAAGAGC CGGCGGCTGC CCGGGCAACT CCCCACTGG AAAGGATTCT GAAAGAAATG AAGTCAGCCC TCAGAAATGA AGTTGACTGC CTGCTGGCTT TCTGTTGACT GGCCCGGAGC TGTACTGCAA GACCCTTGTG AGCTTCCCTA GTCTAAGAGT AGGATGTCTG CTGAAGTCAT CCATCAGGTT GAAGAAGCAC TTGATACAGA TGAGAAGGAG ATGCTGCTCT TTTTGTGCCG GGATGTTGCT ATAGATGTGG TTCCACCTAA TGTCAGGGAC CTTCTGGATA TTTTACGGGA AAGAGGTAAG CTGTCTGTCG GGGACTTGGC TGAAGTCTC TACAGAGTGA GGCGATTTGA CCTGCTCAA CGTATCTTGA AGATGGACAG AAAAGCTGTG GAGACCACC TGCTCAGGAA CCCTCACCTT GTTTCGGACT ATAGAGTGCT GATGGCAGAG ATTTGGTGAAG ATTTGGATAA ATCTGATGTG TCCTCATTAA TTTTCCTCAT GAAGGATTAC ATGGGCCGAG GCAAGATAAG CAAGGAGAAG</p>	<p>4</p>

	<p>AGTTTCTTGG ACCTTGTGGT TGAGTTGGAG AAACATAATC TGGTTGCCCC AGATCAACTG GATTTATTAG AAAAATGCCT AAAGAACATC CACAGAATAG ACCTGAAGAC AAAAATCCAG AAGTACAAGC AGTCTGTTCA AGGAGCAGGG ACAAGTTACA GGAATGTTCT CCAAGCAGCA ATCCAAAAGA GTCTCAAGGA TCCTTCAAAT AACTTCAGGC TCCATAATGG GAGAAGTAAA GAACAAAGAC TTAAGGAACA GCTTGGCGCT CAACAAGAAC CAGTGAAGAA ATCCATTCAG GAATCAGAAG CTTTTTTGCC TCAGAGCATA CCTGAAGAGA GATACAAGAT GAAGAGCAAG CCCCTAGGAA TCTGCCTGAT AATCGATTGC ATTGGCAATG AGACAGAGCT TCTTCGAGAC ACCTTCACTT CCCTGGGCTA TGAAGTCCAG AAATTCTTGC ATCTCAGTAT GCATGGTATA TCCCAGATTG TTGGCCAATT TGCCTGTATG CCCGAGCACC GAGACTACGA CAGCTTTGTG TGTGTCCTGG TGAGCCGAGG AGGCTCCCAG AGTGTGTATG GTGTGGATCA GACTCACTCA GGGCTCCCC TGCATCACAT CAGGAGGATG TTCATGGGAG ATTCATGCC TTATCTAGCA GGAAGCCAA AGATGTTTTT TATTCAGAAC TATGTGGTGT CAGAGGGCCA GCTGGAGGAC AGCAGCCTCT TGGAGGTGGA TGGGCCAGCG ATGAAGAATG TGGAATTCAA GGCTCAGAAG CGAGGGCTGT GCACAGTTCA CCGAGAAGCT GACTTCTTCT GGAGCCTGTG TACTGCGGAC ATGTCCCTGC TGGAGCAGTC TCACAGCTCA CCATCCCTGT ACCTGCAGTG CCTCTCCCAG AAACCTGAGAC AAGAAAGAAA ACGCCCCTC CTGGATCTTC ACATTGAACT CAATGGCTAC ATGTATGATT GGAACAGCAG AGTTTCTGCC AAGGAGAAAT ATTATGTCTG GCTGCAGCAC ACTCTGAGAA AGAACTTAT CCTCTCCTAC ACATAAGAAA CCAAAGGCT GGGCGTAGTG GCTCACACCT GTAATCCCAG CACTTTGGGA GGCCAAGGAG GGCAGATCAC TTCAGGTCAG GAGTTCGAGA CCAGCCTGGC CAACATGGTA AACGCTGTCC CTAGTAAAAA TACAAAAATT AGCTGGGTGT GGGTGTGGGT ACCTGTATTG CCAGTTACTT GGGAGGCTGA GGTGGGAGGA TCTTTTGAAC CCAGGAGTTC AGGGTCATAG CATGCTGTGA TTGTGCCTAC GAATAGCCAC TGCATACCAA CCTGGGCAAT ATAGCAAGAT CCCATCTCTT TAAAAA AAAAAGGAC AGGAACTATC TTAATCAATG TATTAGTCAT GTTTCTCTAG AGGGACAGAA CTAATAGGAT ACATGTATAT AAAAAGGGGA GTTTATTAAG GAGTATTGAC TCACATGATC ACAGGGTTAG GTCCACAAT AGGTCATCTG CAAGCAAGGA AGCCAATTCA AGTCCCAAAG CTGAAGAACT TGGAGTCAA TGTTTGAGGG CAGGAAGCAT TCAGCATGAG AGAAAGATGG AGGCCAGAAG ACTACACCAG TCTAGTCTTT CCATGTTTTG CCTGCTTTTA TTCTGGCAGT GCTGGCAGCT</p>	
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	<p>GATTAGATGG TGCCACCCCA GATTGAGGAT GGTCTGCCTT TCCCAGTCCA CTGACTCAAA TGTTAAATCT CCTTTGGCAG CACCTCACA GATGTACCCG GGAACACTTT GCATCCTTCT ATTCAATCAA GTTGATACTC AGTATTAACC ATCACAGTCC ATTTGGGCAA CTATACCAA TTACCATAGA CCAGGTGACT TAAACAGCAG TTATTTCTCA CAGTTCCGGA GGCTGGGAAA TCCAACATCT AAGTGGTAGC ATATCTGGTG TCTGGTAAGG CATGCTTCCA GATCTTACCA GATGTCAGTC TTTTGATGTT CTCACATGGC AGAAAAAGAG GATGCAAACCT CTCAAGTATA TCTTTAAGGG CACAAATTC ATTCATGAGG GCTCTACCCT CATCACCTAA TTACCTCCCA AAGGCCCCAC CTTCTGATAC TGTCACTTTG GGGATACTGT CTCCCCTTTG AATTCTGGGG GGAATACAAA CATTGAGTTT GTAACAATAG CCTTATGATT TAGAGGTTAC TTGTTTATT ACCTAGACCT CAAATTGCAT TTTACAGCTA GTCAAGTATA TCTTTCTCTG ATTTGATAGT GTGACCTAAA AGGGGACCAT TGTTTGAAAT ATCATTAGAG TTGCTTATTA TTATTATTAT TATTATTATT ATTATTATTA TTATTATTAT TGAGACAGAG TTTCATTCTG CTGCCCAGGC TGGAGTGCAG TGGCATCATC TTGGCTCATT GCAACCTCTG CCTTCTGGGT TCAAGCGATT CTCCTGCCTC AGCCTCCCGA GTAGCTGGGA TTACAGGCTC CTGCCACCAC ACCCGGCTAA TTTTTGATTT TTTAGTGGAG ACAGGGTTTC CACCATGTTG GCCAGCGTGG TCTTGAACCT CTGACCTCAG GTGATTCACC AGCCTCGGCC TCCCAAAGTG CTGGGATTAC AGGTGTGAGC CACTGCACCT GGCCTATTAT TATTTTTAAA TTTTTTTTTT TTAATTGATC ATTCTTGGGT GTTCTCACA GAGGGTGATT TGGCAGGGTC ACAGGACAAT AGTGGAGGGA AGGTCAGCAG ATAAACAAGT GAACAAAGGT CTCTGGTTTT CCTAGGCAGA GGACCCTGCG GCCTTCCGCA GTGTTTGTGT CCCTGGGTAC TTGAGATTAG GGAGTGGTGA TGACTCTTAA GGAGCATGCT GCCTTCAAGC ATCTGTTTAA CAAAGCACAT CTTGCACTGC CCTTAATCCA TTTAACCTG AGTGGACACA GCACATGTTT CAGAGAGCAC AGGGTTGGGG GTAAGGTCAT AGATCAACAG CATCCTAAGG CAGAAGAATT TTTCTTAGTA CAGAACAAA TGAAGTCTCC CATGTCTACT TCTTTCTACA CAGACACAGC AACAATCTGA TTTCTCTATC TTTTCCCCAC CTTTCCCCCT TTTCTATTCC ACAAACCGC CATCGTCATC ATGGCCTGTT CTCAATGAGC TGTTGGGTAC ACCTCCCAGA CGGGGTGGCG GCTGGGCAGA GGGGCTCCTC ACTTCCCAGA TGGGGCGGCC AGGCGGACGC GCCCCCCACC TCCCTCCCGG ACGGGATAGC TGCCCGGGCG GGGGCTGACC CCCCACCTCC CTCCCCGACG GGGCGGCTGG CCGGGCGGGG GCTGACCCCC ACGCCTCCCT</p>	
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	<p> CCCGGACGGG GCGGCTGCCA GGCGGAGGGG CTCCTCACTT CTCAGACGGG GTGGCTGCTG GGCGGAGACG CTCCTCACTT CCCAGACAGG GTGGCTGTGCG GGCGGAGGGG CTCCTCACTT CTCAGACGGG GCAGCTGCGG GCGGAGGGGC TCCTCACTTC TCAGACGGGG TGGCCGGGCA GAGAAGCTCC TCACATCCCA GACGGGGGGG CGGGGCAGAG GCGCTCCCCA CATCTCAGAC GATGGGCGGC CGGGCAGAGA CGCTCCTCAC TTCATCCCAG ACGGGGTGGC GGCCGGGCAG AAGCTGTAAT CTCGGCACCC TGGGGGGCCA AGGCAGGCGG CTGGGAGGGCG GAGGCCGTAG CCAGCTGAGA TCACACCACT GCACTCCAGC CTGGGCAACA TTGAGCACTG AGTGGACGAG ACTCTGCCCC CAATCCCGGC ACCTCGGGAG GCCGAGGCTG GCAGATCACT CGCAGTCAGG AGCTGGAGAC CAGCCCGGCC AACACAGTGA AACCCGTGCT CCACCAAAA AATACGAAA CCAGTCAGGC GTGGCGGGCGC CCGCAATGGC AGGCACGCGG CAGGCCGAGG CGGGAGAATC AGGCAGGGAG GCTGCAGTGA GCCGAGATGG CAGCAGTACA GTCCAGCTTC GGCTCGGCAT CAGAGGGAGA CCGTGGGGAG AGGGAGAAGA GAGGGAGGGG GAGAGGGCTA TTTTAAAAT TTTTAAAAT TGCTGAACAG GGGTACCTCT GGGCAGTGTG TCAGAATACC ACTTTTTAAA TATTTTATGA TTTATTTATT TTTCTATTTT TTGAGGTTTT AACTGATGTG TATCTGTATG TCTATTTGTG TATATTTTGT CATGATCATG TAACAGAGTC TGAAAAGTGT CGAAGAGACA GTTTTTCAGGA ACAACAAGCA ATTATTCCTA CTTTCCAAGT TATTTTGATG CCATGGTGGC TCATACCTAT AATCTGAGTA CTTTGGGAGG CTGAGGTGGA CTGATCACTT GAGCCCAGGA GTTTGAGACC AGCCTGGGCA ACATAGCAAG ACTCCATCTC TACAAAAAAA GACAAAATTT AGCTGAGCGT GGTGGCGTGT TCCTGTAGTC CCAGCTACTT GGGAGGCTGA AGTGAGTGGA TCCCCTGAGC CCAGAGAGGT CAAGGTTGTG ATGAGCTGTG ATCACACCAC TGCAC TTCAG CATGGGAGAC AGAGTGAGAC CCTGTTTTAG AAAAAATAAA TAAATAAAAC CACCAGCACC ACAAACAACA AAAAAAGTT ATTTTGTACT TGTTTTGAGC ACAGGACTCC TGAGGGTATC TTTGCATTTA ATATTACATA GGGGTGCCAG TGGGAAGTAA TGTGTATGCT TGGCCTCATG AGCTAAAACC CTGTGTTAAT TATGACAGAA GGAAAGTGTG TGAGAGAGAT CTAACTACC TAGCAGCTCT AGCTGCCATC TTGAACCATG AAGATACGGG CCACACGTAG GGGTAGCTGG GTAGTGAGCA GCAAGAAGCC TTGTTGGATG AGGGCACGAA GGAGCAGAAT CACTGGAATC ACTGTGTCAG CCCTAATTAC CTACCTCTGG ACTTTTATGT GAGGGGAAAA AAAATTGACA GTTTATATTT ATCTCAACCT AGTTAACCCA AGTGATGCAT TGTTATGAGA TTTAAAATGTT </p>	
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	<p>TGGAGGCCGG GTGCGGTGGC TCACGCCTAT AATCCCAGCC CTTTGGGAGG CCAAGGCGGG CGGATCACGA GGTCAGGAGA TCAAGACCAT CCTGGCTAAC ATGTAAAACC CCGTCTCTAC TAAAAATACA AAAAATTAGC CAGGCGTTGT GGCGGTCCGC TGTAGTCCCT GCTATTTGGG AGGCCGAGGC AAGAGAACGG CATGAACCTG GGAGGTGGAG CTGTCAGCGA GCTGAGATCT TGCCACTGCA CTCCAGCCTG GGCACAGTG CGAGACTCTG TCTCAAAAAT AAATAAATAA ATAAATAATA AATAAAATGT TTGGAATGTT GGCTTCATCC CTGGGATGCA AGGCTGGTTC AACATACGCA AATCAAGAAA CATAATTCAT CACATAAACA GAACTAAAGA CAAAACCAC ATGATTATCT CAATAGATAC AGAAAAGGCC TTCAATAAAA TTCAACGTTG CTTCATGTTA AAAACTCTCA ATAACTAGG TATTGATGGA AAATATCTCA AAATAATAAC CATTTATGAC AAACCCACAG CCATTATCAT ACTGAATGGG CAAAAGCTGG AAGCATTCCC CTTGAAAAC GGCACAAGAC AGGGATGCCG TCTCACCCT CCTATTTAAC ATAGTATTGG AAGTTCTGGC CAAGAAAATC AGGCAAGAGA AACAAATAAG GGGTATTCAA ATAGGAAAAG AGGAAGTAAA ACTGTGTTTG CAGATGACAT GATACTATAT CTAGAAAACC CCATTATCTC CACCCAAAAG TTCCTTAAGC TGATAAGCAA CTTCAGCAAA GTCTCAGGAT ACAAATCAA TGTGCAGAAA TCACAAGCAT TCTATACACC AACAATACAC AAGCAGAGAG CCAAATCATG AATGAACTCC CATTACAGT TGCTAGAAAAG AGAATAAAAT ACCTAGGAAT ACAGCTAATA AGATGTGAAG GATCTCTTCA AGGAGAACTA CAAACCACTG CTCAAGGAAA TAAGAGAGGA CACAAATGAA AAAACATTCC ATTCTCGTGG ATAGGAAGAA TCAATATCAT GAAAATGGCC AACTACCCA AAGTAATTTA TAGGTTCAAT GCTATTCCA TTAACTACT ATTGACATTC TTCACAGAAT TAGAAAAAAA CTACTTTAAA ATTCAAATGG AACCAAAAAA GAGCCCGTAT AACCAAGACA ACAATAAGCA AAAAGAACAA AGCTGGAAGC ATCACACTAC CCAACTTCAA AGTATACTGC AAGGCTACAG TAGCCAAAAT GGCATGGTAC TGGTACAAAA ACAGACACAT AGACCAATGG AACAGAATAG AGACCAGAGA AAGAAGACCA CACATCTACA GCCATCTGAT CATCGACAAA CCTGACAAAA ACAAGCAATG GGGAAAAGAT TCCCTATTTA ATAAATGGTG CTGGGAAAAC TGGCTAGCCA TATGCAGAAA ATTGAAACTG ACCCCTTCT TACACCTTAT AAAAAATTA ACTCAAGATT AAAGACTTAA TGTA AACCT AAAACTATAA AAACCCTAGA AGAAAATCTA TTTAATAACA TTCAAGACAT AGGCACAAGC AAAGGTTTCA TGACAAAAAC ATCAAAAGCA ATTGCAACAA AAGCAAAAAT TACAAATGGG ATCTAATTA ACTAAAGAGC TCCTGCACAG</p>	
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	<p> CAAAAGAAAC TATCATTAGA GTGAACAGGC AACCTACAGA ATGGGAGAAC ATTTTTGCAA TCTATCCATC TGACAAAGGT CTAATATCCA GAACCTACAA GGAACCTAAA ACAAATTTAC AAGGAAAAAA ACAACCCCAT CAAAAAGTGG ACAAAGGACA TGAACAGACA CTTCTCAAAA GAAGACATTT ATGTGGCCAA CAAACATATA AAAAAAAGCT CAACCTTACT GATCATTAGA GAAATGCAA GGAGAACCAC AATGAGATAC CATCTCATGC CGGTCAGAAT GGTGATTATT AAAAAGTCAA AAAACAACAG ATGCTGGCGA GGCTGTGGAG AAGTAGGAAC ACTTTTACAT TGTTGGTGGG AATGTAAATT AGTTCAACCG TTGTGGAAGT GTGTGTGGCT ATTCCTCAA GATCTAGAAC TAGAAATACT ATTTGTCCCA GCAATCCCAT TACTGGGTAT ATACCCAAAG GAATATAAAC CATTTTATTA TAAAGATACA TGCACATTTT TGTTTATTGC AGCACTCTT ACAATAGCAA AGACACAATA GCAAATGCCC ATCAAAGATA GACTGGATAA AGAAAATGTG GTACATATAC ACCATGGAAT ACTGTGCAGT GCAGCCATTA CAGCTTTTGG TGATACAGTG AATCAGATTT TICATTAATT CTTTTAATTG GTTATTACTG AACGTGAAAA AGTAATGTTT GTATTGAAAT CTTGAGTCTG GCCATGTTTC TATTTTAAAT TCATAAAGAA TTCTAACAAG AGGAATTCCA AGAATGTCAT AAATGGATGT TTCTCCATGG ATGAAGGAAC TGTTTTATTC ACTTGCTGAT AATTCAGCCT AATCCAGTTT GACATCATAT AGATAAGTAG TTGAATTATG GATTTAAAAT ACATATCATT TTCTAACTCC AAAGGTAATA CTTATTTAAA TGGTTTTGAA AATATAGAAA GGCACAATTT CTTTTTAAAT CTGTTATTCT CCACCACCAC TCAATCTGTC TATCATCTAT CTCTCCATTC ATTCTTCCAT TTGTTTATAT CTGTTAATCT TTGTATGTGT TCATGTATAG CTTTTACATG ATTGGAATCA TAATGCATAT TCCATTTTGA AGTCTGCTTT TTTTACACA AAAATATGTT GTGAATATTT TCCTATATTA TGAAATATCA TTAGCTGAGC TTTTAGAATT GACTGCATGT TTTGGTACCA TTTAGATATA GTTTAAGATA CTTAGAAGTT ATGTGGCTTT GCCACTATGG ATGAATCTTA TTTACTCAAT ATTAATTAAT TACAAATAAC CTCACCTAAA CACTACTCAG CCATAAAAAG GAATGAATTA ATGACATTCA CAGCAACCTG GAGACTATTA CTCTAAAGGA AGTAACTGAG GAATGGAAAA CCAAACATTG TATGTTCTCA CTCATAAGTG GGAGATAAGC TATGAGGATG CAAAGGCATA AGAAGGATAC AATGGACTTT GGGGACTTAG GGGAAAGGGT GGGAGGGGGG TGAAGGATAA AAGAATACAA ATTGGGTTCA GTGTATACTG CTCAGGTGAT GGGTGCACCA GAATCTCACA AGTAACCACT TAATTACTTA CGCATGTAAC CAGATACCAC CTGTTCCCA AACACCTATG GAAATAAATT TGTTTTTTTT </p>	
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		<p>TTTAAAAAAG GAATGAGATC ATGTCCTTTG CAGGGACATG GATGAAGCTG GAAGCCATTA TCCTCAGCAA ACTAACAGAG GAGCAGGAAA CCAAACACCA CATGTTCTCA CTTGTAAGCG GAAGCTGAAC AATGAGAACA CACGGACACA GGGATGAGAT CAACACACAC TGGGGCCTGA TGCAGGGGCC GTAGCGGGGA GAGCATCAGG ATA ACTAGCT AATGCATGTG GGGCTTAATA CCTAGGTGAT AGGTTGATAG GTGCAGCAA CCACCATGGG ACACGTTTAC CTATGTAACA AACCCGCACA TCCTGCACTT GTATCCAGAA CTTAAAATAT TTTAAAAATC TTTAGAGAAT ACAAAAA AAAAAGAT TCTTCAATGC ATACACAATA AAATTGCAGT TCAGTCAAAC ATTGGAAGTC TTTCTCTGAC TGTCTAGTTG GTATCTTCAT TTTCAGCTTC TTCAAGATCC CACTCCAAAC ACTGTTAGCT CAGCCAAATT GAACAGCTCA TATCTCCTAC CTCTGGATCT TTGGTTCTGG TGATTGTATA TTTCTGGACC ATCTGGAACC CCAGCATATC ACCCTACCCC ACATCTCCAC ATCCCCAAAA TATAACCATA CTTCAAGGGC AGTTCAAATA CCATCTCCTT CTATCCTCCA TGAAGTCAGT TATCTCTTCC ATTGGAATTA TCGCCCCCTC TCCTGAACAG TACTATTTTCG TGTGAATCTC CTCCAAGCCT TCTTTTCATT TTATATCTCA TGCTGTAATT CTTGGAAAGT ATGCTGTAGC TCAAGTGCAG AATTCTCATC AGTTTTATCT TTATATCTCT CCTAAACACT TTACCTGATG AAGAGCCTGG CATAACATA AATATATATT GAATGAATCA GTGATGGATT GAAAAGAGAA ATGATGGATC TCCTAAATTT TAACTTTTAT AAAATATTTT GATACATTCA TGACCTTACT TTAGCAAGCA ATGAACGTGA TGTAACACTAT TGTTGATATA GTTTTTATAT TGGAAGTGTA AGTAGTTTGT GGCATGGGAT TGTGACATAT CCTAGGTTTC CTCATCTTCT TTTTATTGAA ATGTAATTCA CAAGCCATAA AATTTGCCCC TTTAAAGTAA ATGATGCAGT GGATTTTAGT ATATTTACAG AGTTGTGCAA TCATCACCAC TATCTAATTC CAGAACATTT CCATCTACCT AGAAACTCCA TACCAGTGAG CTGCCACTCT AATCCTCCTC TTCCCCCAGC CTCTAGAAAC AATAATCCAT TTTCTGTCTC TATGATTTGC CTGTTCTAGA TATTTTATAA AAATAACAT GTGGCCTTTC GTGTCTGACT TCCTTCACTT AAAAAAAAAA AAAAAAAAAA</p>	
<p>CFLAR -ASI</p>	<p>NR_04 0030.1</p>	<p>CATGCATACA GCCCAGTCTA ACGTACAGGA TCAAAAAGAG ACCTTATTTT GGCTGGCAAT TACCTTACCC GCTTGTTTAT TATCTGACTC CTGCACTCCT CCCAAGTGTT CAATGTAAAC CAGATGGCAA AGTTAACCAG ACAGAGGAAA TCCACATAAT TACTCTAGAT GAAGCCCCCT AACTGCGGAT TAGAAGGAAA AGAGATGAAG GGTGCTGCAC TAATAATAAC AACCATTTTC</p>	<p>5</p>

		<p>AACCAAGGCA GGATTCTGGG CCTAGAATCT TCAGTGACAA ATTACCAGGA GAGGAGGAGG GAAATGAAGA GCTGGAGGCC TAGGGAGATG GTGTGAGGAG AGGAGACTGT GCCTTTCCAC TTCTTAATAA TTAATGTTCT CCTGCTGCTC TGAGTCTGCA TCACCAGTGA TTCAGTTTGA GTCAGACATC CAACAACAAC TGAGCAAAAG AATCATCTCA GATCCAAGCA ATGGTGACGC GCAGATAGAA GAGACTGAGG CCAACGACCA AAGCAAAGGA CAGCAGAACT GACTGACACA GCTCAGAAAA TATCCAAAGC TCTGTATGCT CTGGAAAGGA AGACAAGAAA TGCTAAAGAC TCTGGATATT GCTCTGTTT TATAAGGTCC CACCCATATA CCATCTCTTC TTTTGGATTT CCTTTGGCAG GGTTCCCTTG TATGGAAGTT CTCAGGAGAA CAACACTCTC AGAGTCCCAC TGCTCAAAG AATTCCTAAT GGACAGCTAT ACTTGGCCCC AAGGTTTCAA CCTCACACTG AAGAAAGTGA CATAAGAAAA CACAAGTCCC AGTGCTATCA GGGGCTATCT GGAAAAGCTG CTTAAACAGC AAGTGAAAGA AGGCTCTCCC CTACCCCGTT CTAGACTCAC CCTGAAGTTA TTTGAAGGAT CCTTGAGACT CTTTTGGATT GCTGCTTGGGA GAACATTCCT GTAACCTGTC CCTGCTCCTT GAACTATAAA AGCAGAAGGG AGGTACTGAG TTAAAACCTA CAAATTATTA TCAAATAAAT CTTCAATAGT G</p>	
<p>CHP1</p>	<p>NM_0 07236. 4</p>	<p>ACCACCCCTG GGTTCCCTCC CGGGTCCGCA GTGGAAACAC TGCCCTCTCC CTTCCTGACC CCTAGCCCTT CCTTCCCTCC CTCCTTCCCT CCTGTCGCCG TCTCTTCTGG CGCCGCTGCT CCCGGAGGAG CTCCCGGCAC GGCGATGGGT TCTCGGGCCT CCACGTTACT GCGGGACGAA GAGCTCGAGG AGATCAAGAA GGAGACCGGC TTTTCCACA GTCAAATCAC TCGCCTCTAC AGCCGGTTCA CCAGCCTGGA CAAAGGAGAG AATGGGACTC TCAGCCGGGA AGATTTCCAG AGGATTCCAG AACTTGCCAT CAACCCACTG GGGGACCGGA TCATCAATGC CTTCCTTCCA GAGGGAGAGG ACCAGGTAAG CTTCCGTGGA TTCATGCGAA CTTTGGCTCA TTTCCGCCCC ATTGAGGATA ATGAAAAGAG CAAAGATGTG AATGGACCCG AACCCTCAA CAGCCGAAGC AACAAACTGC ACTTTGCTTT TCGACTATAT GATTTGGATA AAGATGAAAA GATCTCCCGT GATGAGCTGT TACAGGTGCT ACGCATGATG GTCGGAGTAA ATATCTCAGA TGAGCAGCTG GGCAGCATCG CAGACAGGAC CATTGAGGAG GCTGATCAGG ATGGGGACAG TGCCATATCT TTCACAGAAT TTGTTAAGGT TTTGGAGAAG GTGGATGTAG AACAGAAAAT GAGCATCCGA TTTCTTCACT AAAGGAGACC AACTGTTCC TTGCGGTCTA GTATTTAAGA ACTGGAACCTT GAAAGTCCTC CTCTACCAA CTCCACCTCC ACCCCCTCAT TCCCCTTCTC CCAAAGTACT</p>	<p>6</p>

	<p>ACTGCTGTTG CATGACAACC CCAAATATGT TCTGTCAACA CAAACCTGCC TTTGGTGTAT AAACAGGGCA TTACAGAATG GTACACCCTA TATATTTCTG TTCAGTATCC ATTCACTAGT TCTTCATTTA TAAATATCAT CTTCCCCATT CTGCTGCTGA ATGCCACACA TCCATCCAGT CTGAGAAAAGT GAGAGAGGCA ATCATGCCAA GAACAAGCCA GCAAAGCTCT TTCACCAGAT GTAGACTGTA GCCCTGCTGC CTTCCCTCCA GCGAGTCTGC CAGCATGCTT CTTCATCCTT TTTATATGTT CTTTGCTTCC TACTTCCCTG TCTTCCAACA TACTGTTTAC TTA CTCTGCGC AGTCTTTCTG CTTTTTATTA AGCCTCAAAA TCTCCTCTGT TCTACTTGGC ACCACAAGCT ATGTCCTATA TATGTATTTT TGACTTGGCA GGATAGTTCA GGGGTCTGGC AGTTTTTATT TACCTTCATT ATFAAATGGG CCTCTGGGAT GTTGCCCTCT CAGGAGCTTT TTGGTAATCA A TACTTCTCT CAGAAGTATG AGACCATCCT CTGCACTCTG CTCTGTCATC AAAGGCTGCT GGGTGGAGAT ACCCTTTTTG AAAGGTGGCC TTGGTGAGAG GTATGGAGCC AAGTCTTCTA GGTGCTTGC CCACATCACT CTATCTCTGG CCTCTGATTC TCAACTTTGT ACCTGTGTGG CTCCTCTTGT TAGTGCAATG TTGACTGTTG AAAAAGCAGC AGTATGCTTA CAGGTTTGCT TAGTTTGGGG ACACCGTTAC CACCAGAATG GCTGCTCTGA CAATATGCCT AGGGACTTTC TCATGGCTTT TATTTAATAA GGAGGCTGGG CACCCTATAA AGCCTCATGC ATTCACACCT TTGCAGCATG GTTTATGCCT CAGTGTTATG TGCCTGGAA TGTTTTCCAC TTCACATTTT CAAGTAGAAA TATTAGTGTT ACGGAAGTGC CTAATATCCC AGTCCAAATT TTTTTTTTTT TTTTTTTTTT TTTTGTAGAC AGAGTCTTGC TCTGTCACCC AGGCTGGAGT GCAGTGGTGC GATCGCTCAC TGCAACCTCA GCCTCCTGGA TTTAAGTGAT TCTCCTGCCT CAGCCTCCCA AGTAGCTGGG ATTACAGGTG TGCACCACCA TGCCCGGCTA ATTTTTTGTA TTTTATAGTG AGACAGGGTT TCACCATGTT GGCCAGGCTG GTCTCGAACT CCTGACCTCG TGATCCGCCT GCCTCAGCCT CCCAAAGTGC TGGGATTACA GGTGTGAGCC ACCACGCCTG GCCCAGTCC AAAATATTTA AAGATTGTTT CTTAGTGTG TTGAAGTTTT GCACAAAATT CTTTTTTTTG AGATGGAGTC TCACTCTGTC ACCCAGGCTG GAGTGCAGTG GCGTGATCTT GGCTCACTGC AACCTCTGCC TCCTGGGTTT AAGCAATTCT CCCACCTCAG CCTCCCAAGT AGCTGGGATT ACAGACGTGT GCCACCATAC CTGGGTAATT TTTGCATTTT TAGTGGAGAG GGAGTTTAC CATGTTGGCC AGGTGGTCT TGAACCTCTG ACCTCAGGTG ATCTCCTGC CTCGGCCTCC CAAAGTGCTG GGATTACAGG CATGAGCCAC CGTGCTCAGC CGCAAATTC TTTATGAATT</p>	
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		<p>TTACACTTGG CAAATGTTAA TGACGGAAGC CATAGTCTGC TCCTAATACA TGTCCAAAGC ATTGACTGTT GTGTCATTAG CTGCCTGGTT ACATTAGCTC CCTGGCTTCT TGTTTAGACC ACTGCTAATC CCTTAAAAAC AAGAGGTCTG GCACTAGTAG CACAACCTAA GGTGGCATT A CAGATCTTTG AGCGAGCCAC AGCAACTTTT CTGCCAAGTC AGCTTAGTTT AGACTTCAGT GAATCAGGCT ATTGCTATCC TAATGTATGT CTCTATGAGT GTATTTAGCC ACACATCTGC CCTTGGTTGA CTTTCTGACT CATTGCTTGC TTGCTTGTTC CCTTGGCTTTG GAAAACTATT GAAGATTGCT AAAAAATACC ACTGCAAAGT GATGGAAAAG GGTGGAGAAC AGGGGAGTAG CCAGGCTGGA TGGCTCAAAT ATAAATGAAT GAGGAATTCT TTATGAAGTA TCAGTCAGAT TTTATGATTA AGTGATGTAA TATAGGAATT ATGTA AAAAGG GAAGAATGTC TGATACTGAT CTATTAGAGA GGTACTTTAG AGGCTTCTTG ATTGGCATAA AGTTCCTAAG GTTATAGATT TTCCCCCTT TTGGCTGTAT AGCAAAGTGT TTTAATCCAC GGTTGTGCCT TATTGTTCCA TTA AAATTGT ATCTTCGATC CATCAATAAA TACTTGTGGT TGA AACAAAA</p>	
<p>DDX55</p>	<p>NM_0 20936. 2</p>	<p>AGCGGAAGTG CTCGTTGGGG GTGCACAAGG CGCGTTCGAG CAGCGGCGAC CGACGCGGCG AAGGAGCGCG CCATGGAGCA TGTGACAGAG GGCTCCTGGG AGTCGCTGCC TGTGCCGCTG CACCCGCAGG TGCTGGGCGC GCTGCGGGAG CTGGGCTTCC CGTACATGAC GCCGGTGCAG TCCGCAACCA TCCCTCTGTT CATGCGAAAC AAAGATGTGC CTGCAGAAGC GGTCACAGGT AGTGGCAAAA CACTCGCTTT TGTCATCCCC ATCCTGGAAA TTCTTCTGAG AAGAGAAGAG AAGTTAAAAA AGAGTCAGGT TGGAGCCATA ATCATCACCC CCACTCGAGA GCTGGCCATT CAAATAGACG AGGTCTGTGC GCATTTACAG AAGCACTTCC CCGAGTTCAG CCAGATTCTT TGGATCGGAG GCAGGAATCC TGGAGAAGAT GTTGAGAGGT TTAAGCAACA AGGTGGGAAC ATCATTGTGG CCACTCCAGG CCGCTTGGAG GACATGTTCC GGAGGAAGGC CGAAGGCTTG GATCTGGCCA GCTGTGTGCG ATCCCTGGAT GTCTGGTGT TGGATGAGGC AGACAGACTT CTGGACATGG GGTITGAGGC AAGCATCAAC ACCATTCTGG AGTTTTTGCC AAAGCAGAGG AGAACAGGCC TTTTCTCTGC CACTCAGACG CAGGAAGTGG AGAACCTGGT GAGAGCGGGC CTCCGGAACC CTGTCCGGGT CTCAGTGAAG GAGAAGGGCG TGGCAGCCAG CAGTGCCAG AAGACCCCT CCCGCTGGA AACTACTAC ATGGTATGCA AGGCAGATGA GAAATTTAAT CAGCTGGTCC ATTTTCTTCG CAATCATAAG CAGGAGAAAC ACCTGGTCTT CTTCAGCACC TGTGCCTGTG TGGAATACTA</p>	<p>7</p>

	<p>TGGGAAGGCT CTGGAAGTGC TGGTGAAGGG CGTGAAGATT ATGTGCATTC ACGGAAAGAT GAAATATAAA CGCAATAAGA TCTTCATGGA GTTCCGCAA TTGCAAAGTG GGATITTAGT GTGCACTGAT GTGATGGCCC GGGGAATTGA TATTCCTGAA GTCAACTGGG TTTTGCAGTA TGACCCTCCC AGCAATGCAA GTGCCTTCGT GCATCGCTGC GGTGCGCACAG CTCGCATTGG CCACGGGGGC AGCGCTCTGG TGTTCTCCT GCCCATGGAA GAGTCATACA TCAATTTCTT TGCAATTAAC CAAAAATGCC CCCTGCAGGA GATGAAGCCC CAGAGAAACA CAGCGGACCT TCTGCCAAAA CTCAAGTCCA TGGCCCTGGC TGACAGAGCT GTGTTTGAAA AGGGCATGAA AGCTTTTGTG TCATATGTCC AAGCTTATGC AAAGCATGAA TGCAACCTGA TTTTCAGATT AAAGGATCTT GATTTTGCCA GCCTTGCTCG AGGTTTIGCC CTGCTGAGGA TGCCCAAGAT GCCAGAATTG AGAGGAAAGC AGTTTCCAGA TTTTGTGCC GTGGACGTTA ATACCGACAC GATTCCATTT AAAGATAAAA TCAGAGAAAA GCAGAGGCAG AAACTCCTGG AGCAACAAAG AAGAGAGAAA ACAGAAAATG AAGGGAGAAG AAAATTCATA AAAAATAAAG CTTGGTCAA GCAGAAGGCC AAAAAAGAAA AGAAGAAAAA AATGAATGAG AAAAGGAAAA GGGGAAGAGGG TTCTGATATT GAAGATGAGG ACATGGAAGA ACTTCTTAAT GACACAAGAC TCTTGAAAA ACTTAAGAAA GGCAAATAA CTGAAGAAGA ATTTGAGAAG GGCTTGTTGA CAACTGGCAA AAGAACAATC AAGACAGTGG ATTTAGGGAT CTCAGATTIG GAAGATGACT GCTGATTCCA GTGCCACAGA TGAACCCACA AGGACATAGC TGTTCCCTAA CTTGGTGGAT GGCTCCAGTT TGCTTTTAAAC GAAAATCACA ACTTCAGGAG ACATCTGAAA AGAATGATGT CTCTGAAAGC TGTCCTTTCA GATGAGGGAG AAATGAAGGA TTTACACTT CAGAATATTT TACTAAAAAC ATTCCAGTCT TGGCCGGGTG CGGTGGCTCC TGCCATAAT CCCAGCACTT TGGGAGGCTG AGGCAGGAGG ATCACTTGAG CCCAGGAGTT CAAGACCAGC CTGGGAACAC AGCGAGACCC TCTCATTAAC AACAAACAAA CAAAACAATT CCAGTCTTGG AGTAGTCTAA CAGAAGAAAA TGTAATAATTA TTTGAGTGTA AATAATAGAT GTCAGTATTT ATCATGATGG GTCACATATA GACATATGTA CATATTATAT ATATATATAT ATATATATAT ATATATATAT ATATATATAT ATAAGCTCTT TTTTCTGAGG CTATTTTATA GTTATTTTAA AACATAAAGA TACAGAAGTC TTCTTACTT CTGATTTTCA AAACCATTCC TCAGTATCTT CAGGCATTTG ACCTCCTGAA TGTGCTTGGC CCTGGGCTTC AGTTATCCTT TGATGTCTG CAGGGGTGGC TAATGTGCTG GGGTTTTTCT GTGTAAATAG TCACAGTATT GTTTTATTGG TGAATAGCTG AAAACAGAG</p>	
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		GGATTAAGTC ATATTCCGGG AAAGAGAATT ATAGTTTTTA TGCTCCTGT TGAATAAATG GTGTCCTGAT TGCCTGGG	
DMD	NM_0 00109. 3	CGTTAAATGC AAACGCTGCT CTGGCTCATG TGTTTGCTCC GAGGTATAGG TTTTGTTCTGA CTGACGTATC AGATAGTCAG AGTGGTTACC ACACCGACGT TGTAGCAGCT GCATAATAAA TGACTGAAAG AATCATGTTA GGCATGCCCA CCTAACCTAA CTTGAATCAT GCGAAAGGGG AGCTGTTGGA ATTCAAATAG ACTTCTGGT TCCAGCAGT CGGCAGTAAT AGAATGCTTT CAGGAAGATG ACAGAATCAG GAGAAAGATG CTGTTTTGCA CTATCTTGAT TTGTTACAGC AGCCAACCTTA TTGGCATGAT GGAGTGACAG GAAAAACAGC TGGCATGGAA GATGAAAGAG AAGATGTTCA AAAGAAAACA TTCACAAAAT GGGTAAATGC ACAATTTTCT AAGTTTGGGA AGCAGCATAT TGAGAACCTC TTCAGTGACC TACAGGATGG GAGGCGCCTC CTAGACCTCC TCGAAGGCCT GACAGGGCAA AAACCTGCCAA AAGAAAAAGG ATCCACAAGA GTTCATGCC TGAACAATGT CAACAAGGCA CTGCGGGTTT TGCAGAACAA TAATGTTGAT TTAGTGAATA TTGGAAGTAC TGACATCGTA GATGGAAATC ATAACTGAC TCTTGGTTTG ATTTGGAATA TAATCCTCCA CTGGCAGGTC AAAAATGTAA TGAAAAATAT CATGGCTGGA TTGCAACAAA CCAACAGTGA AAAGATTCTC CTGAGCTGGG TCCGACAATC AACTCGTAAT TATCCACAGG TTAATGTAAT CAACTTCACC ACCAGCTGGT CTGATGGCCT GGCTTTGAAT GCTCTCATCC ATAGTCATAG GCCAGACCTA TTTGACTGGA ATAGTGTGGT TTGCCAGCAG TCAGCCACAC AACGACTGGA ACATGCATTC AACATCGCCA GATATCAATT AGGCATAGAG AAACACTCG ATCCTGAAGA TGTTGATACC ACCTATCCAG ATAAGAAGTC CATCTTAATG TACATCACAT CACTCTTCCA AGTTTTGCCT CAACAAGTGA GCATTGAAGC CATCCAGGAA GTGGAAATGT TGCCAAGGCC ACCTAAAGTG ACTAAAGAAG AACATTTTCA GTTACATCAT CAAATGCACT ATTCTCAACA GATCACGGTC AGTCTAGCAC AGGGATATGA GAGAACTTCT TCCCCTAAGC CTCGATTCAA GAGCTATGCC TACACACAGG CTGCTTATGT CACCACCTCT GACCCTACAC GGAGCCCATT TCCTTCACAG CATTTGGAAG CTCTGAAGA CAAGTCATTT GGCAGTTCAT TGATGGAGAG TGAAGTAAAC CTGGACCGTT ATCAAACAGC TTTAGAAGAA GTATTATCGT GGCTTCTTTC TGCTGAGGAC ACATTGCAAG CACAAGGAGA GATTTCTAAT GATGTGGAAG TGGTGAAAGA CCAGTTTCAT ACTCATGAGG GGTACATGAT GGATTTGACA GCCCATCAGG GCCGGGTTGG TAATATTCTA CAATTGGGAA GTAAGCTGAT TGGAACAGGA AAATTATCAG	8

	<p>AAGATGAAGA AACTGAAGTA CAAGAGCAGA TGAATCTCCT AAATTCAAGA TGGGAATGCC TCAGGGTAGC TAGCATGGAA AAACAAAGCA ATTTACATAG AGTTTTAATG GATCTCCAGA ATCAGAAACT GAAAGAGTTG AATGACTGGC TAACAAAAAC AGAAGAAAGA ACAAGGAAAA TGGAGGAAGA GCCTCTTGGA CCTGATCTTG AAGACCTAAA ACGCCAAGTA CAACAACATA AGGTGCTTCA AGAAGATCTA GAACAAGAAC AAGTCAGGGT CAATICTCTC ACTCACATGG TGGTGGTAGT TGATGAATCT AGTGGAGATC ACGCAACTGC TGCTTTGGAA GAACAACCTA AGGTATTGGG AGATCGATGG GCAAACATCT GTAGATGGAC AGAAGACCGC TGGGTTCTTT TACAAGACAT CCTTCTCAA TGGCAACGTC TTAAGTGAAGA ACAGTGCCTT TTAGTGCAT GGCTTTCAGA AAAAGAAGAT GCAGTGAACA AGATTCACAC AACTGGCTTT AAAGATCAAA ATGAAATGTT ATCAAGTCTT CAAAAAGTGG CCGTTTTAAA AGCGGATCTA GAAAAGAAAA AGCAATCCAT GGGCAAAGT TATTCACTCA AACAAGATCT TCTTTCAACA CTGAAGAATA AGTCAGTGAC CCAGAAGACG GAAGCATGGC TGGATAACTT TGCCCGGTGT TGGGATAATT TAGTCCAAA ACTTGAAAAG AGTACAGCAC AGATTTTACA GGCTGTCACC ACCACTCAGC CATCACTAAC ACAGACAAC GTAATGGAAA CAGTAACTAC GGTGACCACA AGGGAACAGA TCCTGGTAAA GCATGCTCAA GAGGAACTTC CACCACCACC TCCCCAAAAG AAGAGGCAGA TTAAGTGTGA TTCTGAAATT AGGAAAAGGT TGGATGTTGA TATAACTGAA CTTACAGCT GGATTACTCG CTCAGAAGCT GTGTTCGAGA GTCCTGAATT TGCAATCTTT CGGAAGGAAG GCAACTTCTC AGACTTAAAA GAAAAGTCA ATGCCATAGA GCGAGAAAAA GCTGAGAAGT TCAGAAAAGT GCAAGATGCC AGCAGATCAG CTCAGGCCCT GGTGGAACAG ATGGTGAATG AGGGTGTAA TGCAGATAGC ATCAAACAAG CCTCAGAACA ACTGAACAGC CGGTGGATCG AATTCTGCCA GTTGCTAAGT GAGAGACTTA ACTGGCTGGA GTATCAGAAC AACATCATCG CTTTCTATAA TCAGCTACAA CAATTGGAGC AGATGACAAC TACTGCTGAA AACTGGTTGA AAATCCAACC CACCACCCCA TCAGAGCCAA CAGCAATTAA AAGTCAGTTA AAAATTTGTA AGGATGAAGT CAACCGGCTA TCAGGTCTTC AACCTCAAAT TGAACGATTA AAAATTCAAA GCATAGCCCT GAAAGAGAAA GGACAAGGAC CCATGTTCTT GGATGCAGAC TTTGTGGCCT TTACAAATCA TTTAAGCAA GTCTTTTCTG ATGTGCAGGC CAGAGAGAAA GAGCTACAGA CAATTTTTGA CACTTTGCCA CCAATGCGCT ATCAGGAGAC CATGAGTGCC ATCAGGACAT GGGTCCAGCA GTCAGAAACC AAACTCTCCA TACCTCAACT TAGTGTCCACC GACTATGAAA</p>	
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	<p>TCATGGAGCA GAGACTCGGG GAATTGCAGG CTTTACAAAG TTCTCTGCAA GAGCAACAAA GTGGCCTATA CTATCTCAGC ACCACTGTGA AAGAGATGTC GAAGAAAGCG CCCTCTGAAA TTAGCCGGAA ATATCAATCA GAATTTGAAG AAATTGAGGG ACGCTGGAAG AAGCTCTCCT CCCAGCTGGT TGAGCATTGT CAAAAGCTAG AGGAGCAAAT GAATAAACTC CGAAAAATTC AGAATCACAT ACAAACCCTG AAGAAATGGA TGGCTGAAGT TGATGTTTTT CTGAAGGAGG AATGGCCTGC CCTTGGGGAT TCAGAAATTC TAAAAAAGCA GCTGAAACAG TGCAGACTTT TAGTCAGTGA TATTCAGACA ATTCAGCCCA GTCTAAACAG TGTC AATGAA GGTGGGCAGA AGATAAAGAA TGAAGCAGAG CCAGAGTTTG CTTGAGACT TGAGACAGAA CTCAAAGAAC TTAACACTCA GTGGGATCAC ATGTGCCAAC AGGTCTATGC CAGAAAGGAG GCCTTGAAGG GAGGTTTGA GAAAAC TGTA AGCCTCCAGA AAGATCTATC AGAGATGCAC GAATGGATGA CACAAGCTGA AGAAGAGTAT CTTGAGAGAG ATTTTGAATA TAAAAC TCCA GATGAATTAC AGAAAGCAGT TGAAGAGATG AAGAGAGCTA AAGAAGAGGC CCAACAAAAA GAAGCGAAAG TGAAACTCCT TACTGAGTCT GTAAATAGTG TCATAGCTCA AGCTCCACCT GTAGCACAAG AGGCCTTAAA AAAGGAACTT GAAACTCTAA CCACCAACTA CCAGTGGCTC TGC ACTAGGC TGAATGGGAA ATGCAAGACT TTGGAAGAAG TTTGGGCATG TTGGCATGAG TTATTGTCAT ACTTGGAGAA AGCAAACAAG TGGCTAAATG AAGTAGAATT TAAACTTAAA ACCACTGAAA ACATTCCTGG CGGAGCTGAG GAAATCTCTG AGGTGCTAGA TTC ACTTGAA AATTTGATGC GACATTCAGA GGATAACCCA AATCAGATTC GCATATTGGC ACAGACCCTA ACAGATGGCG GAGTCATGGA TGAGCTAATC AATGAGGAAC TTGAGACATT TAATTCCTCGT TGGAGGGAAC TACATGAAGA GGCTGTAAGG AGGCAAAAGT TGCTTGAACA GAGCATCCAG TCTGCCCAGG AGACTGAAAA ATCCTTACAC TTAATCCAGG AGTCCCTCAC ATTCATTGAC AAGCAGTTGG CAGCTTATAT TGCAGACAAG GTGGACGCAG CTCAAATGCC TCAGGAAGCC CAGAAAATCC AATCTGATTT GACAAGTCAT GAGATCAGTT TAGAAGAAAT GAAGAAACAT AATCAGGGGA AGGAGGCTGC CCAAAGAGTC CTGTCTCAGA TTGATGTTGC ACAGAAAAAA TTACAAGATG TCTCCATGAA GTTTCGATTA TTCCAGAAAC CAGCCAATTT TGAGCAGCGT CTACAAGAAA GTAAGATGAT TTTAGATGAA GTGAAGATGC ACTTGCCTGC ATTGGAAACA AAGAGTGTGG AACAGGAAGT AGTACAGTCA CAGCTAAATC ATTGTGTGAA CTTGTATAAA AGTCTGAGTG AAGTGAAGTC TGAAGTGGAA ATGGTGATAA AGACTGGACG TCAGATTGTA CAGAAAAAGC</p>	
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	<p>AGACGGAAAA TCCCAAAGAA CTTGATGAAA GAGTAACAGC TTTGAAATTG CATTATAATG AGCTGGGAGC AAAGGTAACA GAAAGAAAGC AACAGTTGGA GAAATGCTTG AAATTGTCCC GTAAGATGCG AAAGGAAATG AATGTCTTGA CAGAATGGCT GGCAGCTACA GATATGGAAT TGACAAAGAG ATCAGCAGTT GAAGGAATGC CTAGTAATTT GGATTCTGAA GTTGCCTGGG GAAAGGCTAC TCAAAAAGAG ATTGAGAAAC AGAAGGTGCA CCTGAAGAGT ATCACAGAGG TAGGAGAGGC CTTGAAAACA GTTTTGGGCA AGAAGGAGAC GTTGGTGGAA GATAAACTCA GTCTTCTGAA TAGTAACTGG ATAGCTGTCA CCTCCCGAGC AGAAGAGTGG TTAAATCTTT TGTTGGAATA CCAGAAACAC ATGGAAACTT TTGACCAGAA TGTGGACCAC ATCACAAAGT GGATCATTCA GGCTGACACA CTTTTGGATG AATCAGAGAA AAAGAAACC CAGCAAAAAG AAGACGTGCT TAAGCGTTTA AAGGCAGAAC TGAATGACAT ACGCCCAAAG GTGGACTCTA CACGTGACCA AGCAGCAAAC TTGATGGCAA ACCGCGGTGA CCACTGCAGG AAATTAGTAG AGCCCCAAAT CTCAGAGCTC AACCATCGAT TTGCAGCCAT TTCACACAGA ATTAAGACTG GAAAGGCCTC CATTCTTTG AAGGAATTGG AGCAGTTTAA CTCAGATATA CAAAAATTGC TTGAACCACT GGAGGCTGAA ATTCAGCAGG GGGTGAATCT GAAAGAGGAA GACTTCAATA AAGATATGAA TGAAGACAAT GAGGGTACTG TAAAAGAATT GTTGCAAAGA GGAGACAACT TACAACAAAG AATCACAGAT GAGAGAAAGC GAGAGGAAAT AAAGATAAAA CAGCAGCTGT TACAGACAAA ACATAATGCT CTCAAGGATT TGAGGTCTCA AAGAAGAAAA AAGGCTCTAG AAATTTCTCA TCAGTGGTAT CAGTACAAGA GGCAGGCTGA TGATCTCCTG AAATGCTTGG ATGACATTGA AAAAAATTA GCCAGCCTAC CTGAGCCCAG AGATGAAAGG AAAATAAAGG AAATTGATCG GGAATTGCAG AAGAAGAAAG AGGAGCTGAA TGCAGTGCGT AGGCAAGCTG AGGGCTTGTC TGAGGATGGG GCCGCAATGG CAGTGGAGCC AACTCAGATC CAGCTCAGCA AGCGCTGGCG GAAATTGAG AGCAAATTTG CTCAGTTTCG AAGACTCAAC TTTGCACAAA TTCACACTGT CCGTGAAGAA ACGATGATGG TGATGACTGA AGACATGCCT TTGGAAATTT CTTATGTGCC TTCTACTTAT TTGACTGAAA TCACTCATGT CTCACAAGCC CTATTAGAAG TGGAACAAC TCTCAATGCT CCTGACCTCT GTGCTAAGGA CTTTGAAGAT CTCTTTAAGC AAGAGGAGTC TCTGAAGAAT ATAAAAGATA GTCTACAACA AAGCTCAGGT CGGATTGACA TTATTCATAG CAAGAAGACA GCAGCATTGC AAAGTGCAAC GCCTGTGGAA AGGGTGAAGC TACAGGAAGC TCTCTCCAG CTTGATTTC AATGGGAAAA AGTTAACAAA ATGTACAAGG</p>	
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	<p>ACCGACAAGG GCGATTGAC AGATCTGTTG AGAAATGGCG GCGTTTTTCAT TATGATATAA AGATATTTAA TCAGTGGCTA ACAGAAGCTG AACAGTTTCT CAGAAAGACA CAAATTCCTG AGAATTGGGA ACATGCTAAA TACAAATGGT ATCTTAAGGA ACTCCAGGAT GGCATTGGGC AGCGGCAAAC TGTTGTCAGA ACATTGAATG CAACTGGGGA AGAAATAATT CAGCAATCCT CAAAAACAGA TGCCAGTATT CTACAGGAAA AATTGGGAAG CCTGAATCTG CGGTGGCAGG AGGTCTGCAA ACAGCTGTCA GACAGAAAAA AGAGGCTAGA AGAACAAAAG AATATCTTGT CAGAATTTCA AAGAGATTTA AATGAATTTG TTTTATGGTT GGAGGAAGCA GATAACATTG CTAGTATCCC ACTTGAACCT GGAAAAGAGC AGCAACTAAA AGAAAAGCTT GAGCAAGTCA AGTTACTGGT GGAAGAGTTG CCCCTGCGCC AGGGAATTCT CAAACAATTA AATGAACTG GAGGACCCGT GCTTGTAAGT GCTCCATAA GCCCAGAAGA GCAAGATAAA CTTGAAAATA AGCTCAAGCA GACAAATCTC CAGTGGATAA AGGTTTCCAG AGCTTTACCT GAGAAACAAG GAGAAATTGA AGCTCAAATA AAAGACCTTG GGCAGCTTGA AAAAAAGCTT GAAGACCTTG AAGAGCAGTT AAATCATCTG CTGCTGTGGT TATCTCCTAT TAGGAATCAG TTGGAAATTT ATAACCAACC AAACCAAGAA GGACCATTTG ACGTTCAGGA AACTGAAATA GCAGTTCAAG CTAAACAACC GGATGTGGAA GAGATTTTGT CTAAAGGGCA GCATTTGTAC AAGGAAAAAC CAGCCACTCA GCCAGTGAAG AGGAAGTTAG AAGATCTGAG CTCTGAGTGG AAGGCGGTAA ACCGTTTACT TCAAGAGCTG AGGGCAAAGC AGCCTGACCT AGTCTCTGGA CTGACCACTA TTGGAGCCTC TCCTACTCAG ACTGTTACTC TGGTGACACA ACCTGTGGTT ACTAAGGAAA CTGCCATCTC CAAACTAGAA ATGCCATCTT CTTGATGTT GGAGGTACCT GCTCTGGCAG ATTTCAACCG GGCTTGGACA GAACTTACCG ACTGGCTTTC TCTGCTTGAT CAAGTTATAA AATCACAGAG GGTGATGGTG GGTGACCTTG AGGATATCAA CGAGATGATC ATCAAGCAGA AGGCAACAAT GCAGGATTTG GAACAGAGGC GTCCCAGTT GGAAGAACTC ATTACCGCTG CCCAAAATTT GAAAAACAAG ACCAGCAATC AAGAGGCTAG ACAATCATT ACGGATCGAA TTGAAAGAAT TCAGAATCAG TGGGATGAAG TACAAGAACA CCTTCAGAAC CGGAGGCAAC AGTTGAATGA AATGTTAAAG GATTCAACAC AATGGCTGGA AGCTAAGGAA GAAGCTGAGC AGGTCTTAGG ACAGGCCAGA GCCAAGCTTG AGTCATGGAA GGAGGGTCCC TATACAGTAG ATGCAATCCA AAAGAAAATC ACAGAAACCA AGCAGTTGGC CAAAGACCTC CGCCAGTGGC AGACAAATGT AGATGTGGCA AATGACTTGG CCCTGAAACT TCTCCGGGAT TATTCTGCAG</p>	
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	<p>ATGATACCAG AAAAGTCCAC ATGATAACAG AGAATATCAA TGCTCTTGG AGAAGCATTG ATAAAAGGGT GAGTGAGCGA GAGGCTGCTT TGGAAGAAAC TCATAGATTA CTGCAACAGT TCCCCCTGGA CCTGGAAAAG TTTCTTGCCT GGCTTACAGA AGCTGAAACA ACTGCCAATG TCCTACAGGA TGCTACCCGT AAGGAAAGGC TCCTAGAAGA CTCCAAGGGA GTAAAAGAGC TGATGAAACA ATGGCAAGAC CTCCAAGGTG AAATTGAAGC TCACACAGAT GTTTATCACA ACCTGGATGA AAACAGCCAA AAAATCCTGA GATCCCTGGA AGGTTCCGAT GATGCAGTCC TGTTACAAAG ACGTTTGGAT AACATGAACT TCAAGTGGAG TGAACTTCGG AAAAAGTCTC TCAACATTAG GTCCCATTTG GAAGCCAGTT CTGACCAGTG GAAGCGTCTG CACCTTCTC TGCAGGAACT TCTGGTGTGG CTACAGCTGA AAGATGATGA ATTAAGCCGG CAGGCACCTA TTGGAGGCGA CTTTCCAGCA GTTTCAAGAGC AGAACGATGT ACATAGGGCC TTCAAGAGGG AATTGAAAAC TAAAGAACCCT GTAATCATGA GTACTCTTGA GACTGTACGA ATATTTCTGA CAGAGCAGCC TTTGGAAGGA CTAGAGAAAC TCTACCAGGA GCCCAGAGAG CTGCCTCCTG AGGAGAGAGC CCAGAATGTC ACTCGGCTTC TACGAAAGCA GGCTGAGGAG GTCAATACTG AGTGGGAAAA ATTGAACCTG CACTCCGCTG ACTGGCAGAG AAAAATAGAT GAGACCCTTG AAAGACTCCA GGAACCTCAA GAGGCCACGG ATGAGCTGGA CCTCAAGCTG CGCCAAGCTG AGGTGATCAA GGGATCCTGG CAGCCCGTGG GCGATCTCCT CATTGACTCT CTCCAAGATC ACCTCGAGAA AGTCAAGGCA CTTCGAGGAG AAATTGCGCC TCTGAAAGAG AACGTGAGCC ACGTCAATGA CCTTGCTCGC CAGCTTACCA CTTTGGGCAT TCAGCTCTCA CCGTATAACC TCAGCACTCT GGAAGACCTG AACACCAGAT GGAAGCTTCT GCAGGTGGCC GTCGAGGACC GAGTCAGGCA GCTGCATGAA GCCACAGGG ACTTTGGTCC AGCATCTCAG CACTTTCCTT CCACGTCTGT CCAGGGTCCC TGGGAGAGAG CCATCTCGCC AAACAAAGTG CCCTACTATA TCAACCACGA GACTCAAACA ACTTGCTGGG ACCATCCCAA AATGACAGAG CTCTACCAGT CTTTAGCTGA CCTGAATAAT GTCAGATTCT CAGCTTATAG GACTGCCATG AAACCTCCGAA GACTGCAGAA GGCCCTTTGC TTGGATCTCT TGAGCCTGTC AGCTGCATGT GATGCCITGG ACCAGCACAA CCTCAAGCAA AATGACCAGC CCATGGATAT CCTGCAGATT ATTAATTGTT TGACCACTAT TTATGACCGC CTGGAGCAAG AGCACAAACA TTTGGTCAAC GTCCCTCTCT GCGTGGATAT GTGTCTGAAC TGGCTGCTGA ATGTTTATGA TACGGGACGA ACAGGGAGGA TCCGTGTCCT GTCTTTTAAA ACTGGCATCA TTTCCCTGTG TAAAGCACAT TTGGAAGACA</p>	
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	<p>AGTACAGATA CCTTTTCAAG CAAGTGGCAA GTTCAACAGG ATTTTGTGAC CAGCGCAGGC TGGGCCTCCT TCTGCATGAT TCTATCCAAA TTCCAAGACA GTTGGGTGAA GTTGCATCCT TTGGGGGCAG TAACATTGAG CCAAGTGTCC GGAGCTGCTT CCAATTTGCT AATAATAAGC CAGAGATCGA AGCGGCCCTC TTCCTAGACT GGATGAGACT GGAACCCAG TCCATGGTGT GGCTGCCCGT CCTGCACAGA GTGGCTGCTG CAGAACTGC CAAGCATCAG GCCAAATGTA ACATCTGCAA AGAGTGTCCA ATCATTGGAT TCAGGTACAG GAGTCTAAAG CACTTTAATT ATGACATCTG CCAAAGCTGC TTTTTTCTG GTCGAGTTGC AAAAGGCCAT AAAATGCACT ATCCCATGGT GGAATATTGC ACTCCGACTA CATCAGGAGA AGATGTTCGA GACTTTGCCA AGGTACTAAA AAACAAATTT CGAACCAAAA GGTATTTTGC GAAGCATCCC CGAATGGGCT ACCTGCCAGT GCAGACTGTC TTAGAGGGGG ACAACATGGA AACTCCCGTT ACTCTGATCA ACTTCTGGCC AGTAGATTCT GCGCCTGCCT CGTCCCCTCA GCTTTCACAC GATGATACTC ATTCACGCAT TGAACATTAT GCTAGCAGGC TAGCAGAAAT GGAAAACAGC AATGGATCTT ATCTAAATGA TAGCATCTCT CCTAATGAGA GCATAGATGA TGAACATTTG TTAATCCAGC ATTACTGCCA AAGTTTGAAC CAGGACTCCC CCCTGAGCCA GCCTCGTAGT CCTGCCCAGA TCTTGATTTT CTTAGAGAGT GAGGAAAGAG GGGAGCTAGA GAGAATCCTA GCAGATCTTG AGGAAGAAAA CAGGAATCTG CAAGCAGAAT ATGACCGTCT AAAGCAGCAG CACGAACATA AAGGCCTGTC CCCACTGCCG TCCCCTCCTG AAATGATGCC CACCTCTCCC CAGAGTCCCC GGGATGCTGA GTCATTGCT GAGGCCAAGC TACTGCGTCA ACACAAAGGC CGCCTGGAAG CCAGGATGCA AATCCTGGAA GACCACAATA AACAGCTGGA GTCACAGTTA CACAGGCTAA GGCAGCTGCT GGAGCAACCC CAGGCAGAGG CCAAAGTGAA TGGCACAACG GTGTCTCTC CTTCTACCTC TCTACAGAGG TCCGACAGCA GTCAGCCTAT GCTGCTCCGA GTGGTTGGCA GTCAAACTTC GGACTCCATG GGTGAGGAAG ATCTTCTCAG TCCTCCCCAG GACACAAGCA CAGGGTTAGA GGAGGTGATG GAGCAACTCA ACAACTCCTT CCCTAGTTCA AGAGGAAGAA ATACCCCTGG AAAGCCAATG AGAGAGGACA CAATGTAGGA AGTCTTTTCC ACATGGCAGA TGATTTGGGC AGAGCGATGG AGTCCTTAGT ATCAGTCATG ACAGATGAAG AAGGAGCAGA ATAAATGTTT TACAACCTCT GATTCCCGCA TGGTTTTTAT AATATTCATA CAACAAAGAG GATTAGACAG TAAGAGTTTA CAAGAAATAA ATCTATATTT TTGTGAAGGG TAGTGGTATT AACTGTAGA TTTCAGTAGT TTCTAAGTCT GTTATTGTTT TGTTAACAAT GGCAGGTTTT</p>	
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	<p>ACACGTCTAT GCAATTGTAC AAAAAAGTTA TAAGAAAAC ACATGTAAAA TCTTGATAGC TAAATAACTT GCCATTTCTT TATATGGAAC GCATTTTGGG TTGTTTAAAA ATTTATAACA GTTATAAAGA AAGATTGTAA ACTAAAGTGT GCTTTATAAA AAAAAGTTGT TTATAAAAAC CCCTAAAAAC AAAACAAACA CACACACACA CACATACACA CACACACACA AACTTTTGAG GCAGCGCATT GTTTTGCATC CTTTTGGCGT GATATCCATA TGAAATTCAT GGCTTTTTCT TTTTTTGCAT ATTAAAGATA AGACTTCCTC TACCACCACA CCAAATGACT ACTACACACT GCTCATTTGA GAACTGTCAG CTGAGTGGGG CAGGCTTGAG TTTTCATTTT ATATATCTAT ATGTCTATAA GTATATAAAT ACTATAGTTA TATAGATAAA GAGATACGAA TTTCTATAGA CTGACTTTTT CCATTTTTTA AATGTTICATG TCACATCCTA ATAGAAAGAA ATTACTTCTA GTCAGTCATC CAGGCTTACC TGCTTGGTCT AGAATGGATT TTTCCCGGAG CCGGAAGCCA GGAGGAAACT ACACCACACT AAAACATTGT CTACAGCTCC AGATGTTTTCT CATTTTAAAC AACTTTCCAC TGACAACGAA AGTAAAGTAA AGTATTGGAT TTTTTTAAAG GGAACATGTG AATGAATACA CAGGACTTAT TATATCAGAG TGAGTAATCG GTTGGTTGGT TGATTGATTG ATTGATTIGAT ACATTCAGCT TCCTGCTGCT AGCAATGCCA CGATTTAGAT TTAATGATGC TTCAGTGGAA ATCAATCAGA AGGTATTCTG ACCTTGTGAA CATCAGAAGG TATTTTTTAA CTCCAAGCA GTAGCAGGAC GATGATAGGG CTGGAGGGCT ATGGATTCCC AGCCCATCCC TGTGAAGGAG TAGGCCACTC TTTAAGTGAA GGATTGGATG ATTGTTTATA ATACATAAAG TTCTCTGTAA TTACAATAA ATTATTATGC CCTCTTCTCA CAGTCAAAG GAACTGGGTG GTTTGGTTTT TGTGTCTTTT TTAGATTTAT TGTCCCATGT GGGATGAGTT TTAAATGCC ACAAGACATA ATTTAAATA AATAAACTTT GGGAAAAGGT GTAAAACAGT AGCCCATCA CATTTGTGAT ACTGACAGGT ATCAACCCAG AAGCCCATGA ACTGTGTTTC CATCCTTTGC ATTCTCTGAG GAGTAGTTCC ACACAGGTTT GTAAGTAAGT AAGAAAGAAG GCAAATTGAT TCAAATGTTA CAAAAAACC CTCTTGGTG GATTAGACAG GTTAAATATA TAAACAAACA AACAAAAATT GCTCAAAAAA GAGGAGAAAA GCTCAAGAGG AAAAGCTAAG GACTGGTAGG AAAAAGCTTT ACTCTTTCAT GCCATTTTAT TTCTTTTTGA TTTTTAAATC ATTCATTCAA TAGATACCAC CGTGTGACCT ATAATTTTGC AAATCTGTTA CCTCTGACAT CAAGTGTAAT TAGCTTTTGG AGAGTGGGCT GACATCAAGT GTAATTAGCT TTTGGAGAGT GGGTTTTTGC CATTATTAAT AATTAATTA TTAACATCAA ACACGGCTTC TCATGCTATT TCTACCTCAC</p>	
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		<p>TTTGGTTTTG GGGTGTTCCT GATAATTGTG CACACCTGAG TTCACAGCTT CACCACTTGT CCATTGCGTT ATTTTCTTTT TCCTTTATAA TTCTTTCTTT TTCCTTCATA ATTTTCAAAA GAAAACCCAA AGCTCTAAGG TAACAAATTA CCAAATTACA TGAAGATTG GTTTTGTCT TGCATTTTTT TCCTTTATGT GACGCTGGAC CTTTTCTTTA CCCAAGGATT TTAAAACTC AGATTTAAAA CAAGGGGTTA CTTTACATCC TACTAAGAAG TTAAGTAAG TAAGTTTCAT TCTAAAATCA GAGGTAAATA GAGTGCATAA ATAATTTTGT TTTAATCTTT TTGTTTTTCT TTTAGACACA TTAGCTCTGG AGTGAGTCTG TCATAATATT TGAACAAAAA TTGAGAGCTT TATTGCTGCA TTTAAGCAT AATTAATTTG GACATTATTT CGTGTGTGT TCTTTATAAC CACCAAGTAT TAAACTGTAA ATCATAATGT AACTGAAGCA TAAACATCAC ATGGCATGTT TTGTCATTGT TTTCAGGTAC TGAGTTCTTA CTGAGTATC ATAATATATT GTGTTTTAAC ACCAACACTG TAACATTTAC GAATTATTTT TTAAACTTC AGTTTTACTG CATTTCACA ACATATCAGA CTTACCCAAA TATATGCCTT ACTATTGTAT TATAGTACTG CTTTACTGTG TATCTCAATA AAGCACGCAG TTATGTTAC</p>	
<p>DNAJC 9</p>	<p>NM_0 15190. 4</p>	<p>GGATGCGCGG CGTGGCCACG CCCCTTCAGT GCTTGTGACG CAGGCGCCCT GGGCTTTTTG GGC GCGAAAA AGAAGCAGTC CTGGGTTGTA CCCGGCGCAG CTGGGAGCGG CTGCTTCCTC CGGGGTGTA TCTCCGCCCG GCATGGGGCT GCTGGACCTT TGCGAGGAAG TGTTCCGGCAC CGCCGACCTT TACCGGGTGC TGGGCGTGCG ACGCGAGGCC TCCGACGGCG AGGTCCGACG AGGCTACCAC AAGGTGTCCC TGCAGGTACA CCCGGACCGG GTGGGTGAGG GCGACAAGGA GGACGCCACC CGCCGCTTCC AGATCCTGGG AAAAGTCTAT TCCGTTCTCA GTGACAGAGA ACAGAGAGCA GTGTACGATG AGCAGGGAAC AGTGGACGAG GACTCTCCTG TGCTCACCCA AGACCGAGAC TGGGAGGCGT ATTGGCGGCT ACTCTTTAAA AAGATATCTT TAGAGGACAT TCAAGCTTTT GAAAAGACAT ACAAAGGTTC GGAAGAAGAG CTGGCTGATA TTAAGCAGGC CTATCTGGAC TTCAAGGGTG ACATGGATCA GATCATGGAG TCTGTGCTTT GCGTGCAGTA CACAGAGGAA CCCAGGATAA GGAATATCAT TCAGCAAGCT ATTGACGCCG GAGAGGTCCC ATCCTATAAT GCCTTTGTCA AAGAATCGAA ACAAAGATG AATGCAAGGA AAAGGAGGGC TCAGGAAGAG GCCAAAGAAG CAGAAATGAG CAGAAAGGAG TTGGGGCTTG ATGAAGGCGT GGATAGCCTG AAGGCAGCCA TTCAGAGCAG ACAAAGGAT CGGCAAAGG AAATGGACAA TTTTCTGGCT CAGATGGAAG CAAAGTACTG CAAATCTTCC</p>	<p>9</p>

		<p>AAAGGAGGAG GGAAAAAATC TGCTCTCAAG AAAGAAAAGA AATAATGGAA TTTTCTCTT CAAAGGTCCT TAGGTGTA TTGATGCCAT CGTAGGCAAG GTGCAGGCAG GATTTGAAGG CAAAAGTCAA TTCAGCTCTT GAGAAAAGGT GTCTTTCCAG CCTGAATTTT TCAGATTGAC TAGACCAAGC AGAATCTCTC AACCTGATCT TAGTATTTCC TAGAAAGCAC TTGACATTGT GTGAGGTCTC ACCTGAAGGA ACTTGGTGGT GACATTTGGG AGGGTGGAGG GAGGCAGTGT CCTTCCGAC AGCACTTGCC TCCATGGATC TTCTGTACAC AGAACTCTTA TCTAGGATGT GGTCTGTTC ATGCTGCTTT CTGCGATGTG CGTGTCTGTT AGAATAGGCT CTCTACCCAG CTAGAACCAC TTCCAGACAC TTGCTGGACA GCTATCTTCC ACATACTTCC CAGTTTACAT TTGGTCTTAA TGATCTTGAA TAGATCCTCT CTTCATTTTA CTCAGCCAGG TTTTGTACTG ATGTACAGGT GTTAAATTAC TTCAAGCATT TTTGTAAGAG GTGTATATAA TTCAATAAAA AAGGTAAAAC ATGATGATTA AGTTCTGGGG GCTTTGTAAA TGATCCCACT AAAATGTGAC CTAGGAAAAA TATGAATGGT GTTTAGGATG AGAGAAAAGG GGAAAACAAT AGCCCTGGTC AGCTTTATAA TAGAGAGCCT GGTTCCTTA GCATGAAGAG ATGTATGTTG TAGTCCTGCC ACTGATTACT GAAGTCCTGC CATTAAATTGC TGAATGTCTT CAGTTGGGCC ACTGAGCTTG TCTGAATCTG TTCCTTTTA TAAAAGAGTT ACTACATCAA AACAAAGAGT GAAATCCAAA TTTGTCAAAC TGTATGTATT AAACATGTCC AGTTTTTTGG TTAATAAAAA ATTGCACTGG CTTTGAGGGA AAACACACAG GGTGAGGGAA TTGGGCTAAA TGACTTCTTA CAGGCCCTT TCTGATTCTT TAACTTTGAA AGGCAAGCCA TATTGATCCA GTTGTTATAG TGAATCATG GTAATGGTTT GTGAGAACAA TAGAGATTTT CATTCTATG TAGATGAGTT GGTATGAGAA TATATGGAAT TTTAAGGGA CTGTTTAAAT CTTTGATTTG TAGACTATTA AATATACCGT ATGCATAAAG TAAGCCTTTA GCTCTAAGGT AAAGACGACA CGTTTTCCGGT TTGTGACTAC AAATAGGTTA AAAATAGATT TTAATTTTAT TAAAAATATA ATTTAATGCA GGTGTGTTGA AGCATCTGTC TTCATATGAT GGCATTAGAA CACCTTGGTA TAATAAAAAG TTACCGTAAT TTATGATTAT TTGAATTTAT CCATTCTGAA AATTAATAAG ATCTAAAACCT GGCATGACAA TCAAGATTTG TATTTAGTGA AATTTAAAAT AAATGTAAGC CATAGTTAAA AAAAAAAAAA AAAAA</p>	
<p>ENOSF 1</p>	<p>NM_0 011261 23.3</p>	<p>CGCCCTCCCG CCGCGCGCTC GGGATCCCGA CCAGTCCTGA CCGCACGGGG GCCGCGGCCA CGGGGCGCAG GGGCCATGGT GCGCGGCAGG ATCTCCCGGC TCTCGGTCCG GGACGTGCGC</p>	<p>10</p>

	<p>TTCCCCACGT CGCTTGGGGG CCACGGCGCG GACGCCATGC ACACGGACCC TGACTIONTCG GCTGCCTATG TCGTCATAGA AACTGATGCA GAAGATGGAA TCAAGGGGTG TGGAATTACC TTCACTCTGG GAAAAGGCAC TGAAGTTGAT TGGTCCAGAA AAGGGCGTGG TGCACCTGGC GACAGCGGCC GTCCTAAACG CGGTGTGGGA CTTGTGGGCC AAGCAGGAGG GAAAGCCTGT CTGGAAGTTA CTTGTGGACA TGGATCCCAG GATGCTGGTA TCCTGCATAG ATTTTCAGGTA CATCACTGAT GTCCTGACTG AGGAGGATGC CCTAGAAATA CTGCAGAAAAG GTCAAATTGG TAAAAAAGAA AGAGAGAAGC AAATGCTGGC ACAAGGATAC CCTGCTTACA CGACATCGTG CGCCTGGCTG GGGTACTCAG ATGACACGTT GAAGCAGCTC TGTGCCCAGG CGCTGAAGGA TGGCTGGACC AGGTTTAAAG TAAAGGTGGG TGCTGATCTC CAGGATGACA TGCGAAGATG CCAAATCATC CGAGACATGA TTGGACCGGA AAAGACTTTG ATGATGGATG CCAACCAGCG CTGGGATGTG CCTGAGGCGG TGGAGTGGAT GTCCAAGCTG GCCAAGTTCA AGCCATTGTG GATTGAGGAG CCAACCTCCC CTGATGACAT TCTGGGGCAC GCCACCATT CCAAGGCACT GGTCCCATTA GGAATTGGCA TTGCCACAGG AGAACAGTGC CACAATAGAG TGATATTTAA GCAACTCCTA CAGGCGAAGG CCCTGCAGTT CCTCCAGATT GACAGTTGCA GACTGGGCAG TGTC AATGAG AACCTCTCAG TATTGCTGAT GGCCAAAAAG TTTGAAATTC CTGTTTGCCC CCATGCTGGT GGAGTTGGCC TCTGTGAACT GGTGCAGCAC CTGATTATAT TTGACTACAT ATCAGTTTCT GCAAGCCTTG AAAATAGGGT GTGTGAGTAT GTTGACCACC TGCATGAGCA TTTCAAGTAT CCCGTGATGA TCCAGCGGGC TTCTACATG CCTCCCAAGG ATCCCGGCTA CTCAACAGAA ATGAAGGAGG AATCTGTAAA GAAACACCAG TATCCAGATG GTGAAGTTTG GAAGAACTC CTTCTGCTC AAGAAAATTA AGTGCTCAGC CCCAACAAC TTTTCTTTC TGAAGTGAAA GGGCTTAAAA TTTCTTGGAA ATAGTTTAC AAAAATGGAT TAAAAAATC CTACCGATCA AGATGAGTTC AGCTAGAAGT CATAACACC TCAGGAATCA GCTAAAGCAA AAAGAAC TTT TACCTCGGCA TCCAGCCCAA CCCCTAAAGA CTGACAATAT CCTTCGAGCT CCTTTGAAAG CACCCTAAC AGCCATTTCC ATTTTAATAG TTGGATGCGG ATTGTACCCT TCAATCTGAA AGTCTTCAGC TTTGAAGTCA TCAATTTTCT CAACTTTTCG AAGAATCCTG AGCTTTGGGA AAGGTCTGGG TTCTCGCTGA AGCTAAAAAC AAAATAAGGC CATTATTTTG CCATAATTGT ACGACCTGTT GTAATTGCTC CTCATGTCCG TGAAACAAGT ACACAGGATG TGATCAACAA AGTTCATTTT TACAGGAGTA TGATCCTGTC GATACCTTGC CGTAGGTTAT</p>	
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	<p>GTAACATGAT TGGAGCGCAA CCAGCTGTTC TCTTGCACAG ATCGAGAGTG AGGGGTATTT TGTGACATTA CACAGCATCA GGAGCCTGGT GCCTCATCAG GTGTAAGTTC TTATAACCAC TCTTGGCAAA TTTATTAAG ACAGGAACAC AGTCAATCTG TAACTCATAG TAGCTCTACG TTTACTTGAA TTCCACAATC CCTAACCCAT CTGTCCCTGG CAGAAAGAAG GAAAGATGAC ATGCATGGAC AGTGAACAGA AAGGGATGAA AGCCAGGATT CCTGGGATGA ACAGACAGTG GCAATTAGGA TGTGAAGACA GGTCAACAAC TATTACTATG TCTAAAAACG ACCAGAGCAG AGAGCCAGAG AGAATAAGCC TGAAGTCACC TCCACTCAAA AGCAGCCAAA CTCCCTCAAA GGAGTAACTT TTAAAACCTG GATCTAACCT GGAAGGGGCT AAAAAGTGTG TGGTTCTGAG TTTTTTCCT TAAGGCTCAT GAAGCAGATG AACTTACATT TTTATTGCCA TTCATATCA ATTGTTGGCT GCTATAACTT AGGGATTICA ACAGACTTTT GAAGTTTGGG CCTAAATATT GTACTTAATG TAAAATTAAC AAAAAATATT TATGGCCAGG GTGGTGGCTT ATGCCTGTAA TTCCAGAATT TTCGGAGGCT GAGGCAGGTG GATCACTTGA AGTCAGGAGT TTGAGACTAG CCTGGCCAAC ATGATGAAAC CCCATCTCTA CTAATAATAC AAAAATTAGC TGGGTGTGGT GGCATGTGCC TGTAATCCCA GCTACCTGGG AGGCTGAGGC AGAAGAATTG CTTGAACCCG GGAGGTGGAG GTTGCAGTGA GCTGAGATCG CACCACGGCA CACTCCAGCC TGGCCGACAG AGAAAGACTC CATCTCAAAA AAAAAAGAAA AGGAAAAACA TTTGCACTTC AATTCTCCTT CAAGTTAAAA TGAGTTAAAA TGCCCCCTTT TGGACAATCC CCTGGCTTGA ATGTGGCTCT TCCCTCTCTG GACTGGTGC TTAGTACCTC ACAGCACCTG ACATGTTAAG TGCCCATGGT TGCTGAGGCA GATGCCTGCC TTGTCCTGCC CACCTGCCCA CCACTTCTCC CTAAACTGAA GCCCCACATT TGGAGCAGTC ATCTTTATCT TGGACACAGC ATTGAGCAGA TGCCTGTTCC ACAGTCAACC TTTTATCAAG AGAAGGTACC AAACCCAAAA GTATAACATC TAATTCTTAC CTGAATTTTC AGTGGCTCGA TGTGATTCAG GTAAATATGT GCATCTCCCA AAGTGTGTAT AAAGTCACCT GGCTATAAAC CCGGGGGAGA AAGCAGAACA GTATGTTAGT TTCAATTCTT TAAAACATCA TTTAAAAACA TTAGAATATG CAGACACCGC AAGGCTTTTT TAAAAAAAT AATTTAGTGT AGCTTTTCCA TTTTTTTGTA GCAACAGCAT CTTGTTATGT TGCCAGGCT GGTATTGAAC TCCAGACCTC AAGCAATTGC TCCTGTCTCA GTCTCCCAA GTGCTGGGAT TACAGGCATG AGCCACCATA CCCAACCTCA GCATAGCTTT TGAGAAAATC CATAGAAGCT GTATCACAAA CAACCTGTAT AGATCTGTTA GTGCGTATAC CACAGGGCCA GAAAACCTTC</p>	
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		<p>CAGAAGAGGA AGGTTTCAA GTAAAAGCTG GTTCATTTCT TACTTACACA TATCAAATTT AAAAGCTAAT CAGAGACTAA ACTCTGCAAT TTGTTTTCCC ATATTAAGA ACTGAAGAGC TCAGTGTGGT AGGCTGGCAA GTCACCTTC CCGAGACAGC CCACCTTCAG GCCCGTGATG TGCGCAATCA TGTACGTGAG CAGGGCGTAG CTGGCGATGT TGAAAGGCAC ACCGAGGCC ATGTCTCCCG ATCTCTGGTA CAGCTGGCAG GACAGCTCAC TGTTACCAC ATAGA ACTGG CAGAGGGCAT GGCATGGAGG CAGCGCCATC AGAGGAAGAT CTGAGGAACC AGCAGAGGAA GATAAGGAGG GATGGTGGTT TGAAAGACCA CAGCTAAAGG CAAAGTAAA CAGGAGAGAA ACAGAAGCCA ACTCATATGG TGGAGACCAG GAGAGAGAGC CACTGGGCTG CAGTGATGTC CATAACAGCC TCTGCAGCGA TGGCACGGAG CTGAGGGGAGA CTATCCATCG GTGCAAGGTT TCTGCAGGTG TCCATTTACG GCTGAAGCAA TGCTCTTCCA TCAGAGCTGA AGGGATCTGG GCTACCTCGT GGCACCAGAT TACAAATACA GCAGGAATAA TTCTGTTTGC CACAGGAAAC TGGTGCTTCT GGTACACCCT CCTATATTAA AAGTCTCTAT TACATGGCAA AAAAAAAAAA AAAAAA</p>	
<p>FANCL</p>	<p>NM_0 011146 36.1</p>	<p>AGCGGACTGC GCATGTGCAG GACCCAGCAG GTCTAGAGCT TTTCTGTGTT TCTCCGACT TCGAGCCATG GCGGTGACGG AAGCGAGCCT GTTGCGCCAG TGCCCCCTGC TTCTGCCCA GAACCGGTCG AAAACCGTGT ATGAGGGATT CATCTCGGCT CAGGGAAGAG ACTTCCACCT TAGGATAGTG TTGCCTGAAG ATTTACA ACT GAAGAATGCA AGATTATTAT GTAGTTGGCA GCTGAGAACA AACTTAGTG GATAACCATCG AATAGTACAA CAGAGAATGC AGCACTCTCC TGATCTAATG AGCTTTATGA TGGAGTTGAA GATGCTTTTG GAAGTTGCCT TAAAGAATAG ACAAGAGCTG TATGCACTAC CTCCTCCTCC CCAGTTCTAC TCAAGCCTTA TTGAAGAGAT AGGAACTCTT GGTTGGGATA AACTTGTGTA TGCGGATACC TGCTTCAGTA CCATCAAGTT AAAAGCAGAA GATGCTTCTG GTAGAGAGCA TTTAATCACT CTCAAGTTGA AGGCAAAGTA TCCTGCAGAA TCACCAGATT ATTTGTGGA TTTTCTGTT CCATTTTGTG CCTCCTGGAC ACCTCAGGTA AATTCTCCTC AGAGCTCCTT AATAAGCATT TATAGTCAGT TTTTGGCAGC AATAGAATCA CTAAGGCAT TCTGGGATGT TATGGATGAA ATCGATGAGA AGACCTGGGT ACTTGAGCCA GAAAAACCTC CACGGAGTGC AACAGCACGC AGAATTGCAT TAGGTAATAA TGTTTCCATA AATATAGAGG TAGACCCAG GCATCCTACT ATGCTTCCTG AGTGCTTCTT TCTTGGAGCT GACCATGTGG TAAAACCCCT GGGAATTAAG</p>	<p>11</p>

		<p>CTGAGCAGGA ACATACATTT GTGGGATCCA GAAAATAGTG TGTTACAAAA TTTGAAAGAT GTTTTAGAAA TTGATTTTCC AGCTCGTGCT ATCCTGGAAA AATCTGATIT TACTATGGAT TGTGGAATTT GTTATGCTTA TCAACTTGAC GGTACCATT CTGATCAAGT GTGTGATAAT TCTCAGTGTG GACAACCTTT CCATCAAATA TGCTTATATG AGTGGCTGAG AGGACTACTA ACTAGTAGAC AGAGTTTTAA CATCATATTT GGTGAATGTC CATATTGTAG TAAGCCAATT ACCTTAAAAA TGTCIGGAAG GAAACACTGA AATAAGAATA CAACATTTTCG GTGAAGAGCT GGAAACTTAA AAAATTATCA AAAGGAATTT TGGTATCATC TTCAGAGAAA AAATAAAGCA AGAAATACTA ACATCAAAG GACAGGTATG ATGATGCGAT AATAATAAAC ATCTGCGTTT GTCTCTTCAC TAAGAGTAAA CTGGGAAATT GTAGGCCAAA GTCCAGTTGA ACTTTCTAAG TCTGTGATCC CCGTGCTGAC TGTGGAAGTG TATTTATACC AAGATGGAGA TCTTGACTTC TTGAATATAT CTGGACTGGT AAAATCTTGA TGAGGCTCAT AAAATGAGTT TGGGAATTGT GTATAGCTGA TTTTTTGTGG GAAACTGTTT ACTTCATTCA AAGGTTCTTG AGACTCTTGA TATTTCTGTC TTCTCCTTGT GCTTTCCTAT GGAAAAATA CATATATAGT TTAGTTTGTG AGACGTGAGT TATCCAAGTA TTTATTTTGT GTAGTGTGTA AGAATGCTAA ATAAAATGTT ATACAAGATC AAAAAAAAAA AAAAAAAAAA AAA</p>	
<p>HJURP</p>	<p>NM_0 18410. 4</p>	<p>CTATTTGAGT TTGTGGCGCG CGAGGCCCTG CAGTCCGGGT TGGCGCTTGG GTACTGGCTG GGTCCGATGC TGGGTACGCT GCGCGCCATG GAGGGCGAGG ACGTGGAAGA CGACCAGCTG CTGCAGAAGC TCAGGGCCAG TCGCCGCCGC TTCCAGAGGC GCATGCAGCG GCTGATAGAG AAGTACAACC AGCCCTTCGA GGACACCCCG GTGGTGCAA TGGCCACGCT GACCTACGAG ACGCCACAGG GATTGAGAAT TTGGGGTGGG AACTAATAA AGGAAAGAAA CGAAGGAGAG ATCCAGGACT CCTCCATGAA GCCCCGCGAC AGGACAGATG GCTCCGTGCA AGCTGCAGCC TGGGGTCCCTG AGCTTCCCTC GCACCGCACA GTCCCTGGGAG CCGATTCAA AAGCGGTGAG GTCGATGCCA CGTCAGACCA GGAAGAGTCA GTTGCTTGGG CCTTAGCACC TGCAGTGCCT CAAAGCCCTT TGAAAAATGA ATTAAGAAGG AAATACTTGA CCCAAGTGGG TATACTGCTA CAAGGTGCAG AGTATTTTGA GTGTGCAGGT AACAGAGCTG GAAGGGATGT ACGTGTGACT CCGCTGCCTT CACTGGCCTC ACCTGCCGTG CCTGCCCCCG GATACTGCAG TCGTATCTCC AGAAAGAGTC CTGGTGACCC AGCGAAACCA GCTTCATCTC CCAGAGAATG GGATCCTTTG CATCCTTCTT CCACAGACAT GGCCTTAGTA CCTAGAAATG</p>	<p>12</p>

	<p>ACAGCCTCTC CCTACAAGAG ACCAGTAGCA GCAGCTTCTT AAGCAGCCAG CCCTTTGAAG ATGATGACAT TTGCAATGTG ACCATCAGTG ACCTGTACGC AGGGATGCTG CACTCCATGA GCCGGCTGTT GAGCACAAAG CCATCAAGCA TCATCTCCAC CAAAACGTTT ATCATGCAAA ACTGGAACTC CAGGAGGAGG CACAGATATA AGAGCAGGAT GAACAAAACA TATTGCAAAG GAGCCAGACG TTCTCAGAGG AGCTCCAAGG AGAACTTCAT ACCTTGCTCT GAGCCTGTGA AAGGGACAGG GGCATTAAGA GATTGCAAGA ACGTATTAGA TGTTCCTTGC CGTAAGACAG GTTTAAAATT GGAAAAAGCT TTTCTTGAAG TCAACAGACC CCAAATCCAT AAGTTAGATC CAAGTTGGAA GGAGCGCAAA GTGACACCCT CGAAGTATC TTCCTTGATT TACTTCGACT CCAGTGCAAC ATATAATCTT GATGAGGAAA ATAGATTTAG GACATTAATA TGGTTAATTT CTCCTGTAAA AATAGTTTCC AGACCAACAA TACGACAGGG CCATGGAGAG AACCGTCAGA GGGAGATTGA AATCCGATTT GATCAGCTTC ATCGGGAATA TTGCCTGAGT CCCAGGAACC AGCCTCGCCG GATGTGCCTC CCGGACTCCT GGGCCATGAA CATGTACAGA GGGGGTCCCTG CGAGTCTGG TGGCCTCAG GGCTTAGAAA CCCGCAGGCT GAGTTTACCT TCCAGCAAAG CAAAAGCAAA AAGTTTAAGT GAGGCTTTTG AAAACCTAGG CAAAAGATCT CTGGAAGCAG GTAGGTGCCT GCCCAAGAGC GATTCATCTT CATCACTTCC AAAGACCAAC CCCACACACA GCGCAACTCG CCCGCAGCAG ACATCTGACC TTCACGTTCA GGGAAATAGT TCTGGAATAT TTAGAAAGTC AGTGTCAACC AGCAAACTC TTTCAGTCCC AGATAAAGAA GTGCCAGGCC ACGGAAGGAA TCGTTACGAT GAAATTAAG AAGAATTTGA CAAGCTTCAT CAAAAGTATT GCCTCAAATC TCCTGGGCAG ATGACAGTGC CTTTATGTAT TGGAGTGTCT ACAGATAAAG CAAGTATGGA AGTTTCGATAT CAAACAGAAG GCTTCTTAGG AAAATTAAT CCAGACCCTC ACTTCAGGG TTTCCAGAAG TTGCCATCAT CACCCCTGGG GTGCAGAAA AGTCTACTGG GCTCAACTGC AATTGAGGCT CCTTCATCTA CATGTGTTGC TCGTGCCATC ACGAGGGATG GCACGAGGGA CCATCAGTTC CCTGCAAAAA GACCCAGGCT ATCAGAACC CAGGGCTCCG GACGCCAGGG CAATTCCTG GGTGCCTCAG ATGGGGTGGA CAACACCGTC AGACCGGGAG ACCAGGGCAG CTCTTCACAG CCCAACTCAG AAGAGAGAGG AGAGAACACG TCTTACAGGA TGGAAGAGAA AAGTGATTC ATGCTAGAAA AATTGGAAAC TAAAAGTGTG TAGCTAGGTT ATTTCCGAGT GTTATTTATC TTCCACTTG CTCTCTGTTT GTATTTTTGT TTTGTTTTTG ATTCTTGAGA CTGTGAGGAC TTGGTTGACT TCTCTGCCCT TAAAGTAAAT ATTAGTGAAA</p>	
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		<p>TTGGTTCAT CAGAGATAAC CTCGAGTTCT TGGTGTAGAA ATTATGTGAA TAAAGTTGCT CAATTAGAAT TTTTAGGGTT CTCTTTGATA GGCCGTGTTT TCTGATGTGT GTGTTTTTTT TGGGGGGGGG TTATTTGTTT GTTTGTGTGT TTGTTTGT GTTTTTGAGA CAGTCTCTCT CTATCGCCCA GGCTGGAGTG CGGTGGCACA ATCTTGGCTC ACTGCAACTT CCGCCTCCCG GGTTCAAGCG ATTCTTCTGC CTCAGCCTCC CGAGTAGCTG GGATTACAGG CGCGCGCCAC CACGCCTGGC TAATTTTGT AGTTTTAGTA GAGACGGGGT TTCACCATAT TGACCAGGCT GGTCTCGAGC TCCTGGCCTC GTGATCCATC TGCCTCGGCC TCCCAAAGTG CTGGGATTAT AGGCGTGAGC CACTGCTCCC AGCCGTGTGT TCTTTTTTAA ATTTAGATAT GTCCAGAGAA TCCTCTCTCC TGTTTCCCAT TTCATTGAG AATATIGTTT GCTTGTGAGA CGTAAGTTCG AGCCCTGCAT GCAATGACCC TTGAAGGAAA ATAAACAGTC CTGGTGGTCC CAGACGCTCC TGCAGCCACA GCGCCTGTGA CTCCTCATGA TTCTTACTGA AGCTGTTGAT GACAGGATAT CATGGTGACG TTTTGTAAAT GAAATATTIC ACATATTGAG AATACATTGG TGAAACTCAT GCTGGAGTAA ATAGTTAATA TATGGCCAT</p>	
<p>HLA- DOA</p>	<p>NM_0 02119. 3</p>	<p>CTTCTTCTTT ACCTCCGCCT TGTTCCGTGC CTCACCACAC GGACTGAGAC TGATTTGATT AAAGCACCAG AGTGTAATGG CCCTCAGAGC AGGGCTGGTC CTGGGGTTCC ACACCCTGAT GACCCTCCTG AGCCCGCAGG AGGCAGGGGC CACCAAGGCT GACCACATGG GCTCCTACGG ACCCGCCTTC TACCAGTCTT ACGGCGCCTC GGGCCAGTTC ACCCATGAAT TTGATGAGGA ACAGCTGTTT TCTGTGGACC TGAAGAAAAG CGAGGCCGTG TGGCGTCTGC CTGAGTTTGG TGACTTTGCC CGCTTTGACC CGCAGGGCGG GCTGGCCGGC ATCGCCGCAA TCAAAGCCCA TCTGGACATC CTGGTGGAGC GCTCCAACCG CAGCAGAGCC ATCAACGTGC CTCCACGGGT GACCGTGCTC CCCAAGTCTC GGGTGGAGCT GGGCCAGCCC AACATCCTCA TCTGCATCGT GGACAACATC TTCCCCCTG TGATCAATAT CACCTGGCTG CGCAACGGCC AAAGTGTAC TGAGGGAGTG GCCCAGACCA GCTTCTATTG CCAGCCTGAC CATTGTGTTCC GCAAGTTCCA CTACCTGCCC TTCGTGCCCT CAGCCGAGGA CGTCTATGAC TGCCAGGTGG AGCACTGGGG CCTGGATGCG CCACTCCTCA GGCATTGGGA GCTCCAGGTG CCTATTCCAC CACCAGATGC CATGGAGACC CTGGTCTGTG CCCTGGGCCT GGCCATCGGC CTGGTGGGCT TCCTCGTGGG CACCGTCTC ATCATCATGG GCACATATGT GTCCAGTGTC CCCAGGTAAT GATCCTTCTG AGAGAAATGA CTTGTGGGAG ACACCCTGCA GATCCTCATG</p>	<p>13</p>

	<p>GGTTTGTGAC AGCCCCTGCG TGCTCAGTGC CCTTTAAGTG CATCCCGCTG TGCTGACTTT GAGTGGGATC AACATCTGTC CTACGGGTCC CCTCTTTTTT GGCCCCAGTA TTCATGGCAG GGTTTGTGG ACACCTACTA GCTTCCCTTC CCATTCAACA CACACACACA TTCTTGCTCT ACCCAAAGCT CTGGCTGGCA GCACTAAATG CTTTGGTGGT GTTTGCCTG TGTCCTTTCC AGGCCTTGGC CAGTTCTTCC AGGGGTGAGG CATGTGGTGC TGGGGATTGG CAGCCATCCT GGGGCCACA CAGGTGTGTC TTGCTCCATT TGGCCATTG TGTGTTACTT TGTGAATGAG CCATTTTACA TGGACTTCAT GAAATTTGCC TCCTGAGTTC AGGTTTACCC TGAAAGGGAT GCAGATTATC CTGTTCTCA CGACCCCTC AGCTAACAAAC AGTTCTGAAG GGTGCTGGGA CAGGACAGGC TCATGGGGAC TCCACTCCTG CCTGGGTTTA CTCTGTATGA AGAGGCCACT GGTATCCTGC CATGATGTTA TCTCTTTTTT CTACTTTTCC CTAGAGTCCC ATGCATGATA AAGAGAGGCC CAAGGCTTGG ATAAGGTGGC CACTTCCCTC AGTGGAGTCA GTCATGTTAG GTAGGAGGTG GTAGAGTCGG TCTGCGAGGT ATCTCGTAAG AGGGGAGGTC CACCTAGACA CACTCTAAAT ATGTGGCCTA GAAGATTTTG GTCTACTTTT CTGTGAACAG AATTTAAAAC ATACAAAGAG ATAAATCACC ATACCACATA GTTTATGTCA GGACCAAAAT GAGCAATACA GATTACGGTT TTCAAACCAG AATGCACATA AGAACTGCTT GGGATCCTTT TAAAAGTACA GGCATTGGCC TGGTGCAGTG GCTCATTCCCT GTAATCCCAG CACTTTGGGA GGCCAAGGGG ACAGGACTGC TTGAGGCCAA GAGGTGGAAA CCATCTTGGG CTACATAGAG AGACCCATC TCTACAAAGA AAGATTTAAA AATTAACCAG GCATGGTGGC TCGCACCTGT ATTCCCAGCC ACTGGGGAGG CTGAGGCCGG AGGAGTGCTT GAGCCCAGGA GTTCAAGGCT GCAGTGAGCC AAGATTGCGC CACTGCACTC CAGCCTAGGT GACAGAGTGA GACCCTGTCT CTAAATAAAT AAATAAATAA AATATAAAAA TAACAGTCAT CACCCAGACC TACTGAATTA GAATCTCGGG AGTGCAGGGG GCAGCAACAG GGAGGCTGTC TTTTCTGAGA TGGGGTCTCA CTCTGTCACC AGGCTGGAGT GCCATGGCAT GATCTCAGCT CACTGCAACC TCCACCTCCT GAGTTCAAGC CATTCTCCTG CCTCAGCCTC CTGAGTAGCT GGGACTACAG GTGTGCGCCA CTACACTCAG CTAATTTTTG TATTTAAGT AGAGACGGGG TTTCATCATG TTGGCCAGGA TGGCCTCCAT CTCTTGACCT CGTGATCCAC CCACCTTCCC TCCCAAAGTA CTGGAATTAC AGGCATTAGC CACTGTGCC AGCCGAGGCT GTCATTTTTA ACCGGCTCTG GATGACTCTG ATGCAGCCAT CCTGGACCTT GGCTGTGGTC TGGTAACTGG AACCCAGTGA CGTAATCAGG TGCCATCGGG</p>	
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		<p>GGTCATGGGA AAGGGGGATC CCCAAGGTCT GAGGTGGACT AGGAAGGCTT TCTGAAGAAC CTGGGTCTGT TAGGGCATCA GCCAATCAAG GTACAAGTAA ATAGAGGCAA AATGAGGGTT TGAAGTGTGA GCAGTTGGTC CTGGAAAAGA AAGAAACCAA GAGATTATGG GGACTIONAATG GGCTTCTTAA GAGAGAATAA GTTGAAATCA ATGACCAGAA GACCCTGATG GAAGTGGAGG AGAATCATCT CAGGCAAAC TTTTGTGTGC CAGTAACAGA AACCCCTCTT GTGTGATCAC ATGCAAAGTA TAGGATATTT GCAATATAGC CATGGGGAGG AGTGCAGGGC CCAAGGGTAG ATTTTAGCCA GGCCCTCCAG GAACAGAACT CGGATCCGAA AAGCCCAGAG AAGCTAGAGC TGCCCCTCCA AACTCTCGG ATCCACATGG TCTGTGTTCT CTAGACCCCC CTGCATGTTA GCGGTGTTCT CTCTCTGTGG ACTGACTGTC CTCTCAGTG AACATGTCCA CCCGACAGCT CCTGAGTTTA TATCATCTCA ACCCACAAA CCCACAGAGG CTGTGTCTCC TAGTCACAGC TTTAAATTAC TGGAAAATA AATGACTGGC CAAACTTGGA GCAGGTGTCC ATCCCAGCCC TGTGTAGTTA GAGCAGGAAT CAAGATCTCA ACACAAATGT GGCTGCCAAG CACTCAGCCC CGGGGCGAGG GGTCAAGTTC TTCTCAGAGA AAGAGGAATA AGTTGGTTCT CAGAAGACAT CACAAGATAC GTGTGTACCC ACAATCTCT GATCTCTGCT GATCTTTTGC TTAGACGTTA ACTTGATGCA TCATTGGAAA GGTGTTTCTC TCATCTCTGT CCTAAGGCTT GATAAAGTCA TTAAAATTGT GTTCTTTTGA CTAAA</p>	
<p>HLA- DRA</p>	<p>NM_0 1911. 4</p>	<p>TTTTAATGGT CAGACTCTAT TACACCCAC ATTCTCTTT CTTTTATTCT TGTCTGTTCT GCCTCACTCC CGAGCTCTAC TGACTCCCAA CAGAGCGCCC AAGAAGAAAA TGGCCATAAG TGGAGTCCCT GTGCTAGGAT TTTTCATCAT AGCTGTGCTG ATGAGCGCTC AGGAATCATG GGCTATCAA GAAGAACATG TGATCATCCA GGCCGAGTTC TATCTGAATC CTGACCAATC AGGCGAGTTT ATGTTTACT TTGATGGTGA TGAGATTTTC CATGTGGATA TGGCAAAGAA GGAGACGGTC TGGCGGCTTG AAGAATTTGG ACGATTTGCC AGCTTTGAGG CTCAAGGTGC ATTGGCCAAC ATAGCTGTGG ACAAAGCCAA CCTGGAAATC ATGACAAAGC GCTCCAATA TACTCCGATC ACCAATGTAC CTCCAGAGGT AACTGTGCTC ACAAACAGCC CTGTGGAAC GAGAGAGCCC AACGTCCTCA TCTGTTTCAT AGACAAGTTC ACCCACCAG TGGTCAATGT CACGTGGCTT CGAAATGGAA AACCTGTCAC CACAGGAGTG TCAGAGACAG TCTTCTGCC CAGGGAAGAC CACCTTTTCC GCAAGTTCCA CTATCTCCCC TTCCTGCCCT CAACTGAGGA CGTTTACGAC TGCAGGGTGG</p>	<p>14</p>

		<p>AGCACTGGGG CTTGGATGAG CCTCTTCTCA AGCACTGGGA GTTTGATGCT CCAAGCCCTC TCCAGAGAC TACAGAGAAC GTGGTGTGTG CCCTGGGCCT GACTGTGGGT CTGGTGGGCA TCATTATTGG GACCATCTTC ATCATCAAGG GATTGCGCAA AAGCAATGCA GCAGAACGCA GGGGGCCTCT GTAAGGCACA TGGAGGTGAT GGTGTTTCTT AGAGAGAAGA TCACTGAAGA AACTTCTGCT TTAATGGCTT TACAAAGCTG GCAATATTAC AATCCTTGAC CTCAGTGAAG GCAGTCATCT TCAGCATTTT CCAGCCCTAT AGCCACCCCA AGTGTGGATA TGCCTCTTCG ATTGCTCCGT ACTCTAACAT CTAGCTGGCT TCCCTGTCTA TTGCCTTTTC CTGTATCTAT TTTCCTCTAT TTCCTATCAT TTTATTATCA CCATGCAATG CCTCTGGAAT AAAACATACA GGAGTCTGTC TCTGCTATGG AATGCCCCAT GGGGCATCTC TTGTGTAAT ATTGTTTAAG GTTTCCTCAA ACTGTGATTT TTCTGAACAC AATAAACTAT TTTGATGATC TTGGGTGGAA AAAAAAAAAA AAAAAAAAAA AAAAAAAAAA AA</p>	
<p>HNRNP A3PI</p>	<p>NR_00 2726.2</p>	<p>CGAGTTGGAA GAGGTGAGTC CTGTCTCAA ATGGAGGTAA AACCGCCGCC TGGTTGCCCC CAGCCCGACT CCGGCAGTCG CCGTCGCCAC TGGGGGGAGG AGGGCCATGA TCCAAAGGAA CCAGAGCAGC TGAGAAAAC TTTTATTGGT GGTCTGAGCT TTGAAACTAC AGATGATAGT TTAAGAGAAC ATTTTGAGAA ATGGGGCACA CTCACAGATT GTCTGGTAAT GAGAGACCCC CAAACAAAAC GTTCCAGGGG CTTTGGTTTT GTGACTTATT CTTGTGTTAC AGAGGTGGAT GCAGCAATGC GTGCTCGACC ATTCAAGGTT GATGGGCGTG TAGTGGAAACC AAAGAGAGCT GTTTCTAGAG AGGATTCTGT GAAGCCTGGT GCCCATCTAA CAGTGAAGAA AATTTTGTG GGCAGTATTA AAGAAGATAC AGAAGAATAT AATTTGAGAG ACTACTTTGA AAAGTACGGC AAGATTGAAA CCATAGAAGT TATGGAAGAC AGGCAGAGTG GAAAAAAGAG AGGATTTGCT TCTGTAAC TTGATGATCA TGATACAGTT GATAAAATTG TTGTTTCAGAA ATACCACACT ATTAATGGGC ATAAC TGTA AGTGAAAAAG GCCCTTGCTA AACAAGTGAT GCAGCCGGCT GGATCACAGA GGGGTCGTGG AGGTGGATCT GGCAATTGTA TGGGTCACAG AGGAAACTTT GGAGGTGGTG GAGGTAATTT TGGCCGTGAT GGAAACTTTG GTGGAAGAGG AGGCTATGGT GGTGGAGGTG GTGGCAGCAG AGGTAGTTAT GGAGGAGGTG ATGTGGATAT AATGGATTAG GAGGTGATGG TGGCAACTAT GGCAGTGGTC CTGGTTATAG TAGTAGAGGC GGGTATGGTG GTGGTGGACC AGGATATGGA AACCAAGGTG GTGGATATGG TGGCGGTGTT GGAGGATATG ATGGTTACAA TGAAGGAGGA AATTTTGACG GTAGTAACTA</p>	<p>15</p>

	<p>TGGTGGTGGT GGGAACTATA ATGATTTTGG AAATTACAGT GGACAACAGC AATCAAATTA TGGACACATG AAAGGGGGCA GTTTTGGTGG AAGAAGCTCG GGCAGTCCCT ATGGTGGTGG TTATGGATCT GGTGGTGGAA GTGGTGGATA TGGTAGCAGA AGGTTCTAAA AACAGCAGGA AAGGGCTACA GTTCTTAGCA GGAGAGAGAG TGAGGAGTTG TCAGGAAAGC TACAGGTTAC TTTGAGACAG TCGTCCAAG TGCATTAGAG GAACTGTA AATCTGTCAC AGAAGGAACG ATGATCCATA ATCAGAAAAG TACTGCAGC TTAACAGGA AACCCCTTCTT GTTCAGGACT GTCATAGCCA CAGTTTGCAG GAAGTGCAGC TATCGATTAA TGCAATGTAG CGTCAATTAG ATGTACATTC CTCAGGTCTT TTATCTGTTG TAGCTTTGTC TTATTCITTT TCTTTTCATT ACATCAGGTA TATTGCCCTG TAAATTGTGG TAGTGGTACC AGGAATAAAA AATTAAGGAA TTTTAACTT TTCAATATCT GTGTAGTTCA GTTTTTCTAC ATTTTAGTAC AGAAACTTTA ACAAAATGCA GTTTGAAGG TGTTTCCTTG TGAGTTAACA AGTAAAGAAG ATCAATTGTT AATTACTATT TTGTATAAAT TTTGCTAAAG TTAACGTAA AGAAACACCT GCTGACTTGC AGTTTAAGGG GAATCTATTC TCCCCATTT CAAACCATGA TATGAATGGA CGCCGACATG TGGAGAGAAC AGATAATTTG TGTGTTTGCA ATGTGTGTTT TAGGTAAATA GGATTGGGTA TTAAATTAG CATTGTGAA TTTAATAGCA TTAAGATTAC CTTCAAATAA AAAAGTCTCA AAATTTCTTT TTGGTTTTTG TGCATTTTCT TTTAAAATGT AATCACATGA TTTTAGTGTG TTAGACTTGC TGAGTCCTAG CTGTGTTTAG AACATCTCCA TTCTACATTT ACCTTGGTCA AATTTGAACT GCTGCCATAG GTTTTGGGTG TAAAGAATGT TACTGCCCT CCATTTAAAT TCTGAAAAGG TATAGTGGAT GTTTTCCCTC TCCTACATTA GAAACCATTC TTA AAAACTT TTCAAATAT AGAACCATTA AGCCTGCTAT ATCTGAGCAA ATTAGTGGGT ACCTTTTTTC CTTTTTTAAA GCACAAGAGG CCCATAAATC TTGAGTTATT TGCATTAGTT TACATTTTTT GATACAACCT TTCAGACCAA GAGAATAAAA ATCATGCGTT ATTAACCCC TAGCTGGCTG GCATGCTTTC CTGTTTGTAC TGTATACATT TTGCTGGATG AAACCAAGGA TAGTTTAGGT ATAATTGTCC AAAATAACCT AACTGCAGCA GAAATGTAGG ACAGTTGCTT AGTACAGGCT TCTCACTTCC TACAGACCTG AATTCAAATT TGGATAGTCT GAGTTATTAA ATTCCCAAAG ACAAAGAACA CACTCTTATT TCTTGTGTAT ATTTCAACAT AAATCATGTT GTTACCAATT TGTTGGGAAG GCCCTGGTTG AGAAGAGTTT TAGATAATAA GGCTGTATAT ATATAGATAT ATATAGATAT ATACCAATGT CTATATATAG AGATATTTTA TATATATATA TACAGGTATA</p>	
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		TATATGTGTG TGTATATATA TAGGTATATA CATATATACA TATATATATA TATATATATG GATATATACC CATGTCTACT GTTTTGCTTC AGCTAGTGCT TACAATTTCA TTCAAGTCCT GAGTATGTGT CCTGCTGTTA CTCCTTCTTT GGTAGTTGAA CGTTGAATTC AAGTCTTTCC TTCTGTTTTA AGAAGTACTA AGCAAACAAG CAATAAAAAG GGAATGGCG CATGCTAGTG TTTGAATATG CTCTCTTGTG GCTCTAATTC TGTGCCTCCG TGCATTAATA TTTGGATGCA TGCAATGCCA GCATGGAAAT TGGCCT	
IL23A	NM_016584.2	AAAACAACAG GAAGCAGCTT ACAAACCTCGG TGAACAACCTG AGGGAACCAA ACCAGAGACG CGCTGAACAG AGAGAATCAG GCTCAAAGCA AGTGGAAGTG GGCAGAGATT CCACCAGGAC TGGTGCAAGG CGCAGAGCCA GCCAGATTTG AGAAGAAGGC AAAAAGATGC TGGGGAGCAG AGCTGTAATG CTGCTGTTGC TGCTGCCCTG GACAGCTCAG GGCAGAGCTG TGCCTGGGGG CAGCAGCCCT GCCTGGACTC AGTGCCAGCA GCTTTCACAG AAGCTCTGCA CACTGGCCTG GAGTGACAT CCACTAGTGG GACACATGGA TCTAAGAGAA GAGGGAGATG AAGAGACTAC AAATGATGTT CCCATATCC AGTGTGGAGA TGGCTGTGAC CCCCAAGGAC TCAGGGACAA CAGTCAGTTC TGCTTGCAA GGATCCACCA GGGTCTGATT TTTTATGAGA AGCTGCTAGG ATCGGATATT TTCACAGGGG AGCCTTCTCT GCTCCCTGAT AGCCCTGTGG GCCAGCTTCA TGCCCTCCCTA CTGGGCCTCA GCCAACTCCT GCAGCCTGAG GGTCACTACT GGGAGACTCA GCAGATTCCA AGCCTCAGTC CCAGCCAGCC ATGGCAGCGT CTCCTTCTCC GCTTCAAAT CCTTCGCAGC CTCCAGGCCT TTGTGGCTGT AGCCGCCCGG GTCTTTGCC ATGGAGCAGC AACCTGAGT CCCTAAAGGC AGCAGCTCAA GGATGGCACT CAGATCTCCA TGGCCAGCA AGGCCAAGAT AAATCTACCA CCCAGGCAC CTGTGAGCCA ACAGGTTAAT TAGTCCATTA ATTTAGTGG GACCTGCATA TGTTGAAAAT TACCAATACT GACTGACATG TGATGCTGAC CTATGATAAG GTTGAGTATT TATTAGATGG GAAGGGAAAT TTGGGGATTA TTTATCCTCC TGGGGACAGT TTGGGGAGGA TTATTTATTG TATTTATATT GAATTATGTA CTTTTTCAA TAAAGTCTTA TTTTTGTGGC TAAAAAAA	16
IQGAP 1	NM_03870.3	GGACCCCGGC AAGCCCGCGC ACTTGGCAGG AGCTGTAGCT ACCGCCGTCC GCGCCTCAA GGTTCACGG CTTCCTCAGC AGAGACTCGG GCTCGTCCGC CATGTCCGCC GCAGACGAGG TTGACGGGCT GGGCGTGGCC CGGCCGCACT ATGGCTCTGT	17

	<p>CCTGGATAAT GAAAGACTTA CTGCAGAGGA GATGGATGAA AGGAGACGTC AGAACGTGGC TTATGAGTAC CTTTGTCAAT TGGAAGAAGC GAAGAGGTGG ATGGAAGCAT GCCTAGGGGA AGATCTGCCT CCCACCACAG AACTGGAGGA GGGGCTTAGG AATGGGGTCT ACCTTGCCAA ACTGGGGAAC TTCTTCTCTC CCAAAGTAGT GTCCCTGAAA AAAATCTATG ATCGAGAACA GACCAGATAC AAGGCGACTG GCCTCCACTT TAGACACACT GATAATGTGA TTCAGTGGTT GAATGCCATG GATGAGATTG GATTGCCTAA GATTTTTTAC CCAGAAACTA CAGATATCTA TGATCGAAAG AACATGCCAA GATGTATCTA CTGTATCCAT GCACTCAGTT TGTACCTGTT CAAGCTAGGC CTGGCCCCCTC AGATTCAAGA CCTATATGGA AAGGTTGACT TCACAGAAGA AGAAATCAAC AACATGAAGA CTGAGTTGGA GAAGTATGGC ATCCAGATGC CTGCCTTTAG CAAGATTGGG GGCATCTTGG CTAATGAACT GTCAGTGGAT GAAGCCGCAT TACATGCTGC TGTTATTGCT ATTAATGAAG CTATTGACCG TAGAATTCCA GCCGACACAT TTGCAGCTTT GAAAAATCCG AATGCCATGC TTGTAAATCT TGAAGAGCCC TTGGCATCCA CTTACCAGGA TATACTTTAC CAGGCTAAGC AGGACAAAAT GACAAATGCT AAAACAGGA CAGAAAATC AGAGAGAGAA AGAGATGTTT ATGAGGAGCT GCTCACGCAA GCTGAAATTC AAGGCAATAT AAACAAAGTC AATACATTTT CTGCATTAGC AAATATCGAC CTGGCTTTAG AACAAGGAGA TGCCTGGCC TTGTTTCAGGG CTCTGCAGTC ACCAGCCCTG GGGCTTCGAG GACTGCAGCA ACAGAATAGC GACTGGTACT TGAAGCAGCT CCTGAGTGAT AACAGCAGA AGAGACAGAG TGGTCAGACT GACCCCTGC AGAAGGAGGA GCTGCAGTCT GGAGTGGATG CTGCAAACAG TGCTGCCCAG CAATATCAGA GAAGATTGGC AGCAGTAGCA CTGATTAATG CTGCAATCCA GAAGGGTGTT GCTGAGAAGA CTGTTTTGGA ACTGATGAAT CCCGAAGCCC AGCTGCCCCA GGTGTATCCA TTGCCGCCG ATCTCTATCA GAAGGAGCTG GCTACCCTGC AGCGACAAAG TCCTGAACAT AATCTCACCC ACCCAGAGCT CTCTGTCGCA GTGGAGATGT TGTCATCGGT GGCCCTGATC AACAGGGCAT TGAATCAGG AGATGTGAAT ACAGTGTGGA AGCAATTGAG CAGTTCAGTT ACTGGTCTTA CCAATATTGA GGAAGAAAAC TGTCAGAGGT ATCTCGATGA GTTGATGAAA CTGAAGGCTC AGGCACATGC AGAGAATAAT GAATTCATTA CATGGAATGA TATCCAAGCT TGCGTGGACC ATGTGAACCT GGTGGTGCAA GAGGAACATG AGAGGATTTT AGCCATTGGT TTAATTAATG AAGCCCTGGA TGAAGGTGAT GCCCAAAGA CTCTGCAGGC CCTACAGATT CCTGCAGCTA AACTTGAGGG AGTCCTTGCA GAAGTGGCCC AGCATTACCA</p>	
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	<p>AGACACGCTG ATTAGAGCGA AGAGAGAGAA AGCCCAGGAA ATCCAGGATG AGTCAGCTGT GTTATGGTTG GATGAAATTC AAGGTGGAAT CTGGCAGTCC AACAAAGACA CCCAAGAAGC ACAGAAGTTT GCCTTAGGAA TCTTTGCCAT TAATGAGGCA GTAGAAAGTG GTGATGTTGG CAAAACACTG AGTGCCCTTC GCTCCCCTGA TGTTGGCTTG TATGGAGTCA TCCCTGAGTG TGGTGAAACT TACCACAGTG ATCTTGCTGA AGCCAAGAAG AAAAAAGTGG CAGTAGGAGA TAATAACAGC AAGTGGGTGA AGCACTGGGT AAAAGGTGGA TATTATTATT ACCACAATCT GGAGACCAG GAAGGAGGAT GGGATGAACC TCCAAATTTT GTGCAAAATT CTATGCAGCT TTCTCGGGAG GAGATCCAGA GTTCTATCTC TGGGGTACT GCCGCATATA ACCGAGAACA GCTGTGGCTG GCCAATGAAG GCCTGATCAC CAGGCTGCAG GCTCGCTGCC GTGGATACTT AGTTCGACAG GAATTCCGAT CCAGGATGAA TTTCTGAAG AAACAAATCC CTGCCATCAC CTGCATTCAG TCACAGTGGA GAGGATACAA GCAGAAGAAG GCATATCAAG ATCGGTTAGC TTACCTGCGC TCCCACAAAG ATGAAGTTGT AAAGATTCAG TCCCTGGCAA GGATGCACCA AGCTCGAAAG CGCTATCGAG ATCGCCTGCA GTAATTCCGG GACCATATAA ATGACATTAT CAAAATCCAG GCTTTTATTC GGGCAAACAA AGCTCGGGAT GACTACAAGA CTCTCATCAA TGCTGAGGAT CCTCCTATGG TTGTGGTCCG AAAATTTGTC CACCTGCTGG ACCAAAGTGA CCAGGATTTT CAGGAGGAGC TTGACCTTAT GAAGATGCGG GAAGAGGTTA TCACCCTCAT TCGTTCTAAC CAGCAGCTGG AGAATGACCT CAATCTCATG GATATCAAAA TTGGACTGCT AGTGAAAAAT AAGATTACGT TGCAGGATGT GGTTTCCCAC AGTAAAAAAC TTACCAAAAA AAATAAGGAA CAGTTGTCTG ATATGATGAT GATAAATAAA CAGAAGGGAG GTCTCAAGGC TTTGAGCAAG GAGAAGAGAG AGAAGTTGGA AGCTTACCAG CACCTGTTTT ATTTATTGCA AACCAATCCC ACCTATCTGG CCAAGCTCAT TTTTCAGATG CCCAGAACA AGTCCACCAA GTTCATGGAC TCTGTAATCT TCACACTCTA CAACTACGCG TCCAACCAGC GAGAGGAGTA CCTGCTCCTG CGGCTCTTTA AGACAGCACT CCAAGAGGAA ATCAAGTCGA AGGTAGATCA GATTCAAGAG ATTGTGACAG GAAATCCTAC GGTTATTA AAA ATGGTTGTAA GTTTCAACCG TGGTGCCCGT GGCCAGAATG CCCTGAGACA GATCTTGCC CCAGTCGTGA AGGAAATTAT GGATGACAAA TCTCTCAACA TCAAAACTGA CCCTGTGGAT ATTTACAAAT CTTGGGTAA TCAGATGGAG TCTCAGACAG GAGAGGCAAG CAAACTGCC TATGATGTGA CCCCTGAGCA GCGCTAGCT CATGAAGAAG TGAAGACACG GCTAGACAGC TCCATCAGGA ACATGCGGGC</p>	
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	<p>TGTGACAGAC AAGTTTCTCT CAGCCATTGT CAGCTCTGTG GACAAAATCC CTTATGGGAT GCGCTTCATT GCCAAAGTGC TGAAGGACTC GTTGCATGAG AAGTTCCTTG ATGCTGGTGA GGATGAGCTG CTGAAGATTA TTGGTAACTT GCTTTATTAT CGATACATGA ATCCAGCCAT TGTTGCTCCT GATGCCTTTG ACATCATTGA CCTGTCAGCA GGAGGCCAGC TTACCACAGA CCAACGCCGA AATCTGGGCT CCATTGCAAA AATGCTTCAG CATGCTGCTT CCAATAAGAT GTTTCCTGGGA GATAATGCC ACTTAAGCAT CATTAAATGAA TATCTTTCCC AGTCCTACCA GAAATTCAGA CGGTTTTTCC AAACCTGCTTG TGATGTCCCA GAGCTTCAGG ATAAATTTAA TGTGGATGAG TACTCTGATT TAGTAACCCT CACCAAACCA GTAATCTACA TTTCCATTGG TGAAATCATC AACACCCACA CTCTCCTGTT GGATCACCAG GATGCCATTG CTCCGGAGCA CAATGATCCA ATCCACGAAC TGCTGGACGA CCTCGGCGAG GTGCCACCA TCGAGTCCTT GATAGGGGAA AGCTCTGGCA ATTTAAATGA CCCAAATAAG GAGGCACTGG CTAAGACGGA AGTGTCTCTC ACCCTGACCA ACAAGTTCGA CGTGCCTGGA GATGAGAATG CAGAAATGGA TGCTCGAACC ATCTTACTGA ATACAAAACG TTTAATTGTG GATGTCATCC GGTTCCAGCC AGGAGAGACC TTGACTGAAA TCCTAGAAAC ACCAGCCACC AGTGAACAGG AAGCAGAACA TCAGAGAGCC ATGCAGAGAC GTGCTATCCG TGATGCCAAA ACACCTGACA AGATGAAAAA GTCAAAATCT GTAAAGGAAG ACAGCAACCT CACTCTTCAA GAGAAGAAAG AGAAGATCCA GACAGGTTTA AAGAAGCTAA CAGAGCTTGG AACCGTGGAC CCAAAGAACA AATACCAGGA ACTGATCAAC GACATTGCCA GGGATATTCG GAATCAGCGG AGGTACCGAC AGAGGAGAAA GGCCGAACTA GTGAAACTGC AACAGACATA CGCTGCTCTG AACTCTAAGG CCACCTTTTA TGGGGAGCAG GTGGATTACT ATAAAAGCTA TATCAAAACC TGCTTGGATA ACTTAGCCAG CAAGGGCAAA GTCTCCAAAAGCCTAGGGA AATGAAAGGA AAGAAAAGCA AAAAGATTTC TCTGAAATAT ACAGCAGCAA GACTACATGA AAAAGGAGTT CTCTGGAAA TTGAGGACCT GCAAGTGAAT CAGTTTAAAA ATGTTATATT TGAAATCAGT CCAACAGAAG AAGTTGGAGA CTTCGAAGTG AAAGCCAAT TCATGGGAGT TCAAATGGAG ACTTTTATGT TACATTATCA GGACCTGCTG CAGCTACAGT ATGAAGGAGT TGCAGTCATG AAATTATTTG ATAGAGCTAA AGTAAATGTC AACCTCCTGA TCTTCCTTCT CAACAAAAG TTCTACGGGA AGTAATTGAT CGTTTGCTGC CAGCCCAGAA GGATGAAGGA AAGAAGCACC TCACAGCTCC TTTCTAGGTC CTTCCTTCTT CATTGGAAGC AAAGACCTAG CCAACAACAG CACCTCAATC TGATACACTC</p>	
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	<p>CCGATGCCAC ATTTTAACT CCTCTCGCTC TGATGGGACA TTTGTTACCC TTTTTCATA GTGAAATTGT GTTTCAGGCT TAGTCTGACC TTTCTGGTTT CTTCATTTTC TTCCATTA TAGGAAAGAG TGGAACTCC ACTAAAATTT CTCTGTGTTG TTACAGTCTT AGAGGTTGCA GTECTATATT GTAAGCTTTG GTGTTTGTTT AATTAGCAAT AGGGATGGTA GGATTCAAAT GTGTGTCATT TAGAAGTGGA AGCTATTAGC ACCAATGACA TAAATACATA CAAGACACAC AACTAAAATG TCATGTTATT AACAGTTATT AGGTTGTCAT TFAAAAATAA AGTTCCTTTA TATTTCTGTC CCATCAGGAA AACTGAAGGA TATGGGGAAT CATTGGTTAT CTTCATTGT GTTTTICTTT ATGGACAGGA GCTAATGGAA GTGACAGTCA TGTTCAAAGG AAGCATTCT AGAAAAAGG AGATAATGTT TTAAATTTT ATTATCAAAC TTGGGCAATT CTGTTTGTGT AACTCCCCGA CTAGTGGATG GGAGAGTCCC ATTGCTAAAA TTCAGCTACT CAGATAAATT CAGAATGGGT CAAGGCACCT GCCTGTTTTT GTTGGTGCAC AGAGATTGAC TTGATTCAGA GAGACAATTC ACTCCATCCC TATGGCAGAG GAATGGGTTA GCCCTAATGT AGAATGTCAT TGTTTTTAAA ACTGTTTTAT ATCTTAAGAG TGCCTTATTA AAGTATAGAT GTATGTCTTA AAATGTGGGT GATAGGAATT TTAAAGATTT ATATAATGCA TCAAAGCCT TAGAATAAGA AAAGCTTTTT TTAAATTGCT TTATCTGTAT ATCTGAACTC TTGAACTTA TAGCTAAAAC ACTAGGATTT ATCTGCAGTG TTCAGGGAGA TAATTCTGCC TTTAATTGTC TAAAACAAAA ACAAAACCAG CCAACCTATG TTACACGTGA GATTAAAACC AATTTTTTCC CCATTTTTTC TCCTTTTTTC TCTTGCTGCC CACATTGTGC CTTTATTTTA TGAGCCCCAG TTTTCTGGGC TTAGTTTAAA AAAAAAATCA AGTCTAAACA TTGCATTTAG AAAGCTTTTG TTCTTGATA AAAAGTCATA CACTTTAAAA AAAAAAAAAA CTTTTCCAG GAAAATATAT TGAAATCATG CTGCTGAGCC TCTATTTTCT TTCTTIGATG TTTTGATTCA GTATICTTTT ATCATAAATT TTTAGCATTT AAAAATTCAC TGATGTACAT TAAGCCAATA AACTGCTTTA ATGAATAACA AACTATGTAG TGTGTCCCTA TTATAAATGC ATTGAGAGAAG TATTTTTATG AGACTCTTTA CTCAGGTGCA TGGTTACAGC CCACAGGGAG GCATGGAGTG CCATGGAAGG ATTCGCCACT ACCCAGACCT TGTTTTTTGT TGTATTTTGG AAGACAGGTT TTTTAAAGAA ACATTTTCTT CAGATTAATA GATGATGCTA TTACAACCTAG CATTGCCTCA AAAACTGGGA CCAACCAAAG TGTGTCAACC CTGTTTCTT AAAAGAGGCT ATGAATCCCA AAGGCCACAT CCAAGACAGG CAATAATGAG CAGAGTTTAC AGCTCCTTTA ATAAAATGTG TCAGTAATTT TAAGGTTTAT</p>	
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		<p>AGTTCCCTCA ACACAATTGC TAATGCAGAA TAGTGTA AAA TGCGCTTCAA GAATGTTGAT GATGATGATA TAGAATTGTG GCTTTAGTAG CACAGAGGAT GCCCAACAA ACTCATGGCG TTGAAACCAC ACAGTTCTCA TTAGTGTTAT TTATTAGCTG TAGCATTCTC TGTCTCCTCT CTCTCCTCCT TTGACCTTCT CCTCGACCAG CCATCATGAC ATTTACCATG AATTTACTTC CTCCAAGAG TTTGGACTGC CCGTCAGATT GTTGCTGCAC ATAGTTGCCT TIGTATCTCT GTATGAAATA AAAGGTCATT TGTTTCATGTT AAAAAAAAA</p>	
LOC49 4127	NR_03 6691.1	<p>GCAGTGCCCG ACTCCGCAGG AGCGCCAGGG CGGCTCCTGC TCTTCCTGGA CTCCCTGAAG AGGCGTTTGT CGAAATGTCC ACAGAAGGAG GATTTGGTGG TACTAGCCGC AGTGATGCCC AGCAAAGCCT AAAGTCCTTC TGGCTTCGGG TCATGGAAGA AATCTGGAAT TTAGCAGTGA AAGATTTCTG AATGCAGGAA CTCCCACTGG CTCGTATTAA GAAGATTATG AAAGTGGATG AAGATGTGAA GATGATCAGT GCAGAGGCC CTGTGCTCTT TGCCAGGGCA GCCCAGATTT TTTATCACAG AGTTGACTCT TCGAGCCTGG ATTCACACAG AGGATAACAA CTGCCGGACT TATGTCGCCA TGGCAATTAC GAAATTTGAT CAATTGGATT TTCTCATCGA TATTGTTCTA AGAGATGAAC TGAAACCTCC AAAGTGTCAG GAGGAGGTGC TGCAGTCTGT AACTCCTGCT GAGCCAGTCC AATACTATTT CACGCTGGCT CAGCAGCCCA CCGCCCCTCC AAGTCCAGGG ACAGCAGCAA GGCCAGACCA CCGCCAGCTC CATGACCACC ATGCAGCCTG GGCAGATCAT CATCGCACAG CTTCAGCAGG GCCAGACCAC GCCCGTGACG ACGCAGGTTG GAGAAGGTCA GCAGGTGCAG ATTGTCCAGG CCCAGCCACA GGGTCAAGCC CAGTAGGCC AGAGTGGTAC TGGATGGACC GTGCAGGTGA TGTAGCAGAT CCTACTAAC ACAGGAGAGA TCCAGCAGAT CCCGGTGCAG CTGAAGGCTG GCCAGCTGCA GTGTATCCGC TTAGCCCAGT CTGTATCAGG CACCCACGTT CTGCAGGGAC AGATCCAGAC ACTTGCCACC AGCGCTCAAC CGATTACACA GACAGAGGTC CAGCAAAGAC AGCAGTAGTT CAGCCAGTTC ACAGATGGAC AGCAGCTCTA CCAGATCCAG CAAGTATCCA TACCTGCGGG CCAGGACCTG CCCAGCCCAT GTTTTTCCAG TCAGTCAACC AGCCCTCTGA TGGGCAGGCC CCCCAGGTGA CTGGCGGCTG AGGGCCGGAG CTGGCAAGGC CGAGGACACT CAACACAATT TTGCGGTAC AGCCCCAGGT CATGAACACA GCCTTCTTCC CCAGAGGACC CGGCCGACCT CAGCTCCTCC TGCAGGCTAG GACAATGGCG CACTAGGCCT CATGCCTGGG GGCCGAGATT CTCCAACAGA AAGATGCAAT ATTTTTTGT TCTTTTTTTC TCCAAGGAAT</p>	18

		CAATATTTCA ATATGTTGAG CTGTGTGTCC AATGCTATGA AATTAAAATA TTAAATCACA AAAAAAAAAA AAAA	
LOC64 6471	NR_02 4498.1	AATGAATCCA ACTACTGTCC TTGTCCTCTC CTCTCTAGGT TTGTGGAGAC TGAATTCCAC AAACCTAGGG ACAGGGCACT CTCTAGGGGC AAGTCAACTC TCAATTATAG TGAGGGCAGG TTCCCCAGTT GCCAGCCTAC ACCCTGGCCA GCCACCCAAG GGAATACCTG CTGCTGCTAA GGCAGTCAAT GTTGGGAGGG TCAGGGAAGG GGAGAGGAAG TAGCTGAGTG TAGAGATTAT CCAGGCTTTC CTTCCCGTCC TCTGTACAGA AAGGCAGACA TACTACTGACT CCTGAAGTGC CCCAGGATCA TAGTTGGTTC TTCTTAGGGG GAGGGAGTGA AACGGTGGCC TTGCTCCAAT TCCAGGCCTC TGGAGAAAGG AGTGCTCAAC TGGAGAGTCT CAGAACCTTT TACAGTACTT CAGCAGGCCC AGCACCCAAC CTTCCCACC ATACCCCTGC CCTGGGAATG CCCCTTACCA CAGTGGCTTC CATTCTGAC ATTTGAACCA GCTCCTGCAC TTGAAGGACA AGGCAGCCAC TGGGCAGCTC CGGGAGGCGG GAGGCGTGAC TCCTTCATAT CAGGCCATCT GGAGGAGCAC CAGATTTCCC TCTTGTA AAA CAACCCCCA CAGTGGGGAT TTAATGGTCA CCACACAAGC CCCAAAAAAC CAGGCCCTGC AGAGCCATGA CTTACAGGG GAGAAAACCA GCCCAAGAGG CTGCCCTCAC CCAACACTTA GCTCTAAGTC CAGACTCTTA AAGGTTCTGT TAAAGATCTG CGGCCAGGCG CAGTGGCTCA CATCTGTAAT CCCAGCACTT TGGGAGGCCG AGGTGGGCGG ATCACGAGGT CAGGAGAGCG AGACCATCCT GGCTAACACA GTGAAGCCCT GTCTCTACTA AAAATACAAA AAATTAGCCA GGTGTGGTGG CGGGCACCTG TAGTACCAGC TACTTGGGAG GCTGAGGCAG GAGAATGGCG TGAACCCAGG AGGCAGAGCT TGCAGTGAGC CGAGATCACG CCACTGCACT CCAACCTGGG TGACAGAGCG AGACTCCGTC TCAAAAAAAAA AAAAAAAAAA AAAAGATTTG CCTAGTGAGC TTGGCCAACC CAAGAGCCTT CCTGGGCTCT CAGGTTTCTT GAGGCAAAAC TTCTAAAGAT GCAAATAGAA TCCTTGGTAG CTACAAAGCT CATCTAGCCA AATTATAAGT ACTATCCAAG CCCAGGATTC ATTCTTGAAG ATGAGATGTA AAAGCACTGC TGGGCCCTTG CAAAGGGAGA GATCAGAGAA AATAGGAGAA CGAACACCAG CTCATGCCAA TCTCAGTCTC ATGAGCTACA GAGAAAAAGA GCCAAAAAAG TTTATCCTGG CTCAGCCACC TTCTGGCTTT GTAGCTACAG GCAAGTTATT TAACTTTTCT GAATGTTGGT TTTTCCATCT GGAAAAAGGA GACAGGAAA GCTGCCACTG GAGGCTACCT TAAGGGTCAA TGAGTGAAAT GGGGTGAGT GCTTAATGAA CTGTTAAAAC ATGACACAGA TGGGTGAGGT GACATCTGAC	19

	<p>TGCTCTTCAC AGCAGTCCTT GCATGGTGAG CAGATGCAAT CAGTGTGCTC GTTCATTAC TTGTTTCATCA AACACCAACA GTACCCACTT GTGCCAGGCC TTGTGCTGGG CACCGTAGGG GCTATGGAAG GAGTAAGACA CAGTCATGCT CTCAAGGAAC TCGCTGTCCA GCAGAGTGAT AGGACATGAA TACATACTTC TAGCAAAGGC TGAGTAGCAT TAGGCAGGTG GCCAGATGAG GTCAGAGCTC TGGCCCTAGT GCCCTGTAGT AAGTTGCGAG GGAGTAGGAA ATGAAGACAA AACAGCAAGA GCACAGGTCC AGCACAAGAA GGTGAAGGAC ATGGAAAGAG GAGCAGAGCA TGATAAAGCT ACAAAGTAG GGGGCAAGTC ATGAGGTCCG GGTCACTGGG GAGCCCTGGA CAGGCTCTGA ATGGGAGATA GACTAGACTG ACAAAGCAGC TCTTTCTGGA GACCTCCTGT GGAGGGTCCC CTCTTCTACC CTTGGGAGAC GAGGTGTCAT TCACTTCACG ATACACATGG GGAAAGGAAG TTGGACAGGG GAAGGCACTT GTCTAAGGTC ACTCAGTGGT CATGCCAGTG CAAAGCACCC AAAGCCTGGT CCAGGTCCCTG CCACCTCCTG CGTGGGCTCC ATCTTCTCGG CTCAGTTCAA GCCTTTATTA TCAGAGTCCC TCTGTGTGCC AGGCTCTGTG CTGAATGTCC GTGTGCGCCA CAAATTGCAC TAACGGTGCT TACTGCCACT TCCTCTCCGA CCTTACCTCC AAAGACCCAG CCGGGCCTAA CCCTCACACA CAGCCAGAGG AATCCTTCCC AAACACAGTT AATCACATCG CTCATCCGTT TAAAACTGGC TTAGTAGAGC CAACACAGGA TAAAATCCAC AATCAGTCCC AAACCCATCT TTCTCTGCCG TCTCTAGACC ACATTCACTG AGCACCAAGG CCTGCACGCT TCGCAGCCAG CCCTCCTGCT CTCCCCCTTG CCCTCCAGCC AACTGGCCC TCAACCCCA AACTGGGCC CACCTCTGAG CCCTGCATCA AACACCAGCA CCCCCTTCG CGAGCTCATT CTTCATCATT CCTCAGAGCT CAGCTTAAGC ATCACCTTCC TTCCCTGCGA CAGGGTCAGG TCCTTTCTG TCGGCTCCGC AGCACCTGC CTGTCCCTCG GAGTCCTCG GAGAGGAACT AGATAACTGT GATCCCCGCG CACAGCACGG TGCTGCCTA GGACACAGTG AGCGCTTCAA CGCGCTGAAC TAAACGAATG AAGGGGAACA CCAACTCAGC TGTCACCCCC GCCCCAAGT GTCAGTGGCT AGAAGATCTT CCCCCGGTTC GCCGGCAGCA CCGGTGGGTC TTGTCACCTC TGCGCCCCGC AGCCCTCCGG AGGCTGCAGT TGCTGTCTGC TGACAGGCGT CCCCTCAGGA GAACGGGAGC TCCTCCAGCG CAGGGCCCTG CTCAGCACCC CTGGTCCCGG GGGCCCCAC GGAGGCCGCC GCGCTCAGTG AAACCTTGCC GCGCGCAGCA GTGGCCGGAG ACCCGCGCGG ACCCTCTCCC CGCGGGGACC CGCCAGGGCG AGTCGCCCT CCCCCGCGG AGACGACCCG ACCTGGGCCA AGGTCGGCCA AGTGCTCGCC GCCTGCCAG CGGCCCTTGG</p>	
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		ACCCACAGGA CCCGGACCCG AGTGCGCGGA AGCCTCCGAG CCGCGGCCAG GTCTCAGAAT GCGCGGCGGG GCAGCCCGGC CCGTCAAGCA GCGAAGCCGA AACTAGAAGC CTGAGGTCTG GCTTGGTAC TGTTTGCTGG GGA ACTCCGA CGAGCAACTA CCCTTCGGGG ACTCAGTTT CCCC GGCCA AGTGAGACGG ACAGCTGCGG CTCCC GCGAG AACTCGCGCG GGAACAGGAG CTGCGGGCTC CGCACAGCGC CCGGCACGTA GACCCAGTCT ACTAACGGGC TGGAAGGTCG CGCTCCCGCC TCCGCGCCTC CCTAGAACCT TTCTCACCTC AAGCTCGGGC TCCCGGACTC TCTCACCTCA CGGCCCTCC TTCCACCTTG GGCCGCCAG CCTCAACCGC CCT	
LOH12 CR	NR_02 4061.1	GGCTACGCAA GACCTTCAGT TCCGGATTAG GAGGCCCGC CCCCGGCCC GAGGGAGGGG CGGAGAGACC CGCTCCTGCG ACTTAGGGCG ATGCCACCTT AAAGGGCTTG ACCTCCTCGA AGCCAGA ACT GCGGAAGAGG ATGGAGAAAG AAACGCTAGA TAGACCCAGG ATTCGAACCC ACAACAGCTT GGATGCAAAG CTCATTTTGA ATTCTGAAGA GCTGGGAGTT TAGCAGTGGA CGACCAAACA AAAAAATACC AGAAGACGGT GAAAGCAGCA GCTGGAAATG AGAGAAAAAT TAAGAAATAG AAAATGCTGT TTCATGTATT AGGGAGTGCA CATCTTTAAA GAGAAGTGAA GAACCTTTTT GGCTAGGCGC TTTCTGGGGA CTTGGCAGT TATCAACAAG TCACTCTGAA ACTAACAGAA GGTGAGGAGG GAGTAATCCA GAAAGGAAAC TGCCATTTTT GCTAGGGCAA GAAAGTAGAC CTAGCAATGC AATGCCATCA AGGAGTTAGC TGGCTCTAGC GTTCTCCGAA CTTTGCAACT CATTITATAT TACATTGTCT GCGTCAAGAA ATTTCAAGTA AATGCCTTGA GATTTTATAT GTATAAAATG TAGTCTCTGG CCAGGCGGGG TGGCTCACGC CTGTAATCCC AGCACTTTGG GAGGCTGAGG CGAGTAGATC ACGATGTCAG GAGATCAAGA CCATCCTGGC TAACATGGTG AAACCCCGTC TCTACTAAAA AATACAAAAA ATTAGCCGGG CATGATGGTG GGCGCCTGTA GTCCCAGCTA CTCGGGAGGC TGAGACAGGA GAATGGTG TG AACCCGGGAG GCAGAGCTTG CAGTGAGCCG AGATCGCGCC ACTGCACTCC AGCCTGGGCG ACAGAGCGAG ACTCCGTCTC AAAAAAAAAA AAAAAAAAAA AAGTCTCAGA AGAAGTGTC AAGTTGACTAA ATGATTTTTA AGTCCCTTCC ACTCTACTAT ATCACAATAA TTTCCCCTAT TTTACCTTAT TTAATTTTCA CAACAACCCT GCAGCCTGTA AGTAGGTGAC TCCTCCATGG TCAGCAGGTG GTCAGTGGTC GAACTATTTA GGA CTATCTG ATGCTCTTTC CTCTGCTCAC CATTGCAATA CAATGAGAAA AATGCAAATG AATCTAAGTA CAGTGTTCTC TGGGAATATG AAGATAGAAC	20

		<p>TAAAACTGC CTGAGAGATC AGGGCAGGCT TCTAGAAAGT TGTGTCTAAC TAGTCTTGAG GTTTACATAA ACCACACCAG GCTGATGGGA CTGGGAAGGG CCCACTAAGC AGTGAGACCA TTCCCTTTTG GAGAGTCCTT GCATCCCCTC CCAGATTTCC TCTTTGAGGG GAAGGTGAGA GAGGAGGTAA AAGGGGTGAG GAATGGAGAA TAACTCATTC TGGTTCTTGT TTCCCTTTT CCACATAAAA GTATATTTGT CTTGTGTTCC ATACACCAGT CCATACTGAT GTGATGGTGT TTTTATGCT TCCTTTTGAA TAAACATTC ATTCTTAA</p>	
PBXIP1	NM_020524.3	<p>GGTCAGTTTC TGGTCACATG ATTTTCTTCT CGGGCTGCAA ACAAAGGGAA GCCTGCAACA AGTTAAGCTG AAGACCGAAG CAAGAGCTGG TTCAGGTGGC AGCCACAGCA GCCTCAGGGA CCTCAGCAAC TATGGCCTCC TGCCCAGACT CTGATAATAG CTGGGTGCTT GCTGGCTCCG AGAGCCTGCC AGTGGAGACA CTGGGCCCGG CATCCAGGAT GGACCCAGAA TCTGAGAGAG CCCTGCAGGC CCCTCACAGC CCCTCCAAGA CAGATGGGAA AGAATTAGCT GGGACCATGG ATGGAGAAGG GACGCTCTC CAGACTGAAA GCCCTCAGTC TGGCAGCATT CTAACAGAGG AGACTGAGGT CAAGGGCACC CTGGAAGGTG ATGTTTGTGG TGTGGAGCCT CCTGGCCAG GAGACACAGT AGTCCAGGGA GACCTGCAGG AGACCACCGT GGTGACAGGC CTGGGACCAG ACACACAGGA CCTGGAAGGC CAGAGCCCTC CACAGAGCCT GCCTTCAACC CCCAAAGCAG CTTGGATCAG GGAGGAGGGC CGCTGCTCCA GCAGTGACGA TGACACCGAC GTGGACATGG AGGGTCTGCG GAGACGGCGG GGCCGGGAGG CCGGCCACC TCAGCCCATG GTGCCCTGG CTGTGGAGAA CCAGGCTGGG GGTGAGGGTG CAGGCGGGGA GCTGGGCATC TCCCTCAACA TGTGCCTCCT TGGGGCCCTG GTTCTGCTTG GCCTGGGGGT CCTCCTCTTC TCAGGTGGCC TCTCAGAGTC TGAGACTGGG CCCATGGAGG AAGTGGAGCG GCAGGTCTC CCAGACCCCG AGGTGCTGGA AGCTGTGGGG GACAGGCAGG ATGGGCTAAG GGAACAGCTG CAGGCCCCAG TGCCTCCTGA CAGTGTCCCC AGCCTGCAA ACATGGGTCT TCTGCTGGAC AAGCTGGCCA AGGAGAACCA GGACATCCGG CTGCTGCAGG CCCAGCTGCA GGCCAAAAG GAAGAGCTTC AGAGCCTGAT GCACCAGCCC AAAGGGCTAG AGGAGGAGAA TGCCCAGCTC CGGGGGGCTC TGCAGCAGGG CGAAGCCTTC CAGCGGGCTC TGGAGTCAGA GCTGCAGCAG CTGCGGGCCC GGCTCCAGGG GCTGGAGGCC GACTGTGTCC GGGGCCAGA TGGGGTGTGC CTCAGTGGGG GTAGAGGCC ACAGGGTGAC AAGGCCATCA GGGAGCAAGG CCCAGGGAG CAGGAGCCAG AACTCAGCTT CCTGAAGCAG</p>	21

	<p>AAGGAACAGC TGGAGGCTGA GGCACAGGCA TTAAGGCAAG AGTTAGAGAG GCAGCGACGG CTGCTGGGGT CTGTACAGCA GGATCTGGAG AGGAGCTTGC AGGATGCCAG CCGCGGGGAC CCAGCTCATG CTGGCTTGGC TGAGCTGGGC CACAGATTGG CCCAGAAACT GCAGGGCCTG GAGAACTGGG GCCAGGACCC TGGGGTCTCT GCCAATGCCT CAAAGGCCTG GCACCAGAAG TCCCACCTCC AGAATTCTAG GGAGTGGAGT GGAAAGGAAA AGTGGTGGGA TGGGCAGAGA GACCGGAAGG CTGAGCACTG GAAACATAAG AAGGAAGAAT CTGGCCGGGA AAGGAAGAAG AACTGGGGAG GTCAGGAGGA CAGGGAGCCA GCAGGAAGGT GGAAGGAGGG CAGGCCAAGG GTGGAGGAGT CGGGGAGCAA GAAGGAGGGC AAGCGACAGG GCCCGAAGGA ACCCCAAGG AAAAGTGGTA GCTTCCACTC CTCTGGAGAA AAGCAGAAGC AACCTCGGTG GAGGGAAGGG ACTAAGGACA GCCATGACCC CCTGCCATCC TGGGCAGAGC TGTTGAGGCC CAAGTACCGG GCACCCCAGG GCTGCTCAGG TGTGGACGAG TGTGCCCGGC AGGAGGGCCT GACTTTCTTT GGCACAGAGC TAGCCCCAGT GCGGCAACAG GAGCTGGCCT CTCTGCTAAG AACATACTTG GCACGGCTGC CCTGGGCTGG GCAGCTGACC AAGGAGCTAC CCCTCTCACC TGCTTTCTTT GGTGAGGATG GCATCTTCCG TCATGACCGC CTCCGCTTCC GGGATTTTGT GGATGCCCTG GAGGACAGCT TGGAGGAGGT GGCTGTGCAA CAGACAGGTG ATGATGATGA AGTAGATGAC TTTGAGGACT TCATCTTCAG CCACTTCTTT GGAGACAAAG CACTGAAGAA GAGGTCAGGG AAGAAGGACA AGCACTCACA GAGCCCAAGA GCTGCGGGGC CCAGGGAGGG GCACAGCCAT AGCCACCACC ACCACCACCG GGGCTGACAC CCTGCCCCAC AGGGAATGGC CTTGGCCTGG CCCAGCCAA GATCCCAGCG TTATCTAACT CCTGGAGGGT GGACTCTGTC CTGGCTTGTT TGGTGTCCCTC AGATATCTTT CACACAGTAG AGCAAAATCA CCAGCCCTGC ACTGATGTCA CTTTATGTAG AAAAAGGCCT TAGCTGGACC TGC GTTGCCG TCTATGCAA TGCA TGCAA TACTCCAGGC CCTGGGATGT GGGCTTGTGT TTTGTCACTG TGAAGGGGGA GATGGGAGAG GAGCCTGTTT TGGGGTGGGG TCTGGGGAAG GCAATCTGAT TCTGAAGCTA AAGAGCTTTC ATCCTCTTGA GTGTATGTCC CCATAGTGGG CCCCTTGACC CACATGCTGA CCGGTGCCCT GGGATTTGAC TAGAGTTGCT GGCTCGAGGC CCAGCACGAG GACTTACCCT GGGGTTTTGT TAGGTTTGGGA AGCAGCTGTC CCTAGGGGGT GAAGTCCCCC CCCTTTTTTT TTTTACCCT GCTTCTCCA CGGCTTACC TCCCTATGTG AACTGTAGAC TCAGATCCA ATAAAGTGCT GTTGCAGCTA TGATGCTAGG TGGTTTCTAA GCACAGGGGA CACCCACAC CCCCTGCCTG</p>	
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		<p>AATGGATGGG TCCATCCCAG GCACTGGTAC TTGCCCCCTT GTTCTGTATC CCCCTTTGCC CTTGCCTTGC CCTTCCAACA AACCCTAGGC CTTGAGAAG CTGATACTTC TCCTTTTGCT CACAGCTGCC TTGGCCCCAC CCCTGGGAGA TGTAGCAAAT TGAGTGTGGG TTTTGGAGTC TGAGCCTCAG GCTCAAATCC AGGCCAAGTG ATCTTGGGCA AGTTAATCTC TGGGAACTTT GGGTTTCTTA TCCTCAAAA AGGCGATGGA AGGGCTGGGG AAGTGATTAA ATAAAAGCAA CGCAAGAAAA AAAAAAAAAA AAAAA</p>	
<p>RNF5</p>	<p>NM_0 06913. 3</p>	<p>AATAGTGATT AGGAAACCTT GAAGCCTGCC CAACGATCGT GGGCAGGAGG TGGTTTCTGG TTTGTTGGGG CGTGTGTATG TGTATTTGGG GGGACTGAAG GGTACGTGGG GCGAAACAAA ACCGGCCATG GCAGCAGCGG AGGAGGAGGA CGGGGGCCCC GAAGGGCCAA ATCGCGAGCG GGGCGGGGCG GGC GCGACCT TCGAATGTAA TATATGTTT GAGACTGCTC GGGAAGCTGT GGTCAGTGTG TGTGGCCACC TGTACTGTTG GCCATGTCTT CATCAGTGGC TGGAGACACG GCCAGAACGG CAAGAGTGTC CAGTATGTAA AGCTGGGATC AGCAGAGAGA AGGTTGTCCC GCTTTATGGG CGAGGGAGCC AGAAGCCCCA GGATCCCAGA TTA AAAACTC CACCCCGCCC CCAGGGCCAG AGACCAGCTC CGGAGAGCAG AGGGGGATTC CAGCCATTG GTGATACCGG GGGCTTCCAC TTCTCATTG GTGTTGGTGC TTTTCCCTTT GGCTTTTTCA CCACCGTCTT CAATGCCCAT GAGCCTTTCC GCCGGGTAC AGGTGTGGAT CTGGGACAGG GTCACCCAGC CTCCAGCTGG CAGGATTCCT TCTTCTGTT TCTCGCCATC TTCTTCTTT TTTGGCTGCT CAGTATTGA GCTATGCTG CTTCCTGCC ACCTCCAGCC AGAGAAGAAT CAGTATTGAG GGTCCCTGCT GACCCTTCCG TACTCCTGGA CCCCTTGAC CCCTCTATTT CTGTTGGCTA AGGCCAGCCC TGGACATTGT CCAGGAAGGC CTGGGGAGGA GGAGTGAAGT CTGTGCATAG ATGGGAGAGC CTCTGCTCA GAGGCTCACT CAGTAACGTT GTTTAATTCT CTGCCCTGGG GAAGGAGGAT GGATTGAGAG AATGCTTTT TCCTCTCCTA AGTCTTTGCT TTCCCTGATT TCTTGATTG ATCTTCAAAG GTGGGCAAAG TTCCCTCTGA CTCTTCCCC ACTCCCATC TTA CTGATTT AATTTAATTT TTC ACTCCC AGAGTCTAAT ATGGATTCTG ACTCTTAAGT GCTTCCGCC CCTCACTACC TCCTTTAATA CAAATTCAAT AAAAAAGGTG AAATATAAAA AAAAAAAAAA AAAAAAAAAA AAAAAAAAAA AAAAA</p>	<p>22</p>

<p>SERTA D2</p>	<p>NM_0 14755. 2</p>	<p>CTCCTGCACG GCGAGTGCTG GAGCACGACG TACCGCTCGC TCGGTCAGGG CGCCCCCTCC GCCCGCCTCC TGCTTCCTCC TCCGCTGCCT GCCGCCGCCG CCTCCACCAT TGTATAATGC TCGGGGCGCG CAGGCAGAGA ACGGCGGAGT CTTAGCTTCA GCCTCGCCTG CTGCCCGCTC CCCGGCGCCA CCCTCGGGCC CCTGGAGCGG GGCACCTCCGC ATGGAGCGGG AGTAGCTGAG GAGTGGGCGG AAACCCCTCC TGATGCGTTA GTTCCCAGGT GGAGCTGCAT GTGATATATG TTGGGTAAAG GAGGAAAACG GAAGTTTGAT GAGCATGAAG ATGGGCTGGA AGGCAAAATC GTGTCTCCCT GTGACGGTCC ATCCAAGGTG TCTTACACCT TACAGCGCCA GACTATCTTC AACATTTCCC TTATGAAACT CTATAACCAC AGGCCCTGA CAGAGCCCAG CTTGCAAAAAG ACCGTTTTAA TTAACAACAT GTTGAGGCGG ATCCAGGAGG AACTCAAACA GGAAGGCAGC CTGAGGCCCA TGTTACCCCC CTCTCCAG CCCACCACCG AGCCAGCGA CAGCTACCGA GAGGCCCGC CGGCCTTCAG CCACCTGGCG TCCCCGTCCT CCCACCCCTG CGACCTCGGA AGCACTACGC CCCTGGAGGC CTGCCTCACC CCGGCCTCAC TGCTCGAGGA CGACGATGAC ACGTTTTGCA CCTCCAGGC CATGCAGCCC ACGGCTCCCA CCAAACTGTC ACCTCCAGCC CTCTTGCCAG AAAAGGACAG TTTCTCCTCT GCCTTGGACG AGATCGAGGA GCTCTGTCCC ACATCTACCT CCACAGAGGC GGCCACGGCT GCGACTGACA GTGTGAAAGG GACCTCCAGC GAGGCTGGCA CCCAGAAACT CGACGGTCCT CAAGAGAGCC GCGCAGATGA CTCAAAACTG ATGGA CTCTC TGCTGGGAA TTTTGAAATA ACGACGTCCA CGGGTTTCCT GACAGACTTG ACCCTGGATG ACATCCTGTT TGCTGACATT GATACGTCCA TGTATGATTT TGACCCCTGC ACTTCCTCAT CAGGGACAGC CTCAAAATG GCCCCTGTGT CTGCCGACGA CCTCCTCAA ACTCTGGCTC CTTACAGCAG TCAGCCTGTC ACCCCAAGTC AGCCTTTCAA AATGGACCTC ACAGAGCTGG ACCACATCAT GGAGGTGCTT GTTGGGTCCT AAGACCCAGG GACCCAGCGA CTATGCCAC CCAGACCCCA GAGCGTTCCC ATAACCCTGA CAGTTCTCCA CACTGTGCAT GCACCCTTGC TTGCCTTTTT CAGAGAAAA GAAAATTTTA CAACAGGATC AACTAGTTT TTGCTTTGAG CAGAGTTGGA GTGCCTTCAT CCAAGTATGA CCACTTTTAA TACACTTTTT TGAGTGGTTC CTCAGAGACC TACTACCCTG GTATAGGAAA GAATCCATTT GAAGACAATG TTGCAATGTT GAATGACAAA AATAAACAGT TCAAGTGAAG CACAAGGATT AAGTTGGAAA AGCTGTAAAT TGCATGTGCA TATTTGTCTA TTTTTCTAT AAGTTTTATT GCAAGAGGTA AAGAAGAAAA CTATATATAT ATATCTTATT TAGATAATCT CAGTACCTTT TCTGGCATT</p>	<p>23</p>
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	<p>TTGCCCTGTA TAGGTTGACT TGGCAATTCG GCCTTTTTAG AGGCATTAAC TACTCCTCGT AAGTGTTGCA TTTACATGGC TGTTTAGAAA ACTGCTGCCC AAATTTATTT TATATTTTTG TACAGATTCT GCAGTTTATG ATATTGTTTT TCTAAAAACA AATGCTGTTT ATACATATGA GATAGCTATT TTGATAGGAT TTGCTCACAT AGTTCCTGCA AACTTCAGAT GTACAAGTTG CACTTGTACT TTTATAGAGT TGTAATGTTT TATATGTGTA TGGTGCAAGA GAAAATTGGA TCAAATCAAT CTGCAGTTGA TGTCCCCAAA TGCAAACACA GGCACACACA TGCACACACC CATAAACACA CACACAGTGC TTTAAGAAAAG GGCCAGGTGA TATCACACCC AAATTCACA AGCACTGACC CCCTGGCACC AACACCCGCC AGTACTGTGA CTTCCAAAGC CAGAGCCACA TGTGCTCATC AAAC TTGCAT TAAGCAGTTG GCGGGAGATG GCTGTGGAGC TGGGGGTTTA AGTGATGGTT CTCTTTTGCT CCCTCTTTTG AGGGTAAAGC TACTGTCTTT CTTAAGAGTG TATTTATGCC AAGTTTGCGC TTTAATTTGT TTTTATTTG TTTTTAATG AAAACCCAGA TCTTTCCTTT TTGGCATAAT TTTTATGATG ACCTGAAATT TTACATCCGA ACAAATTTT ACATCCGAAA AGCAACCAAC TTCTTCATGG AACTCAGCCC TGTTGCAATG CTTAGGGCCC TAAAGAAGA AAATCTCCCC AGAAGGCATC CATCATGTTG CTTAATTGTC TTCTGCAGCT TCCTTTCCTT AGAGCTTTC CTGTGTTGCT AAGAGCTGAA AATGGCATCT TCGTGATCAC CACAGTGAGC TTGGCTCGCC TCGGCCGGCC CGGGATGCAC TCTTACAACA TGTGTGACTC TTGAACCTGG AGTTCATCAC ATTACGTCAC AGCTTCCCAT CTGGTTGCTT TCCTGAGTCA GCTACTTAC ACTTGTCAAG GCTGTTTTAC CCCAAACTC AGACAGGACT TTCTATGCAT GTTTTCCCTC CTCCCCCAA TTCCCCCCC CATCACCTTA TCTCCAGGA CACTTTGAG AAGTAGCTTT TTATTCCTAG TGGTGTACAT TTAATTTTAA AAAGGTTGCA ATGTATCATG CTTGTTGCCG AAAC TTGTTA TGGCCTTCTT GTTTCAGTTT TTTCTTTTCT TCCAATGGTA CTTTAGCTGT TGAGTGCAGG TTACAACCTA TATTGTTATG CAGATGGCTT CTTTAGGAAT AACTTTTATA TTTATTTAAA AATTTTTAAA TTATGGGATG TTTTGTIGTT GTTGTGTCT TTGTTGTGG TCATTTGTCA ATATTCAGTC ACCAATTCTG CTCACTTCTT GCCATGGATA AAATTGGGTC TTTCTGGCTA ATTA AAAAAG ACAACTTTAT AAAATGGCAC TTTAAGCAAG CCATAGTTAG TTTTATTTTT GTAATGCACA TGGCAAAGCA AAGACGTTTG TGATGAAGGA ACTGCTCATC TAAGCAAAG ATTIGAGTAT GATATGATAA AGGCTTTCTA CATTCTAATT TACTTTTTCC CCCCACTTGA ATGTGTTTTA AAGGCTAATT ATCAGCTCAG TAGAGCAGTG</p>	
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	<p>AGAAACTGAT CAAATTGCAC TTGTTCTCCT ACAAGCAACC TCCACGCAGA CACCTCGTAC TGCTACAGGT GTGTCATTTC CTTTAATAGG ACCAGGGACC ATGTAAGTGA GGTGAGGGTT GTAGTAGATG CTTCCAGTGT CAGTATGCCT GTTAATTTTA AGAGCTTCCC TTTCTTGCAG AGAACAAGTC TGCCCAGATT CCATGCTTTC TATAACTGGA GGACCTGGCA AACCTGCCGC ATGCTGCACA CATCTACCTA CGTACACATA TACAATAGTA TTGATGATTTC TGAACAATAA CAGGGTAAAA CAGTTGGTTT GCCATTGTTA AAAACTGATT TACAGTAACT TACAACAAC GTACTTTTGT TGGATTAGCA AATCATGTGT TTAAACAAAT CCCATATGTT GGGCAACAGT TCAAATAAGC ACGGAGAAGT GTTGCCCAA CTTGGTTCTC TGACTCTTAT GTATTTGTAA GGCTGGGCTT CAAAATCAAA AAAAAACCC CAAAAACAGC AGGCAAATGC TTTTAACTC TGACACCGTT GCCATAAATC CCTGATACTC AAAGTCTAAC AAGAAAGACA TGGAAAATTA GCAGCCCATT TTCAGAAAGA TCAAAATGAT CTAGGGTTCT AATTGCTTTT GCATCCTATT CTTACAAAGT GATGTCCCAA CAGGGAACAG TAGGAGCTGG AGTGGGATCT CCAAGTCCCA GTTTGAGTGT GGGATGTGCT TCCAGCAGTG CCTTCCCTTT ATGAAAGACA TCACATGGCA TCCAGGGCCA GGCAGGCAGC TTGAGGTGCC TTTACGAGAA AACCGAGCTG GGGCTGGGAG AGGACAGTTA TTGACACTGA TGTGCAATGA AGTGACAAGA TGAGAGCAGA ATCGTAAGAG CTTTGAATTT GAAGTGAGTT TTTTTCCCC CATAAGTTAT TTATTCCTTT TTTCTGTGTA AATATATTTA TTTTACTGTG GAGCGCTAAC ATCTGGATCG TAACATGTGC AGAATGTATG GTAGGAATGT ATTCTCTTGT AGGAATGTAA ATCTGTATTA AAAGGGGGTC CAAGCCAGGC CCCCAGGTCT TCTCATTGTA TGCACAGTCC GCATTCATTT TTACTCTTCT CTAATATGGG TCTATTTGAA ATATGCAAAA GGTATGAGGA ATGTTTTAAT ACCTCCAAAT TTTTAAGAAA AGCATCAAAG GGTGATATT TTTTAAAGTT TTTTAGTAG CACTTCTCTT GGATGACAGA AGGAGCAACC ACATGGGCAC CCTTGTTTAT ACCAAAGGGT GAGCAGTGGC CAGAGCCTCC TCTGCACCTC TCGAGTGTCT TTACCAATTG AGCTTTTTAT CGCCATAGCC CCTTGGAGTG CCCAGCTGC CCTGAGGTCA ATCAAGGAAA ATTTCTTAAT GAAATAAGCT CCAAAGAGCC AAAGTATCAA CTTACAGATC GTTTTTAAAG CTTAAATTTA TGAACCACCT TTGTGGTAAA CAATGAATTA TGAATACCGC AGGGCAGCCT TCTTAAATGA CAAATGTAAA AAAAAAAAAA AAAAAGACTC TACTTCGTGC AGCAATTGCT ACTCTATACG AATTGTCTTA ATTTGAAAAC CTTGCTGTTA CAAATTGGAC CTTTATACAT TTTCTGAAAA CAATGAAAAG AGTATATTTA</p>	
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		<p>ACCTTTTCTG GCTGTAAATG GTTACCTTCC TGTAACCTGCC CCGCACCTGG AGGCATGGAG TTGTGTGCAT CCTGCTTATG TACAATTGTT TTCAGTGTTT CTAAGAATGA GTCTGAATGG TTCTTGAAAA TTAGCCAGGA TCAAATGCTA TTGCAGACAA AGCCAATAAA AAGTTGGACT TCTTTTGGGG ATAACAAGTT TTGGAAGAGA AATGCAGGCC ATATGTGCGC ATGACCGAGA TTTTGAAAAA AGATGTACAT AGTGACATGT TTGGTGCATG GTTTTTGAGG AGGGCTTTTG TCAAAAAGGA GGTATAACCT TTCCCCACA GACCTGAGAG CTGTGCCTTT TCTATGCAAT ATTACAGACG TTACATCGGA ACCCAGATGG CTGTATTAC ATGTAGGTTT GGGCTGTAAT CTAACAATT GGACAGATTA AATGTACATG GAAATGAGCA GTCTTACTTT TGTAGTTTTA TATTATACAA TAAACAGTTA AAAGATGAAA AAAAAAAAAA AAAAA</p>	
<p>SLC35 G5</p>	<p>NM_0 54028. 1</p>	<p>TACCACCTCA GGAGAGTTC AGGGAAGAAC CCCACCCGCA CTCCAATGAG GTCACAATGG CTGGAGCTCT GAGGGGCCA GGCTCCCTGA GCCAGGAGGA GAGGAGAAAG TCCAAGGAAA GATGGCTGGC AGTCACCCCT ACTTCAACCT GCCTGACTCC ACACACCCAT CGCCGCCCTC CGCTCCACCC AGCCTCCGCT GGCACCAGCG CTGCCAGCCC TCTGGTGCCA CCAATGGCCT GCTGGTGGCC CTGCTGGGTG GGGGCCTGCC TGCTGGCTTC GTGGGCCCCC TTTCTCGTAT GGCTTACCAG GGTTCCAACC TGCCCTCGCT GGAGCTGCTC ATCTGTGCGAT GCCTCTTCCA CCTCCCTATT GCCCTGCTAC TTAAACTGCG TGCCGACCCC CTTCTGGGAC CTCTGACAT CCGAGGCTGG GCCTGCTTCT GTGCCCTGCT CAACGTCCTC AGCATTGGAT GTGCCACAG TGCAGTTCAG GTGGTGCCCG CTGGCAACGC TGCCACTGTT CGCAAAGGTT CTTCCACCGT ATGCTCCGCT GTCCTACCC TCTGCCTTGA GAGCCAGGGT CTCGGTGGCT ACGAGTGGTG TGGACTGTTG GGCAGCATCC TAGGACTAAT CATCATTCTG GGACCTGGAC TCTGGACACT ACAGGAGGGG ACCACAGGTG TCTACACCAC CCTGGGCTAT GTGCAGGCTT TCCTGGGAGG CCTGGCGCTG TCCCTGGGGC TTCTGGTCTA TCGTTCTCTG CACTTCCCT CCTGCCTCCC AACAGTGGCC TTCCTATCTG GCTTGGTGGG GCTGCTGGGC TGTGTGCCAG GCCTCTTTGT GCTGCAGACC CCCGTGTGTC CCAGTGACCT CCTGAGTTGG AGTTGTGTGG GGGCAGAGGG GATCCTCGCC TTGGTCTCCT TCACATGTGT GGGCTATGCG GTCACCAAGG CCCACCCTGC CCTGGTGTGC GCTGTCCTGC ATTCCGAGGT GGTTGTGGCC CTTATACTGC AGTATTATAT GCTCCATGAG ACTGTGGCAC TTTCTGACAT CATGGGGGCA GGGGTTGTGC TGGGCAGCAT</p>	<p>24</p>

		TGCCATCATT ACAGCCCGGA ACCTCAGCTG TGAGAGGACA GGGAAGGTGG AGGAGTGAGA TAGAACTTGG GAGCCCGGGG GTTGGGAGGG ACAGGGATAA ATAAAGACAA AGACTGAAGA C	
SPATS 2L	NM_0 011004 22.1	AGTGCTGAGG GAGCAAAGTT CATTTCCTCG GGTAGGAGAA GATGATTCTC TTGCAACACG TGCGGATTGT GACAAAATCT TTCATTAACA AGGGGAGTTT CGGTGAAGTG GAGGTTTGGG GAAAGGCGAG GAAGTCGGTC TGGAGCAAGC AAGCAAAGTG CGGAAGCTGT ACTGGGATTC TTCTAGAAAG TGGGGTGGGA AAGGAGCTAG GGAGGGCGTG TGGAGGGACG AGATCTGTGT CAGAACGTGC GTGTGAGCGG ATACAAAACC CGAGAGAGGC GTGAGCAGCG CTGTGTTTGC GAGCGGGAGC GAGGGGCGCC GGCTGGGGTG TGTGCTCCTG AGCTCTTCAG AAACCAGGCT GCTTTCAGGA ACATTGCTGT GGATTCCTCAG GGCCTATTCC ACTAGAAGCA AGATGGCTGA ACTCAATACT CATGTGAATG TCAAGGAAAA GATCTATGCA GTTAGATCAG TTGTTCCCAA CAAAGCAAT AATGAAATAG TCCTGGTGCT CCAACAGTTT GATTTTAATG TGGATAAAGC CGTGCAAGCC TTTGTGGATG GCAGTGCAAT TCAAGTTCTA AAAGAATGGA ATATGACAGG AAAAAAGAAG AACAATAAAA GAAAAAGAAG CAAGTCCAAG CAGCATCAAG GCAACAAAGA TGCTAAAGAC AAGGTGGAGA GGCCTGAGGC AGGGCCCCTG CAGCCGCAGC CACCACAGAT TCAAAACGGC CCCATGAATG GCTGCGAGAA GGACAGCTCG TCCACAGATT CTGCTAACGA AAAACCAGCC CTTATCCCTC GTGAGAAAAA GATCTCGATA CTTGAGGAAC CTTCAAAGGC ACTTCGTGGG GTCACAGAAG GCAACAGACT ACTGCAACAG AAACTATCCT TAGATGGGAA CCCCAAACCT ATACATGGAA CAACAGAGAG GTCAGATGGC CTACAGTGGT CAGCTGAGCA GCCTTGTAAC CCAAGCAAGC CTAAGGCAA AACATCTCCT GTTAAGTCCA ATACCCCTGC AGCTCATCTT GAAATAAAGC CAGATGAGTT GGCAAAGAAA AGAGGCCCAA ATATTGAGAA ATCAGTGAAG GATTTGCAAC GCTGCACCGT TTCTCTAACT AGATATCGCG TCATGATTAA GGAAGAAGTG GATAGTTCCG TGAAGAAGAT CAAAGCTGCC TTTGCTGAAT TACACAACCTG CATCATTGAC AAAGAAGTTT CATTAAATGGC AGAAATGGAT AAAGTTAAAG AAGAAGCCAT GGAAATCCTG ACTGCTCGTC AGAAGAAAGC AGAAGAATA AAGAGACTCA CTGACCTTGC CAGTCAGATG GCAGAGATGC AGCTGGCCGA ACTCAGGGCA GAAATTAAGC ACTTTGTCAG CGAGCGTAAA TATGACGAGG AGCTCGGGAA AGCTGCCCGG TTTTCCTGTG ACATCGAACA GCTGAAGGCC CAAATCATGC TCTGCGGAGA AATTACACAT CCAAAGAACA ACTATTCCTC AAGAACTCCC TGCAGCTCCC	25

	<p>TGCTGCCTCT GCTGAATGCG CACGCAGCAA CCTCTGGGAA ACAGAGTAAC TTTTCCCGAA AATCATCCAC TCACAATAAG CCCTCTGAAG GCAAAGCGGC AAACCCCAAA ATGGTGAGCA GTCTCCCCAG CACCGCCGAC CCCTCTCACC AGACCATGCC GGCCAACAAG CAGAATGGAT CTTCTAACCA AAGACGGAGA TTTAATCCAC AGTATCATAA CAACAGGCTA AATGGGCCTG CCAAGTCGCA GGGCAGTGGG AATGAAGCCG AGCCACTGGG AAAGGGCAAC AGCCGCCACG AACACAGAAG ACAGCCGCAC AACGGCTTCC GGCCCAAAAA CAAAGGCGGT GCCAAAAATC AAGAGGCTTC CTTGGGGATG AAGACCCCGG AGGCCCCGGC CCATTCTGAA AAGCCCCGGC GAAGGCAGCA CGCTGCAGAC ACCTCGGAGG CCAGGCCCTT CCGGGGTAGT GTCGGTAGGG TTTCACAGTG CAATCTCTGC CCCACGAGAA TAGAAGTTTC CACAGATGCA GCAGTTCTCT CAGTCCCGGC TGTGACGTTG GTGGCCTGAG CTAGGAGGAA AAAGAGCAGT TTCACTCAG TTTTGGTTCC CTGCCGAGG TGCTGACCCA ATTCGCTGCC AAAAGAGTGT CAATCAGAAT ATACAAATCC CGTATGGTTG TGTCATCCTC TCTAATCAT TTTTACTAAT TCTAATAATC AGCTCTAGCT TGCTTCATAA TTTTCATGGC TTTGCTTGAT CTGTTGATGC TTTCTCTCAT CAAGACTTTG CAGCATTTTA GCCAGGCAGT ATTTACTCAT TATTAGGAAA ATCAAGATGT GGCTGAAGAT CAGAGGCTCA GTTAGCAACC TGTGTTGTAG CAGTGATGTC AGTCCATTGA TTGTCTTTAG AGAGTTAATG TTACAAAAAA GAATTCTTAA TAATCAGACA AACATGATCT GCTGAGGACA CATGCGCTTT TGTAGAATTT AACATCTGGT GTTTTTCTGA AAAAATATAT ATACATATAT TGCTTTATTT GAAACAAATT AAAATATGCT GCATTTGACA CCTGGCTAGT TTCTTTTATT GATACCCACC TAGTTATTGA ATGFACTGTT TAGTGCTTTC AAAAAAACT TTAGAGACTA GAGGTTGTGG TGCAAAGCTG TGTACAATAA ATACTGTTTC TGTTGAGACA AGTACTCTTT CAGGAAAATA TATATATGCC CTTTCAATTA GATTACACAA ATAGATGGAT ATGCACCCTG ATCATCTTAG ACACACCATG TGGCAGTTGG GCAGTTGGAA ACATTGGTTG CAAGTACTAC AAACCTCAGC TGAGCCTAGC TTCAACAATA AGAATTTATT GGTTCAAGTGA GTGAAAAGTC CAGTTGGTCC AATTIGATCA GGATTTGAGC TCCCATCTCT TTTTITTTTT TCTTTTTTCT TTTTCTTTTT TTGAGACAGT CTTGCTTTGT CACCAGGTG GAGTGCAATG GCATGATCTC GGCTCACTGC AACCTCCGCC TCCTGGGTTC AAGCCATTCT CCTGCCCGGC CTTCCGAGTA GCTGGGATTA CAGGCATGTG CCACCACGCC CTGCTAATTT TTATAATTATT AGTACAGACA GGATTTACC ATGTTGTCCA GGCTGGTCTC AAACCCCTGA CCTCAGGTGA</p>	
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	<p>TCCACCTGCT TTGGCCTCCG AAACCTGCCGG AATTACAGGC ATGAGCCACC GTACCCGGCC AGCTCCCATC TCTTAATTAT CTTAGCTCTC CTTTCCTCCT TGGGTIGACA TTGCAITCAG AATTGTGGCA AAGGGCCACC ATGCTCTTAA GACTCAAGTC TATTTTCCAC ACTGTCCAGA GGAGAGAAGT TATCCTAGTA GCTTCTACAC AGAGCAAGTG TCTTTCTCAG AAATCCCAGC AAAGGTCTTG CATAACATTG CTGGTAGGCC TGTCTCTTAA ACCAATCATG AAAGGGGGGA GAGAAGTGAA TGGGATGAAC TAATTTGCAT AGACTAATTA GAGCCCACCC CTGGAGCCTG GGTGTGGCCA TCTTCCCAGA GTTCCTGAGC TAGATGGAGA AGGGGTACTT CTCAGAAAGG GAAAATGGAT ACCCAATGGC CCAAACCCAA AATAGCCCCA GTTCCCCTAA CTTTGACTAC AGGGCAGTCC AGTTTGGGTG CCGCTTCCGT TGCACTCACA TGTCTACAT ATCTGTTGAC TACTCACAAG TGCAAATGCT TATTCTCAAC TCAACATTA CTTTTTTTCT GGCAACCCAG GTTCACTGGT TCTCTTCCAC AGAGCGGCCC TGAGCAGCTG AGCCTGCAAG CCACGCAAGC ATCTGTTTCT TCTTTTGCCA AGTACAGGAG GATGTTTGTCT CTCTCTGTAG AGAGCTTTCT GAGGTCTCTG GGTGTACCCA GAGATTTAAT AGAAATTCTT AACGTTAAGT CACATTCCAG GAAGGAAGGA AGAGTTGTTT GTTCAAATAA GAAAGATAAA TGTTCCGGCAC TGTAGGCCCT GTTTACCCCA TCTGAGGCCC TGAATTCATA TATTACAAGA CGGAAGGATT TTGCACAGTT TTTTATGTAG CAAGATTTTG CTCACCACTG AAAAATGTCA GTGTAAATGT GACCGCTTTA AAGATGAGTC AAGTAATTCT TGGAACAGGG AAAAAAATGA ATTTGCCAGG TCAGGAGTTC ACCTGCCTTT GTCAGAGTTG AACCCAACCA CTCTTGACCT CGACTCACTC CTTAGGGTT AAGAAAGCCC AAACACATTC CTGAGCACAG AGCAAACACT CCCATGTCAC TGAAAAGAAA CAAAAGAATG CTA AAAAAGTG CTCAGACCTG ATCACATTTT TTCCAATATT TTTTCCTTTT TTTTTTTTTT TTTGAGACAG GGTCTTGCTC TATCACCTAG GCTGGAGTGC ATGTTTTTGA GACAGGGTCT TGCTCTATCA CCTAGGCTGG AGTGCATGAT CATGGCTCAC TATAGCCTTG AACTCCTGGG CTCAAGCCAT CCTCAGCCTC CCAAGTTGCG AGGACTACAG GTGTGCACCA CAACGTCCCA GCTACTTTTT AAATTTTTAG TAGAAACGAG GTCTCACTAT GGTGCCCAGA CTGGTCTCGA ACTCCGGAGC TCAAGTGATC TTCTTGCTTT GGCCTCCAA AGTGTTAGAA TTACAGGCAT GAGCCACCTC GCCTGGCTGG TTTTGCCTT TCTTATAGAC CCTGGGCATG TAAGCATTTA TTAGTTTGCA TTTTGA AAC AGTAATTICA ATATTTTAGT GCCAATGTCA GGCCGCTTAA ACACGTGATT ACATATCTTC ATCTGTCTGG TGGA ACTATT GGTGTGATCC</p>	
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		<p>TAGAGA ACTG AGTCCTATTC TGCCATTCAT TTAAAGTGTT TTAAACTCTA ATCTCTCTAC TTAATGCACA GTAGTCAGAT TATTCTCTTA AACATTG GCC TAGTAGAGGT TAAAATAGTT TAATCCTTAT GAAGATGGAA TAACTTCAAA CTCACATTGT GGCACCTAGA TCTTCCACCA AGACTTCATC CGTGAAATCC ACACCTCCCT GTTGGGTTCC CAATTACATT CCAAATTTAC ATTTCTTTTG AGAATCTCTG CATACTCCAG CTCTGTCTG TTGATCCTAT TCTAGAAGTG CTTAATGCAG CAAGACACAG AAAGTTAAAC GCAAATTGCT GCAA AATTCA CCCTCAGTGG AGGACTAGAA ACACAACATG TCCAATTTAA AGCTCAGTTC ACAAGCAGTT CAATTCTGCT GGCATCAGAA AAGGAGATTC TAATTAACA TTCTTAGGGA AGGACATCAA ATGAGGTTAA TGGGAAACGT TACCAGATTA AAAGCAGTTT TTTGACAAAG TAACAGATTT GGAAATTCTG ACTCTCTGAA AGCCTTGATT TGAACCTCAA ACTTGATTTC ACCATGAGAA GTGGGGATCA AGGGCCTGCG CAGTTCTTTT CCTAAATCGA TTCGGTGCTC CCCACCCCGA CGCAGGCACA GGTC CGCAAC CAGATAGGGA GATGCCTGAA TTTCAGGCTA CCTTTGACAA AGCTTTCTTC CTCCTCCCT CCCTTTGCAC GCTGCATCCC ACGCTGCCTG CTAAAGCGC CCTCATGTGT CTACAGATGG TAAAACGTTT ATTTCTCAAC AGACATTCCA GTGATAGCAT CCAATGACCT ATGTAACGGC ACCCTTTTTT GCTGTACGCG TTTTTGAGA TGGAGTCTCG CTCTGTCCG CAGGCTGGAG TGCAGTGGCG CGGTCTCGG TC ACTGCAAG CTC CGCCTCC CGGGTTCACG CCATTCTCCT GCCTCAGCCT CCCGAGTAGC TGGGACTACA GGCACCCGCC ACCACACCTG GCTAATTTTT TGTATTTTTA GTAGATACAG GGTTTCACCG TGTTAGCCAG GATGGTTTTG ATCTCCTGAC CTCGTGATCC ACCCCCCTCG GCCTCCCAA GTGCTGGGAT TACAGGGGTG AGCCACTGCG CCCGGCCCCA GTCACTTGTT CTTAAGTTTC TTAAGCAAAC TATAAAATAG CAAAAGACCA AAAAAAAGG AAAAAAAGCA GTTCGCCTAA TACATTGTT CAGCATTTC TTGAAAGTAC TGAGCCATCT CAATTGCTCT GATTTTGTGA GAAAATTATG AAGAGTTGCA AAGTCCCAGT GATTTCTTTG TTACTTAGCT AAGAATTTGA AATGTAATTT AAATACTATT CTTTACAATC CATTATAAGG ATTTTAAAAAT CTTTTTGCTT CTTTAATAAA TTCTAACAAA GAAAAAAAAA AAAAA</p>	
TDRD7	NM_013028 84.1	<p>AGAGCCGAGG CCAGGCTGCC CTCGAAGCGG GGCGGGGCGA AGCGGGGCGG GGCCGAGCAG GGCGGGGCGG GGGCTTGAGG TGATTCCCAA GCCGCGGGGC GGCTCCGGTG GTGCGGGGAA ACCGAAAGTG GGCGGCGGCC GCGGCGGGGC CCCTGGCGGA</p>	26

	<p>GACGGCGGCA GGAGCTGGGC CCAGAGACGC GGGGACGGGC CGTGGGCCCC CGGAACGAGA TTACCTGCTA TGCCATGGCC TGCACAGAAA CTGCAAGAAT TGCTCAGCTT GTGGCTCGTC AAAGGAGTTC TAAAAGGAAA ACCGGGCGTC AAGTTAATTG TCAGATGAGA GTGAAGAAAA CCATGCCATT TTTTCTAGAA GGAAAACCAA AAGCAACCCT CAGACAACCA GGATTTGCTT CAAATTTTTTC TGTTGGCAA AAACCTAATC CAGCACCGTT AAGAGACAAA GGAAACTCTG TTGGAGTTAA GCCTGATGCT GAAATGTCTC CTTATATGCT ACACACAACCT CTTGGAAATG AAGCATTCAA AGACATTCCA GTGCAAAGGC ATGTGACCAT GTCCACCAAC AACAGGTTTA GCCCAAAGGC GTCCCTTCAA CCACCTTTGC AGATGCATCT CTCAAGAACC TCTACTAAGG AAATGAGTGA TAATTTAAAT CAGACTGTTG AAAAACCCAA TGTC AAGCCT CTGCCTCTT ACACTTATAA AATGGATGAG GTTCAAAATC GCATAAAGGA AATACTAAAC AAGCATAACA ATGGCATTG GATATCTAAG CTTCCACATT TTTACAAAGA GTTATATAAA GAAGACCTTA ATCAAGGAAT TTTACAACAG TTTGAACACT GGCCCTCATAT TTGCACGGTG GAGAAACCTT GCAGTGGTGG CCAAGATTTA CTTCTTTATC CAGCTAAGAG AAAGCAGCTT TTGAGAAGTG AACTGGATAC TGAGAAAGTA CCTCTATCCC CACTACCTGG TCCCAAACAA ACACCACCGT TGAAAGGGTG TCCAACAGTT ATGGCAGGAG ACTTTAAAGA AAAAGTGGCA GACCTGCTGG TGAAATACAC AAGTGGCCTT TGGGCCAGTG CACTTCCGAA AGCATTGAG GAAATGTACA AAGTGAAATT CCCTGAGGAT GCCTTAAAAA ATCTTGCCTC ACTTTCTGAT GTATGCAGCA TAGACTACAT TTCTGGAAAT CCCCAGAAGG CCATTCTCTA TGCTAAACTT CCATTGCCCA CTGACAAAAT CAAAAGGAT GCAGGGCAAG CACATGGTGA TAATGATATC AAGGCTATGG TTGAACAAGA GTATTTGCAG GTAGAAGAAA GCATTGCTGA AAGTGCTAAT ACCTTTATGG AGGACATAAC AGTTCCTCCT TTAATGATT CAACTGAAGC ATCACCATCT GTATTGGTGG TTGAAGTGA CAACACAAAT GAAGTGGTTA TCAGGTATGT GGGCAAAGAC TATTCTGCTG CTCAGGAATT AATGGAAGAT GAGATGAAGG AATATTACAG TAAGAATCCT AAGATCACAC CAGTCCAGGC TGTGAATGTT GGGCAGTTGC TGGCCGTA AA TGCCGAGGAG GACGCCTGGT TACGGGCACA GGTATCTCA ACAGAAGAGA ACAAATAAA GGTATGCTAT GTTACTATG GTTTTAGTGA AAATGTTGAA AAAAGCAAAG CATACAAAT AAACCCGAAG TTTTGTTCAC TCTCATTCA AGCTACAAA TGTAAGCTTG CAGGCTTGGA AGTCTAAGC GATGACCTG ATCTAGTGAA GGTGGTTGAA TCTTTAACTT GTGGAAAGAT CTTTGCAGTG GAAATACTTG</p>	
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	<p> ACAAAGCTGA CATTCCACTT GTTGTICTGT ACGATACCTC AGGAGAAGAT GATATCAATA TCAATGCCAC CTGCTTGAAG GCTATATGTG ACAAGTCACT AGAGGTTTAC CTGCAGGTTG ACGCCATGTA CACAAATGTC AAAGTAACTA ATATTTGCTC TGATGGGACA CTCTACTGCC AGGTGCCTTG TAAGGGTCTG AACAAGCTCA GTGACCTTCT ACGTAAGATA GAGGACTACT TCCATTGCAA GCACATGACC TCTGAGTGCT TTGTTTCATT ACCCTTCTGT GGGAAAATCT GCCTCTTCCA TTGCAAAGGA AAATGGTTAC GAGTAGAGAT CACAAATGTT CACAGCAGCC GGGCTCTTGA TGTTTCAGTTC CTGGACTCTG GCACTGTGAC ATCTGTAAAA GTGTCAGAGC TCAGGGAAAT TCCACCTCGG TTTCTACAAG AAATGATTGC AATACCACCT CAGGCCATTA AGTGCTGTTT AGCAGATCTT CCACAATCTA TTGGCATGTG GACACCAGAT GCAGTGCTGT GGTAAAGAGA TTCTGTTTTG AATTGCTCGG ACTGTAGCAT TAAGGTTACA AAAGTGGATG AAACCAGAGG GATCGCACAT GTTTATTTAT TTACCCCTAA GAACTTCCCT GACCTCATC GCAGTATTAA TCGCCAGATT ACAAATGCAG ACTTGTGGAA GCATCAGAAG GATGTGTTTT TGAGTGCCAT ATCCAGTGGA GCTGACTCTC CCAACAGCAA AAATGGCAAC ATGCCCATGT CGGGCAACAC TGGAGAGAAT TTCAGAAAGA ACCTCACAGA TGTCATCAA AAGTCCATGG TGGACCATAC GAGCGCTTTC TCCACAGAGG AACTGCCACC TCCTGTCCAC TTATCAAAGC CAGGGGAACA CATGGATGTG TATGTGCCTG TGGCCTGTCA CCCAGGCTAC TTCGTCATCC AGCCTTGGCA GGAGATACAT AAGTTGGAAG TTCTGATGGA AGAGATGATT CTATATTACA GCGTGTCTGA AGAGCGCCAC ATAGCAGTGG AGAAAGACCA AGTGTATGCT GCAAAGTGG AAAATAAGTG GCACAGGGTG CTTTTAAAAG GAATCCTGAC CAATGGACTG GTATCTGTGT ATGAGCTGGA TTATGGCAAA CACGAATTAG TCAACATAAG AAAAGTACAG CCCCTAGTGG ACATGTTCCG AAAGCTGCCC TTCCAAGCAG TCACAGCTCA ACTTGCAGGA GTGAAGTGCA ACCAGTGGTC TGAGGAGGCT TCTATGGTGT TTCGAAATCA TGTGGAGAAG AAACCTCTGG TGGCACTGGT GCAGACAGTC ATTGAAAATG CTAACCCTTG GGACCGGAAA GTAGTGGTCT ACTTAGTGGG CACATCGTTG CCAGACACCG ATACCTGGAT TCATGATTTT ATGTCAGAGT ATCTGATAGA GCTTTCAAAA GTTAATTAAT GACTGCCTCT GAAACCTTGA CAACTAATTC AGATTTTTTA GCAATAACAA AATGTAGTAG GCTTAAAAAA AATCTTAACT CTGCTACATG GCTCTGACTG CTGTGGGGGA TTGAAAAGAA TATGCTTATG TTTGATGAAA GATATTTAAC AAGTTTTGTT TTAACAGAGT TGACTTTTCA AAGAAAATG TACTTGAATT ATTACTATAA </p>	
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		TATTAGAATA AAAATGTTTA TCAATATAAA AAAAAAAAAA AAAAAAAAAA AA	
TOX4 (housekeeping gene)	NM_013035 23.1	AGCAGAGAGA ACACACGTCC TTGCGGAAGT GACGGCAGTT CCGAGTCCAG TGGGGGCGGT GGGAGCGATG AGGGTCTGAG ACGGTGGGAG CGGTTGTGTG AAGATGGAGA CATTCCATAC ACCAAGCTTG GGTGATGAGG AATTTGAAAT CCCACCTATC TCCTTGGATT CTGATCCCTC ATTGGCTGTC TCAGATGTGG TTGGCCACTT TGATGACCTG GCAGACCCTT CCTCTTCA GGATGGCAGT TTTTCAGCCC AGTATGGGGT CCAGACATTG GACATGCCTG TGGGCATGAC CCATGGCTTG ATGGAGCAGG GCGGGGGGCT CCTGAGTGGG GGCTTGACCA TGGACTTGA CCTCTATA GGAATCAGT ATAGTGCCAA CCCACCTGTT ACAATTGATG TACCAATGAC AGACATGACA TCTGGCTTGA TGGGGCATAG CCAGTTGACC ACCATTGATC AGTCAGAACT GAGTTCCAG CTGGGTTTGA GCCTAGGGGG TGGCACCATC CTGCCACCTG CCCAGTCACC TGAAGATCGT CTTTCAACCA CCCCTTACC TACTAGTTCA CTTACGAGG ATGGTGTGGA GGATTTCCGG AGGCAACTTC CCAGCCAGAA GACAGTCGTG GTGGAAGCAG GGAAAAAGCA GAAGGCCCA AAGAAGAGAA AAAAGAAAGA TCCTAATGAA CCTCAGAAAC CAGTTTCAGC ATATGCTTTA TTCTTTCGTG ATACACAGGC TGCCATCAAG GGACAGAATC CTAATGCCAC TTTTGGTGAG GTTTCAAAA TTGTGGCCTC CATGTGGGAT AGTCTTGGAG AGGAGCAAAA ACAGGTATAT AAGAGGAAAA CTGAGGCTGC CAAGAAAGAG TATCTGAAGG CACTGGCTGC TTACAAAGAC AACCAGGAGT GTCAGGCCAC TGTGGAAACA GTGGAATTGG ATCCAGCACC ACCATCACA ACTCCTTCTC CACCTCCTAT GGCTACTGTT GACCAGCAT CTCCAGCACC AGCTTCAATA GAGCCCCCTG CCCTGTCCC ATCCATTGTT GTTAACTCCA CCCTTTCATC CTATGTGGCA AACCAGGCAT CTTCTGGAGC TGGGGGTCAG CCCAATATCA CCAAGTTGAT TATTACCAA CAAATGTTGC CCTCTTCTAT TACTATGTCT CAAGGAGGGA TGGTFACTGT TATCCAGCC ACAGTGGTGA CCTCCGGGG GCTCCAATA GGCCAAACCA GTACAGCTAC TATCCAGCCC AGTCAACAAG CCCAGATTGT CACTCGGTCA GTGTTGCAGG CAGCAGCAGC TGCTGCTGCT GCTGCTTCTA TGCAACTGCC TCCACCCCGA CTACAGCCCC CTCCATTACA ACAGATGCCA CAGCCCCCGA CTCAGCAGCA AGTTACCATT CTGCAGCAGC CTCTCCACT CCAGGCCATG CAACAGCCTC CACCTCAGAA AGTTCGAATC AATTTACAGC AACAGCCTCC TCCTCTGCAG ATCAAGAGTG TGCTCTACC CACTTTGAAA ATGCAGACTA CCTTAGTCCC	27

	<p> ACCAACTGTG GAAAGTAGTC CTGAGCGGCC TATGAACAAC AGCCCTGAGG CCCATACAGT GGAGGCACCT TCTCCTGAGA CTATCTGTGA GATGATCACA GATGTAGTTC CTGAGGTTGA GTCTCCTTCT CAGATGGATG TTGAATTGGT GAGTGGGTCT CCTGTGGCAC TCTCACCCCA GCCTCGATGT GTGAGGTCTG GTTGTGAGAA CCCTCCCATTT GTGAGTAAGG ACTGGGACAA TGAATACTGC AGCAATGAGT GTGTGGTGAA GCACTGCAGG GATGTATTCT TGGCCTGGGT AGCCTCTAGA AATTCAAACA CAGTGGTGTG TGTGAAATAG TCCTTCCTGT TCTCCAAGCC AGTGAAGAGT TATCTGCTGG GAAAGTGTCC AAGAGCCTGT TTTTGAAACA CAAGCTGGGC TTCTGGTAGT GCCTCATCAC AACCCATGAT GGCTGTTTTCAT GTTTCACCCC TTTTCTTCT TCAGCAGAGG CCAGGCTATG GAGCAGGGCC ACTGAATTTG CTGTAATCTG GAGATGCTTT TFACTTTCAA CCATAAGCGG TAATAGCAGA GGAAAGGGTG AAGGGAGTCT GGGCAAGCAA AGCATAGAGA TGGTGGGGTG GTGGTGGGGT TGAAGAACT TGTTGGTATA ATTGTCATAG GACTTGCCTA AAATATTATT AAAATTACGG GAGTGTACTC AGCTTTGAGC CTAGGAGAAA ATGCCACTGT GTGCATCCAT TTTAAAGGGT TCCCTCATAA AAAAATGTTA TTCCCATT TAACATCAGT AACTGCTTT GAAAACAAA CTTTTCAACA TGGGCATACT GGGCTACATG GAAAATGACA TCACCAGGA GTGATTTCTC TTTATATATA TTATTTCTGC AGTTACCATC CTTATCTGAG TTATCACAGT TCATGAATCT AAGAGGCGGA ACTCTACATC ATTAGTAAGA GGTTCCACCA AAGTCTAAAG TTGTATTAC TTGTGTTTGA TGAATATCT TAAAAGACC ATAGGTCTAT CATTATTTCT TAGACATAAT CAAAAGAAA ACAGACTAGA GAAGCCACCT GGTTGTAACA GAATAAGCAG AAGTTTACAG CATGATAGTC CAAGTGGTGA TAACTTTAAA TAAAAC TCAA ATTTTACTG TTTGTAGACA GGAATGCTGT CCTAGAGAAC CTCCTCCTCA ACCAGCTACG TACATAGTTT TATCCTATGC ATTCCTGTTT TCTGIGTGTT TTTIGTTTTT TTTTTTTTTT TTTTTTTTIG AGACAGAGTC TCGCTCTGTC ACCCAGGCTG GAGTGCAGTG GTGCGACCTC AGCTCACTGA AACCTCTGCC TCCGGGGTTC AAGCGATTCT CCTGCATCAG CCTCCCGAGT AGCTAGGATT ACAGGCGCCC GCCACTACGC CCAGCTAATT TGTGGTATTT TTAGTAGAGA CAGGGTTTCA CCATGTTGGC CAGGCTGGTC TCGAACTCCT GACCTCATGA TCCGCCCGCC TTGACCTCCC AAAGTGCTGG GATTACAGGC ATGAGCCACC GCACCCAGCC TGCATTCTG TTTTTTAAT GGTTTTGGAG GGTAGCAGTA GAGATGGGGT CTCACTATGT TGCCAGTCT AGTCTTGAAC TCCTGGGCTA CAGTTACCCT CCTACCTCGG CTTCCTAAAG </p>	
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		<p>TGCTCGGATT ACAGGTGTGA GCCACTGTGC CTAGCCTATA ATGATCATT TAATGTTTCC CATGCACTCA TTTAGTTTGA ACCTTCACAG CAACCCAATG AGGTAATACT CCCATTTTAC ATATAATACT GAGAGATGAG TTGCACAAGA TTATACACTG TTAAGTAGCA GAGCCAGAAT GGACTTCAGA ATCCCAACTA CAATACAAAT GTTTATTTAA ATAAAGAAGA AAGCTATTGT ACAAATATCA CTCTTCAGGT TTAGCTTACA GAGCCATGGC TATGGATTCT TAGCTCTGTA AGGAAGTGCT TCTATAAATT CTTAGGTTTA GAGATGATAC CATCTGGGTA CCTTTGCTTG AACCGTGCAA CCACATCTGG GTCTAGTAGG TGGATCCCAT CCAGTTGGTT TCCAAGGGTG ATCCTGAAAC AGTGTA AAAAG GAGGGGCAA CCAGAAATCC TGGAATTAGA GGGTTAATA TTGTTAAAA ATGCATACCA AATGAAGACT GCCTATCATC ATATCAAATA TGCCAATTCT AAAAAGAGCT TAACATTAGA ATAGTATATG GTAGAATTAC TAGTTCAGAA TTGGCATAGA TTCTGGTGTT AAAATAGACT GGATCTGTAT TATCTGAGGG TTAGTAACTA ATGCTTAGCC AGGCCTGCTT CACAGAGTTG CTACCAGGGA GTATTCTTTG GATAAGCAAA ATGCTAGCAG CATGTGTTTT AAGCTCTGTT AAGGGGTGAA AGATGTAATT ATTGACAGAT TAAATAGATA ACTTCGTAAC CACCAGGGGG CAGATTCAAT ACATCACAGA ATGGCTGAGG AAGATCCTTG GGTGTGAAG AGAGTAGAAA CCCTAGGGAG CAGTGCTTTT GGGTCCTAGA ACCTGTTGAG TTTCTAATGA ATATTTGTAG AATCTCATAA AACAGTTTAA ATACAAGCTT AAGTGGCTTA TGAATCCTGT GAAGCTCATT TATGGACTAG TGTA AAAACAA TGTGAAGCTC TACTAAGTTC TGTCCTTAAT CATAAATAAT AGCCCCTTGA GGACTAGCCT GTTCTCTGGT CACCTTACCA GTTGGGTTGC ACATTGTGTG GTCGTCCAAA TAACTCAATC TTGCGAGTGC CAGGAGATAG TCTTTCAATC ATGCCATAGA TTTTATCTGG TTTATGACTG GTGGAACGAA CCTAGGAAAT AAAA ACTAGC TGCTTTTAA GTTACACAAG AAAAA</p>	
<p>TPTI (housekeeping gene)</p>	<p>NM_012862 72.1</p>	<p>CTTCGTGCCA CGTCACCGCC TCGTTCGCTT CCGGAGGCGC AGCGGGCGAT GACGTAGAGG GACGTGCCCT CTATATGAGG TTGGGGAGCG GCTGAGTCGG CCTTTTCCGC CCGCTCCCCC CTCCCCCGA GCGCCGCTCC GGCTGCACCG CGCTCGCTCC GAGTTTCAGG CTCGTGCTAA GCTAGCGCCG TCGTCGTCTC CCTTCAGTCG CCATCATGAT TATCTACCGG GACCTCATCA GCCACGATGA GATGTTCTCC GACATCTACA AGATCCGGGA GATCGCGGAC GGGTTGTGCC TGGAGGTGGA GGGGAAGATG GTCAGTAGGA CAGAAGGTAA CATTGATGAC TCGTCAATTG GTGGAAATGC CTCCGCTGAA GGCCCCGAGG GCGAAGGTAC</p>	<p>28</p>

	<p>CGAAAGCACA GTAATCACTG GTGTCGATAT TGTCATGAAC CATCACCTGC AGGAAACAAG TTTCACAAAA GAAGCCTACA AGAAGTACAT CAAAGATTAC ATGAAATCAA TCAAAGGGAA ACTTGAAGAA CAGAGACCAG AAAGAGTAAA ACCTTTTATG ACAGGGGCTG CAGAACAAAT CAAGCACATC CTTGCTAATT TCAAAAATA CCAGTTCTTT ATTGGTGAAA ACATGAATCC AGATGGCATG GTTGCTCTAT TGGACTACCG TGAGGATGGT GTGACCCCAT ATATGATTTT CTTTAAGGAT GGTTTAGAAA TGAAAAATG TGATGCAAAA GAAAGAAATC CCTGCGCTTT CTGTCTGTCT TTGTGGCGGC CCAGATTGAA TTGGGGAATA CATCTTTAGC CTGGAAATGT AGGCTGCATG TTAATGGTAA TGAACTTTT GCAGTGTAAT GTTTGAAAAA TATTAATGTA GTTTTIGCTT TTACAGTAAC AAATGTGGCA ATTATTTIGG ATCTATCACC TGTCATCATA ACTGGCTTCT GCTTGTGATC CACACAACAC CAGGACTTAA GACAAATGGG ACTGATGTCA TCTTGAGCTC TTCATTTATT TTGACTGTGA TTTATTTGGA GTGGAGGCAT TGTTTTTAAG AAAAACATGT CATGTAGGTT GTCTAAAAAT AAAATGCATT TAAACTCATT TGAGAGAATG CCTTTTAGTT TAATGCATAT TAAACTAAA TTGATCCTGT AGTGTTCTG GAGAAGCTAG AGCCTGATTG TAGGCTACTA CTCATCAATT AACTTCTACA GTGGAGACTA CTTCTGGGAC TGGAATATAA AAAAGAATCA AAGGTTCTGA TTTTGAGTTG CAATAAAGGG AAAGACCATG CTCATAGCAG TGCCAACATC TGAAGTGTGG AGCCTTACCC ATTTTCATCAC CTACAACGGA AGTAGTTAAC TGGAAGAGAT TACCAAGAGA ATAAAAAGAG ACTCATTAG TGGAAGCAAC TTTGTCTCAG CTTATTTTAC ATAAAGAGAG CGAAGTCTTT TGGGATGAAT GTTAATTTAA CTCCCTGGTA ACTAGAACAG GGACTGGCAA ACTAGCCTAT CTGACCACCT GTTTTGTACA CTTTAAGGTG GTTGGTTGCC TTTTTAAATG GTTGAGGGGA AAAGAATACC TTGTGGGATA TGGAATTTAA GTTCGAGTCC AGTTTTATTG GAACGTGGCT ATGCTTATTC ATTTATGGAT TGACTIONG TGTGTGTCAGT GCATGAGCAG AGTTGTGTCT AACAGACTAG AGCCTGCAAG TTTGCCAGCC CCTGATTTAA AAGATGAAGG TACACAGAAT GTGGGCTGGC TGGTGGGCAA AGGGGTAAAA ATGTTCTCTA TATTGTATCT GAAAAGATGG GGTGTCTGAA TAAGAAAATG CATCTATTTG ACAGACCTGG AGCAGTTGCT ATCTGCTGCT ATGGTTTCCA CCACAGATGC AAGAAGAACA TGTCCTTGCG CTTTCCGTCT GTCTAATIGT GGCAGCTGAG ATTGAATAGA GGAATACAGG AGGAAAAAAA GCGGGAAGAG TTTTGTAGGC AGGTCGGTCA CCCAGGCTTG TAGTGCAGTG GCACAAGCAA CTCACTGCAT TCTCTGCATC CTGTGCTCAA GCCATTTTCC</p>	
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	<p>CACCTCAGTC TCACTAGTTG CTGGGACTGC AGGCATGCAC CCCTATGCCC AGCTAATTTT TGTAGAGACC GAGTATCGCT TAGTTGCCCA GGGTGGTCTC AACTCCTGGG CTCAAGGAGA TCTGCCACC TCAGCCTCCC AAAGTGCAGG CCTAGCCTGG GAGGGGAATT TTCAAACGT GAGTTTTGGG AAATAGTCTA TCAGCCTTAC CTGGTTGATT ACACTTGTA AAGAAAGATT AAAAGCAGGC CAGTGA CTCT GGTCTGCTTG AACATGTGAA TGTAGTGGTT TGAGCAATCT GGAGTTTGCC CTAGTGTCAA ATTCCAGACT GTCCATAGTG TCCAAAACCT GAGGCAGATA CTAATGTAA CCCCAGCAC CCCGTGATTG GAAACAAACC TAAATACGTA TTGGGAACTT AATAGCAATT TTAAGCATT TGATAGATTT TTTGTAGGGA TGGGGTCATG CCATGTGGCC CAGGCTGGTC TGAAAACCTCT GGCCTCAAGT GATCTCAAGC TTTGGCCTTC TAAAGTGTG GGATTACAGG TGTGAGGCAT TGCACCTGGC TTAGCGTTCT GATTTGACAT TGTAATGAAA AGTGTGAGTC TCATCTACAG GGCCTTTTGT CCTCTGAAAT GATAGCAGGA AGGGAATTTT CAGGCAGTGG TCAAAGCTGG GGAAACCAGG ATAGTGAAGA AGGCCTTGAG GTGAGAGATG GAAGCTAATT GGTGAACTAG CCTTGGAAGC CTGAAACAGA CAAGTAGCAA TTCAGAGACT TTGTGGGCTC CACTGCTCCA ACTTGTTTTG AAGATTTTCA GTTCTGCAGA AGAGGTATTT CCCAGTTGT CCTTTCAGTG CTCTTAGCTG TTTTCCCAAC ATCCAGATCC AATCAAGGCT GGGACATAGC ATTTTATCAT GTCTATTTAA GTCAGAAGTG ATGAACCCCA GCTGTTTACC TCATGGTAAA CCTTGAAGA TTCCAGGTAG AATCTTCTCA GACTTTGAAG ACTGTCTCAT TTTATATCTT TTTCTCGTTA TTCCTAGGGT CAAGACGTTT TGGGCAAGAA TAAGGATGTG AACATCAGAA AGCTCATAAC ATTTTGTFTT TGATGCTAAG TTTAACAAG GCATGCTTTA GTAGCCTGTG GGCCCTAGGG TTTGTAAAG TGTGGAGAAC AACTGAGTGG AGCAAGAGGA CTTTTCTAGG AAGGTCCTTG TAATGTGACA TTTGAAAACA AATGAAGGTG TGGAAGTAGG CCATGTGGAT ATCAGGACAA ACCATTCCAG GCCAAGACAA CAGCAGTTAG TCTGGAGTGT GATGTGTTCT GGGAAAAAAG TGGCCACTTT GCTAACCCAA GAAGACAGGA AGGGTTGTAA AGCAGTGGGA GTGTGCAAGG AAGGAAGACC AGACCTCAAG GAAACCACAG GCGCTCTGAG CAGAAGAGTT ACATGATATG ACTCAAATTT TAAAGGATC ACTTTGGCTG CCAGGTGGCA GGGTAAAAGC ATAGAATAAT TGTGTATAAT GTGTTTTTAA GGCAAAGATA GTGGCTTAGT CTAGGGTAGT AGACTGAGGT GGTAGGAAAT GAAGATAGAG ACAACAGGAT ATGCTGGTGG GTGAGGATGG ATTTAATGTT GATACAAGTA TTTTGGTCTG AGCGTTTGA AGAAAGTTGG</p>	
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		<p>CACTGAGGTG GGAAGTCGAG TTTAGTTTTG TTAGTTTTGG ATGTGTTAAG TTTGAGATGC TGATTCCTCA GAGAAGTCTA AGCTGGAGAA CTATATAGAG AGTGGAAAGA TAACAATAGA CATTGAAAGC CATGATACAG GATAAGGTCA TTTGGAGAGA GGATAGACTG CATTCCAACA TGAGATTGGT TGACAAAGAG AAACCAACAA AGGTAATTAA GAGGTGCTCC CACTGCACTT GTACTCAGAA GGCTGAGGTA GGATTGTTAG AGGCCAGCCT GGCACCACA GGGAGACCCC ATCTCTAAAA TTTAGCCAGG AACCATGGCT CATGCCTGTA GCCCCAGGAA TTTGGGAGGC TGAGTGGGGA GGATCGCTTG AGGTCAGGAG TTTGAGACCA GCCTGGGCAA CATAGGGAGA CCTAAAAAAA TTAATTGGGC ATCTGTAGTC CCAGCTACTC AGGCGGCTGA GCTGAGAGGA TGGCTTGAGT CCGAGAGATT GAGGGTGCAG TGAGCTGTGA TCATAACACT GCACTCCAGC CTGGGCGGCA GTGAGACACT ATCTGAAAAA AGTTTAAAAA TTTTAAAAAA GAAGGAACTG CCCCTGAGGT AAGAACCAAG GGAGGGCCTC CCAGAGGTCA GGTGGAAAAA GTTTTAGGAA GGAGGAAGTA GTCAACAGGG TTACCTGTTG CAAAGTACTT AAGTAATATG AGGCCTGATA GTGGTAAACT TGACTACCGT TGGATTTTAC TAGTGGGAAA GGAAGTCTAA TAAAATGCA CTCAAGAGAC TAACAGTCGC AGGCATGAAA TACAATACAG GTACATGGTT TTTTATTATG TGTGCATCTG CTTTCAGTAAT AGGTGTGAAT TACTCATTG GATCATTAGG AGTTTCAAAA TCTAGTTAAA TGACTAGATT TTTGTTGATG TAAATTCTGT CATTCTGAAC TGCAGGGATT GTCAGTAACT TAACTGCAAA CTAAACTGGT GATAATTATG GTAAAATTGC AAGACGAGCA ATAAATCTCA ACCAACTTGA GAGAACACTG ATAA</p>	
<p>TXK</p>	<p>NM_0 03328. 2</p>	<p>GATTCAGTT GAAAGATGTG TTTTTGTGAG TAGAGCACCG CAGAAGAACT GAAGACTGTT GTGTGCTCCC CGCAGAAGGG GCTACCATGA TCCTTTCCTC CTATAACACC ATCCAGTCGG TTTTCTGTTG CTGCTGTTGC TGTTTCAGTGC AGAAGCGACA AATGAGAACA CAGATAAGCC TGAGCACAGA TGAAGAGCTT CCAGAAAAAT ACACCCAGCG TGCAGGCCG TGGCTCAGCC AATTGTCAAA TAAGAAGCAA TCCAACACGG GCCGTGTGCA GCCGTCAAAA CGAAAGCCAC TGCCTCCCCT CCCACCCTCT GAGGTTGCTG AAGAGAAGAT CCAAGTCAAG GCACTTTATG ATTTTCTGCC CAGAGAACCC TGTAATTTAG CCTAAGGAG AGCAGAAGAA TACCTGATAC TGGAGAAATA CAATCCTCAC TGGTGGAAAG CAAGAGACCG TTTGGGGAAT GAAGGCTTAA TCCAAGCAA CTATGTGACT GAAAACAAAA TAACTAATTT AGAAATATAT GAGTGGTACC ATAGAAACAT TACCAGAAAT</p>	<p>29</p>

	<p> CAGGCAGAAC ATCTATTGAG ACAAGAGTCT AAAGAAGGTG CATTATTGT CAGAGATTCA AGACATTTAG GATCCTACAC AATTTCCGTA TTTATGGGAG CTAGAAGAAG TACGGAGGCT GCCATAAAAC ATTATCAGAT AAAAAAGAAT GACTCAGGAC AGTGGTATGT GGCTGAAAGA CACGCCTTTC AATCAATCCC TGAGTTAATC TGGTATCACC AGCACAATGC AGCCGGTCTC ATGACTCGTC TCCGATATCC AGTTGGGCTG ATGGGCAGTT GTTTACCAGC CACAGCTGGG TTTAGCTACG AAAAGTGGGA GATAGATCCA TCTGAGTTGG CTTTTATAAA GGAGATTGGA AGCGGTCAGT TTGGAGTGGT CCATTTAGGT GAATGGCGGT CACATATCCA GGTAGCTATC AAGGCCATCA ATGAAGGCTC CATGTCTGAA GAGGATTTCA TTGAAGAGGC CAAAGTGATG ATGAAATTAT CTCATTCAA GCTAGTGCAA CTTTATGGAG TCTGTATACA GCGGAAGCCC CTTTACATTG TGACAGAGTT CATGGAAAAT GGCTGCCTGC TTAACCTATCT CAGGGAGAAT AAAGGAAAGC TTAGGAAGGA AATGCTACTG AGTGTATGCC AGGATATATG TGAAGGAATG GAATATCTGG AGAGGAATGG CTATATTCAT AGGGATTTGG CGGCAAGGAA TTGTTTGGTC AGTTCAACAT GCATAGTAAA AATTCAGAC TTTGGAATGA CAAGGTACGT TTTGGATGAT GAGTATGTCA GTTCTTTTGG AGCCAAGTTC CCAATCAAGT GGTCCCCTCC TGAAGTTTTT CTTTTCAATA AGTACAGCAG TAAATCTGAT GTCTGGTCAT TTGGAGTTTT AATGTGGGAA GTTTTTACAG AAGGAAAAAT GCCTTTTGAA AATAAGTCAA ATTTGCAAGT CGTGGAAAGCT ATTTCTGAAG GCTTCAGGCT ATATCGCCCT CACCTGGCAC CAATGTCCAT ATATGAAGTC ATGTACAGCT GCTGGCATGA GAAACCTGAA GGCCGCCCTA CATTGCCGA GCTGCTGCGG GCTGTCACAG AGATTGCGGA AACCTGGTGA CCGGAAACAG AATGCCAACC CAAAGAGTCA TCTTGCAAAA CTGTCAATTA TTGTGAATAT CTTACCATA TGGGGTCACT TATGGTGAAT ATCTTTCTTC AGAGTTGCTG ACTCTTGAAA ACAGTGCAAA GATCACAGTT TTTAAAAGTT TFAAAAATTT AAGAATATTC ACACAATCGT TTTTCTATGT GTGAGAGGGA TTTGCACACT CTTATTTTTT TGTAATAAT TTCACATCCC AAATGTGAAG AAGTGAAAAA GACTTCGCAG CAGTCTTCAT TGTGGTGCTC TTCATGATCA TAGCCCCAGG AACCTTGAG GTTCTTCTTC ACAAGGCTGA GAGTGCTTCC TTCTTGAAGA CGAGTGACAT TCATCACTTC AGTGATCCAT GCATAGAATA TGAAAATAAA TTCTTCCAAC TCATGGGATA AAGGGGACTC CCTTGAAGAA TTTTATGTTT TTGGGCTGTA TAGCTCTTTA CAGAAAATGC ACCTTTATAA ATCACATGAA TGTTAGTATT CTGGAAATGT CTTTTGTAA TATAATCTTC CCATGTTATT TAACAAATTG </p>	
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		<p>TTTTTGCACA TATCTGATTA TATTGAAAGC AGTTTTTTGC ATTCGAGTTT TAAACACTGT TATAAAATGT AGCCAAAGCT CACCTTTGAA CAGATCCCGG TGACATTCTA TTCCAGGAA AATCCGGAAC CTGATTTTGT TTCTGTGATT TTACACTTTT TACATGTGAG ATTGGACAGT TTCAGAGGCC TTATTTTGTG ATACTAAGTG TCTCCTGTAA TTTTCAGGAA GATGATTTGT TCTTTCCAGA AGAGGAGACA AAAGCAAGAT AGCCAAATGT GACATCAAGC TCCATTGTTT CGGAAATCCA GGATTTTGAA TTCGAGATGA AACCAACCAGC AATCACAGTT AAATCTTAAC TTTGCCCTGCA CTCTTTGTAG GAATGATCAG AAATTTATCT TTATCATTCT GAGTGCTTCA GGAGTACAAT AGGAAGAAAG ATACTGGAGA AAGCACTAAT GTAATCACCA TGAAGTCTGA CAACAGGAGC CCATTATTTG CGTACTGTCC CACCCTGTAT CATGGTTCTC TGGGAACAAG CTTTATGATT CTCATTAGAG TTTATTTGTT GATTGTGAGT AGTTGCGACT TTTAAATTAT ATTTCCCCCA CTCAAAGAAT GGTATCTTTA TATATCAATG ACATTCAATA AATGTGTATT ATTTCTAATG AGAA</p>	
<p>YY2</p>	<p>NM_2 06923. 3</p>	<p>ATTAAATAAG AGAAAAAAGC ATAAAAGGAG TAGGCTTTTT TGGCAGATAC TGTTATATGG GATGGACAGG AGGGGAAGAC ATGAAAATCA ACTAAAATGA AATACATAAA AAATCAGTAA AGAGCCTGCT TACGAAAGTG GAAAAACAAC AGCTGGGCGG GGTGCGAGGG TGGCAAACGT ACGCGGGCAC GTGCACGTGC TTTTGGGGCC GACAGACGCG CCAGTTGCCT AGGCGCATGC GTCTGGCTAT CCCAGAAGCA CCTGCGCCTT CCGGCTCGTG CTTTCCTCAG TCTCGCGCCT TTCTCTGCAG CTCGCGCCTT TCTCTGCAGC TCGCGCCTTT CTCTGCAGCT CGCGCCTTTC TCTGCAGCTC GCGCCTTTCT CTGCAGCTCG CCCCTTCCTC TGCAGCTCGC CCCTTCCTCT GCAGCTCCA CCTCACTCCC CTCAGCGTTC TTTTCCCAC GGTCTTCCCG TTGCCGCTAA CCTAACTAAC TCTCAGCCAT GGCTCCAAC GAAGATTCT CCATCACACA AGACCTGGAG ATCCCGGCAG ATATTGTGGA GCTCCACGAC ATCAATGTGG AGCCCCTTCC TATGGAGGAC ATTCCGACGG AAAGCGTCCA GTACGAGGAT GTGGATGGCA ATTGGATCTA CGGTGGCCAC AACCATCCGC CATTGATGGT GTTGCAGCCG CTCTTACGA ACACGGGCTA TGGCGACCAC GACCAGGAAA TGCTTATGTT GCAGACACAA GAGGAAGTGG TGGGCTATTG CGACTCAGAC AACCAGCTAG GCAACGACTT GGAGGACCAG TTGGCCCTCC CGGATAGCAT TGAAGACGAG CACTTCCAGA TGACCCTGGC CTCTCTGTGCG GCCTCGGCGG CATCAACATC AACATCAACC CAGAGCCGCA GCAAAAAGCC CAGCAAAAAG CCCAGCGGCA AGAGTGCCAC CAGCACTGAG</p>	<p>30</p>

	<p>GCCAACCCGG CAGGCAGCAG CTCCAGCCTG GGCACGAGGA AGTGGGAGCA GAAGCAAATG CAGGTCAAAA CGCTGGAGGG TGAGTTTTCC GTGACTATGT GGTCCCCTAA CGATAACAAT GACCAAGGGG CAGTGGGTGA AGGCCAGGCT GAAAACCCAC CTGATTATTC CGAGTACTTG AAAGGGAAGA AACTTCCTCC TGGGGGGTTA CCAGGCATTG ATCTCTCAGA TCCTAAACAG CTGGCAGAAT TTAATAAAGT GAAGCCCAA AGGTCCAAAG GAGAACCTCC CAAAACAGTC CCTTGCTCTT ATAGCGGCTG CGAAAAGATG TTCCGGGATT ACGCCGCCAT GAGAAAACAT CTCCACATCC ACGGGCCCAG AGTCCACGTA TGTGCAGAAT GTGGCAAAGC TTTTCTTGAG AGCTCAAAGC TGAGACGACA CCAGCTGGTC CACACCGGCG AGAAGCCCTT TCAGTGCACA TTCGAAGGCT GCGGGAAACG CTTTTCCCTT GATTTCAATT TGCGCACACA CTTGCGCATC CACACCGGCG ATAAGCCCTT CGTGTGCCCC TTCGATGTTT GCAACAGGAA GTTCGCTCAG TCAACCAACC TGAAAACCCA CATATTAACG CATGTGAAGA CCAAAAACAA CCCGTGAAAA GGAGAAGACC CCTCTCAGAC TTGGGAATTA TCTTCCAGGA CTGCGGTAGG GAATAAATAT GCCTCTCAA GCTTTGTATG TTGTTTCTAA GAGTTTTAAA AAAAAATGAA TCCTGCACAT TTAAGGTTTCG TGTTTTGTTA GAGTAGTAAA AATAGAATTT AAACGTTTTT AAAAAGGTAA ACCTTGACAT AAGATAATAG TGCTAAGATG CCATAGCTTG TTCTGTA ACT ATTTTTGTAA AGTTTGGTCC CAACAGGAGA AAAATTCGTA GACTTCACAT CAAGAGACGG TTCTTACAAA CTGTTTTAAA TGGGACTTTT CACATTCTTA GAAATAGGAA GTTCATTTAT TGTTTACAAT GTTTTTAAAA AACTTGTTAA AAAATTCAAA GTGTTTATGT TTATACTTTT AGGAATATGC TTAATAAGTC TATGTATGGT TTTTCTGGAG GTTGATAACT TTGGGAAAGA TTTACTTTAA AAGAGTGAAC AATTATATGC ATACGTGAAG TATTTTCTG CTAAAAAAG TTATATAGGT GTTATTTGTT TTAATCTTGG TTGTAGTCTT GGATGTTAAC ACATCTTGCA TTTTAGCTGT ATTAGGTCAT GTAGTATTGA TATTAGGTGA TTTAATAGTA CTAGTTTAAA CCTATTTTAG TCATTTTATT TTCCCAA AAA TACTACCAGA TGCTGTTGTT TAGTGTAATT TCTTTGCCTG TTCAGTTAAA GTAGTGCTTG CTTGTAGAAT ATATTGTGTA TATGTTGACT TTAACACTTA AGAAGTACAT CCTGTGTAAT AGAAAAAGCA AAATAAAACA CCTCTTCTAA AGAAGGAAAA AAGTAGTTTG CCTATATCAG TACAGAAGTT AGAACTAAAG AAAAGGGGGA GGGTGCTACT GGCATCTGGT GAGTGGAGGG AGATCAGGAA TGCCACTAAA CATCCTACAA TGCACAGACA GCCCCACGAA ACACATAATT ATTTGGCTAA AATGCCAATA GTGTCAGGTG CAGGGGCTCA</p>	
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		<p>GGCCTGTAAT GTCAACACTT TGGGAGGCCG AGGTGGGTGG ATCGTTTGAG CTCAGGAATT GAACATCAGC CTGGGCAACA TGGCATAACT CGGTCTCTAC C</p>	
<p>ALG9</p>	<p>NM_0 010776 90.1</p>	<p>GTCTTTTGTC CCTCGGCGGA CACCGTTTGC CAGCCAAAGC TATGTCTGCG CGCTCACCGA CTTCATAGGG TGCCGAATTC TTTTTTCCCC AGGCTTGCCA TGGCTAGTCG AGGGGCTCGG CAGCGCCTGA AGGGCAGCGG GGCCAGCAGT GGGGATACGG CCCCCGGCTGC GGACAAGCTG CGGGAGCTGC TGGGCAGCCG AGAGGCGGGC GGC GCGGAGC ACCGGACCGA GTTATCTGGG AACAAAGCAG GACAAGTCTG GGCACCTGAA GGATCTACTG CTTTCAAGTG TCTGCTTTC GCAAGGTTAT GTGCTGCTCT CCTGAGCAAC ATCTCTGACT GTGATGAAAC ATTCAACTAC TGGGAGCCAA CACACTACCT CATCTATGGG GAAGGGTTTC AGACTTGGGA ATATTCCCCA GCATATGCCA TTCGCTCCTA TGCTTACCTG TTGCTTCATG CCTGGCCAGC TGCATTTTAT GCAAGAATTC TACAACTAA TAAGATTCTT GTGTTTTACT TTTTGCATG TCTTCTGGCT TTTGTGAGCT GTATTTGTGA ACTTTACTTT TACAAGGCTG TGTGCAAGAA GTTTGGGTTG CACGTGAGTC GAATGATGCT AGCCTTCTTG GTTCTCAGCA CTGGCATGTT TTGCTCATCA TCAGCATTCC TTCCTAGTAG CTTCTGTATG TACTACTACGT TGATAGCCAT GACTGGATGG TATATGGACA AGACTTCCAT TGCTGTGCTG GGAGTAGCAG CTGGGGCTAT CTTAGGCTGG CCATTCAGTG CAGCTCTTGG TTTACCCATT GCCTTTGATT TGCTGGTCAT GAAACACAGG TGGAAGAGTT TCTTTCATTG GTCGCTGATG GCCCTCATA TATTTCTGGT GCCTGTGGTG GTCATTGACA GCTACTATTA TGGGAAGTTG GTGATTGCAC CACTCAACAT TGTTTTGTAT AATGTCTTTA CTCCTCATGG ACCTGATCTT TATGGTACAG AACCCCTGGTA TTTCTATTTA ATTAATGGAT TTCIGAATTT CAATGTAGCC TTTGCTTTGG CTCTCCTAGT CCTACCACTG ACTTCTCTTA TGGAATACCT GCTGCAGAGA TTTCATGTTT AGAATTTAGG CCACCCGTAT TGGCTTACCT TGGCTCCAAT GTATATTTGG TTTATAATTT TCTTCATCCA GCCTCACAAA GAGGAGAGAT TCTTTTTCCC TGTGTATCCA CTTATATGTC TCTGTGGCGC TGTGGCTCTC TCTGCACTTC AGAAATGTTA CCACTTTGTG TTTCAACGAT ATCGCCTGGA GCACTATACT GTGACATCGA ATGGGCTGGC ATTAGGAACT GTCTTCCTGT TTGGGCTCTT GTCATTTTCT CGCTCTGTGG CACTGTTTCA AGGATATCAC GGGCCCCCTG ATTTGTATCC AGAATTTTAC CGAATTGCTA CAGACCCAAC CATCCACACT GTCCCAGAAG GCAGACCTGT GAATGTCTGT GTGGGAAAAG AGTGGTATCG ATTTCCAGC AGCTTCCTTC TTCCTGACAA TTGGCAGCTT CAGTTCATTC CATCAGAGTT CAGAGGTCAG TTACCAAAC CTTTTGCAGA AGGACCTCTG GCCACCCGGA TTGTTCTTAC TGACATGAAT GACCAGAATC TAGAAGAGCC ATCCAGATAT</p>	<p>31</p>

	<p>ATTGATATCA GTAAATGCCA TTATTTAGTG GATTTGGACA CCATGAGAGA AACACCCCGG GAGCCAAAAT ATTCATCCAA TAAAGAAGAA TGGATCAGCT TGGCCTATAG ACCATTCCTT GATGCTTCTA GATCTTCAA GCTGCTGCGG GCATTCTATG TCCCCTTCCT GTCAGATCAG TATACAGTGT ACGTAAACTA CACCATCCTC AAACCCCGGA AAGCAAAGCA AATCAGGAAG AAAAGTGGAG GTTAGCAACA CACCTGTGGC CCCAAAGGAC AACCATCTTG TTAAC TATTG ATTCCAGTGA CCTGACTCCC TGCAAGTCAT CGCCTGTAAC ATTTGTAATA AAGGTCTTCT GACATGAATA CTGGAATCTG GGTGCTCTGG GCTAGTCAAA GTCTATTTCA AAGTCTAATC AAAGTCACAT TTGCTCCCTG TGTGTGTCTC TGTTCTGCAT GTAAACTTTT TGCAGCTAGG CAGAGAAAGG CCCTAAAGCA CAGATAGATA TATTGCTCCA CATCTCATTG TTTTTCCTCT GTTCAATTAT TTAGTAGACC GGAGAAGAGC AGAACCAACT TACAGGAAGA ATTGAAAATC CTGGTACTGG ATGGCTGTGA TAAGCTGTTT TCCACACTCT GGCCTGGCAT CTGAGAACTA GCAAGCCTCT CTTAGGCCAT ATGGGCTTCT CCACCAAAGC TGTTTGGCAG CTCCTAGCAG ACCTTCTTAT TGAAATCCTC ATGCTGAAAA TGAACACAGC CTAGTTGCCA ACCCACATGT CCTTTTCACC TCCAGCAAGA CTAAGCTTCT TAAAGCACT TCACAGGACT AGGACCCTGT CCTGGAGCTA TCTCAGGAAA AAGGTGACCA TTTGAGGAAC TGTGACCTAA TTTTATTATA ATGATGCCTC TAATTTTCAT TTCCTTTACA ACCAACTGTA ACTATAAGGT TGTATTGCTT TTTTGTTCAG TTTTAGCATG CTATTTTTTG AATTCTAGAC TCCTCCATGT GAAGATATCA ACAGACAAAA CTACAAC TGT ATAGGACATA TTTGGAGAAA ATTCTATCAA TTGATACATT TGGATGACAT CACATTTTTA AGTAATGTAA TCTGAGGCCA TTGCTGAGGA AATTAAGAAT TTTCCTTTTT TTTAACCAC CCCAGTGAA AAGGATCAGT GTATATTTAT AGCACCTATT TTTTAGTTCT GTCGTGTGAG AGGCACATCC TGCATGGGGC ACTTCTAGTC AAATAGGCAA TGATAAGGAC CTAATTAATA TGTGATAAGT GTATACTATT ACTTTAAAAG CCTTTACAGT CAGTACTTCA GTTTACAAGG CACTTTCACA GCATCTCGTT TGATCCTCAC AGTCACAACA TGTGGTAGAC AAGGCAGGTG ATTTTTATCC CCATTTTACA GATAAGGAAA CAGGCTGCGG GTGGGGAGTG AGGGGAGGTA AAGATAGTTA GTTGCCTAAG GTCACACAGC CAGTAAGTAA TAGAGCTGGG ACTGGAACCC AGGTTTCCTT ACTCTCATCT ATTGCTCCTC CATATTCCTC ACTCAACCAT GAAAACATTA CTTGAAAGGA CTGATGAGGT TAACCAGAGA CCTAACTGAT ATTGTAACCTT TCTATTTTAA GGAAGAATTG TGTCTGTATT TGAGTTCTTT GGAGCCTCCA GTCTGCCTGT GTGTTAGACC AGCACAGCAG TGCTGTGTGA TGCAGCCTGA CCTGTGGCAG GAAAGTAGTG CTTCTGTTG GAAGTCATGT TCTTTTGCAG CCACACAGGA TCCAAATATC AGTACTATTC CTGTAGTCAA TCTGGGGTCA CATTATAGGT GCCTTATTTT CCTAAGGGTA ACTGATCTGA ATATCTGCAA ATAGGATGAA TCTATTTTTT AGAAGTTCCA TCTTTCATT</p>	
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	<p>TTCTTTTTTT TTTTGAGACA GAGTCTCATT CTGTGCGCCA TGCTGGAGTG CAGTGGCGCG ATCTCGGCTC GCTGCAACCT CTGCCTCCCA GGTTGAAGCA ATTCTCATGC CTCAGCCACC CGAGTAGCTG GGATTACAGG CATGCGCCAT CATGCCCAGC TAATTTATGT ATTTTTAGTA GAGTTGGAGT TTCACCATGT TGGCCAGGCT GGTCTTGGAC TCCTGACCTC AGGTCATCCA CCCGCCTCAG CCTCCCAAAG TGCTGGTATT ACAGGCGTGA GCCACCGCAC CCAGCCCCAT CTTTCATTTT CAAAGAGAAG GGCATTCTAA TAGGAACTGG TGCCAAGAGA GAAGAAAAGA AGTGATAACA GAAGAAATGG CTAGTTACAA TATTA AAAAG CTCCTCTTTG AGATCTCCTC TGCAGGAATA TCAGAGACGG AGTTGAAGCG CTGGAGAGGT AATAGGTCTA GACAGTACAG ACAATAACT GGGGAGTGTG TGAGGATAGA CTGGGCTCCC CCTTGCTTGA AAGATCTCTG GCATTTAATT CTC AATTCTT GATTACTATT TTCCAGTGTA AACTAGCAC ATATGATCTG ACTACAGGAC AGAGAATTTT AAGTGAAACA TTTGCCTTAC TTGCAGTAAT AATGTGCTGT TCTTCACAGT AGCTAAGGCC CTCTATGTTT CCCAGAGGTA AATAAGAATC CAGGAATGGA GGTCCATCTG TGATGAATGG CTTTTTCTA ATCAAAGTAG TATAATGCTG TTTTATCTGT TTTGTCATCT TGTTTTTTTT TTTTTTTAAA AAAACAAAAC CTTAATTATA ATATAGCGCA AAGAAAGGCC AGGACTGATG CAGGGATTCC TTGGAAATAT CAGTTCCTAT CACTTTTAAA ACCTGATTTT GGATCTCTCT GTTCTATGTA TGTCTTTAGT GAGAGCACAA TACATGGCAG AACGCTGTGC CAAATGTTAT AGGTAAGGAA TATAGAAATG AATGTTTTTT GTTGTGAAGG TGTTTTCATG TGATATTTTA TAAACACATT TTA AAAAATC TCCATCACTT TTTAGTATAG GAAGGATAGC TTTGCCTGGG AAAACAGTT TCAACACACC TGCTCAGAGT AGCAGTTCTC CCTCAAAAAA GCAGTGTICA GCCTGCACTG ACTGTTCTGC TTGCCAAAAG GAGGAAGCAT GCAAGATACT TATTTCTCCA TAGATTGTGG AGTATAGAGG GATGTGGGAC TACAGATTAT TATTTTTTTT CCCCAGACA GAGTCTTGCT CTGTGCGCCA GGTTGGAACA CAATGGCAGC ACCTCAGCTC ACTGCAACCT CTGTCTCCCG GGTTCAAGCA ATTCTCCTGC TTCAGCCTCC TGAGTAGCTG GGATTACAGG CACACACCAC CACCGCACTC AGCTAATTTT TGTATTTTTA GTAGAGGTGG GGTTTTACCA TGTTGGCCAG GCTGGTCTTA AACTCCTGAC CTTGTAATCA TCCCGCCTCG GCCTCCTAAA GTGCTAGGAT TACAGGCATG AGCCACCGCA CCCGGCCCAG ATAATTTTTA ATAGCCTTTG ATCATGGGGT GAGTGAGGGA GTAGGTATAC TTGGCAAATG CATGGTTCTC TGATTTCTAG CTCTAAAGCA GCCTTATCTG AATCCCCAAA TCTTGTGATG CTGAGTACCA TTA CTGAACC AGTCTGCACG GTAGGCATCT GCTACCAAAA TTTACCTCCT ACCTGGTAGG TGTCATCTGA TAAGAAAGAA GACAGGTTAT TTTAATTTTT TGAGATAATC ACAGAAAATT GCAGCCATA CTCTTTATTA CCGAATTCAA GTTTGAAAT AGACCCTTTG TTTTAAATCA TGATGGGTCT TTATCCCAAT CATTATCTG GGTCATTTT CCAACTTTGG</p>	
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		<p>AGTTCTAGGA AAGAACCTTG AAAACCTGAT ATGATTCTGC AGCATGAGGT CTACGGTGAC CATTGGGCA AAGCTCCAGT GGCAATCATT TATTGTGTTT TGCATTTCT GGGATTTATT GAAATAAGAA TTCACTGTGA TTATGTAGTC TTCTGGCTAG TATCAGGCAG CTCTGCTTTT AATTTGGTTA ATTTATTTT CTCTGAAGAG GGAGAAGAGG TACAATTTAA TCTTGGCCTC CACAAGCATA TTAAAGCTCA CGTGTTAATC AGTGCATTCT TATGCTCCTA CATTAAATGC CTTGGGTAAA TGGATAAATG GACATGTGCC CAGCTTTAAT TTTTTTGGCA ACAGAAAGAT CAGACTTCCG TATGGCATCG TTGGATTTCA GAGGCTTTCT GGTGTATCTG TAAATCTGAA TGTTGCCTTC TGCCAGTCTG TATAACCAGG TGATTATGTC TGCAAATGAA ATCAGGAAGC AGTAAAGTGT TAAAGCAAGA GTATTGTCCA ATTCACTTGT CTTCTGATC CTGTACTTT ATTTACGTG TCGGTGTTA CATTACATAC TTATATTTCC TGTGAAAGAA AGAGTAAAT AAATTGTAGC AGTTTGA</p>	
<p>SEPN</p>	<p>NM_0 31475. 2</p>	<p>AGCGGAGCGC CAGGCAGCGC GGAGCGGAGG CCAGGCCAC AGCCGCTCCG CCTCCCGGCC CGCAGATCCC CGACGGCCGC ACCGCGGGCT CCTCTGGCCC GCAAGAACAC GTGCATGGCG TCCTGGGGAA GCGCGTGAAT GCGGAGTCGC GCGCCGCAC GCGGCACCAT GGCCCTGGAG CAGGCGCTGC AGGCGGCGCG GCAGGGCGAG CTGGACGTGC TGAGGTCGCT GCACGCCGA GGCTCCTGG GGCCCTCGCT GCGCGACCCG CTGGACGCGC TGCCCGTGCA CCACGCGGCC CGCGCTGGGA AGCTGCACTG TCTGCGCTTC CTGGTGGAGG AAGCCGCCCT CCCC GCCGCG GCCCGCGCCC GCAACGGCGC CACACCGGCC CACGACGCT CCGCCACCGG CCACCTCGCC TGCCTGCAGT GGCTGCTGTC GCAGGGCGGC TGCAGAGTGC AGGACAAAGA CAATTCTGGT GCCACAGTCT TGCATCTGGC TGCCCGCTTC GGCCACCCCG AGGTGGTGAA CTGGCTCTTG CATCATGGCG GTGGGGACCC CACCGCGGCC ACAGACATGG GCGCCCTGCC TATCCACTAC GCTGCCGCA AAGGAGACTT CCCCTCCCTG AGGCTTCTCG TCGAGCACTA CCCTGAGGGA GTGAATGCC AAACCAAGAA CGGTGCCACG CCCCTGTACC TGGCGTGCCA GGAGGGCCAC CTGGAGGTGA CCCAGTACCT GGTGCAGGAA TGCGGCGCAG ACCCGCACGC GCGCGCCCAC GACGGCATGA CCCCCTGCA CGCCGCGGCG CAGATGGGCC ACAGCCCAGT CATCGTGTGG TTGGTGAGCT GCACCGACGT GAGCCTGTCC GAGCAGGACA AAGACGGCGC CACCGCCATG CACTTCGCGG CGAGCCGCGG CCACACCAAG GTGCTCAGCT GGCTGCTGCT GCACGGCGGG GAGATCTCGG CTGACCTGTG GGGCGGGACC CCGCTGCACG ACGCCGCCGA GAACGGGGAG CTAGAGTGCT GCCAGATCCT GGTAGTGAAC GCGCGGAGC TGGACGTCCG CGACCGCGAC GGGTACACGG CCGCCGACCT GTCGGACTTC AACGGCCACA GCCACTGCAC CCGTACCTG CGCACGGTGG AGAACCTGAG CGTGGAGCAC CGCGTGCTTT CCCGGGATCC ATCCGCAGAG</p>	<p>32</p>

	<p>CTGGAGGCTA AGCAGCCGGA TTCAGGCATG TCCTCACCCA ATACCACGGT GTCGGTCCAG CCGCTGAACT TTGACCTCAG CTCGCCTACC AGCACCTCT CCAACTACGA CTCCTGCTCC TCCAGCCACT CCAGCATCAA GGGCCAGCAC CCTCCATGTG GGCTTTCCAG CGCTAGAGCT GCAGACATAC AGAGCTACAT GGACATGCTG AACCCGGAGC TGGGCCTGCC TCGGGGCACG ATTGGGAAGC CCACACCCCC ACCACCCCCA CCCAGCTTCC CCCCGCCACC CCCGCCCCA GGCACCCAAC TGCCCCCACC CCCACCTGGC TACCAGCTC CCAAGCCTCC TGTAGGACCA CAGGCAGCTG ACATCTACAT GCAGACCAAG AACAACTCC GCCACGTGGA GACAGAGGCC CTCAAGAAGG AGCTGAGCTC CTGTGACGGC CACGACGGGC TCGGGAGGCA GGA CTCCAGC CGCAAGCCCC GCGCCTTCAG CAAGCAGCCC AGCACGGGGG ACTACTACCG GCAGCTGGGC CGCTGCCCCG GCGAGACGCT GGCCGCACGC CCGGGCATGG CGCACAGCGA GGAGGTGCGT GCCCCGCCAGC CCGCGCGCGC CGGCTGCCCG CGCCTCGGCC CTGCCGCCCG CGGCTCACTC GAAGGCCCT CCGTCCCCC GCAGGCGGCG CTGCTTCTTG GGAACCATGT TCCTAACGGC TGCGCCGCGG ACCCAAGGC GTCCAGGGAG CTGCCACCGC CGCCCCACC GCCGCCCGCG CCCCTGCCGG AGGCCGCGAG TTCGCCACCG CCGGCCCGCG CTCTGCCCT CGAGAGCGCT GGCCCTGGCT GCGGGCAGCG CCGTCTCC TCGTCCACCG GCAGACCAA GTCCTTCAAC ATGATGTCCC CGACGGGCGA CAACTCGGAG CTACTGGCTG AGATTAAGGC AGGCAAGAGC CTGAAGCCGA CGCCCCAGAG CAAGGGGCTG ACCACAGTGT TCTCAGGCAT CGGGCAGCCG GCCTTCCAGC CCGATTCCGC GCTGCCTTCT GTGTACCTG CACTGTCACC AGTCCGGAGC CCCACACCGC CAGCTGCGGG GTTTCAGCCG CTGCTCAATG GAAGCTTGGT TCCCGTGCCG CCCACTACTC CTGCGCCGGG AGTGCAGCTG GACGTGGAGG CTCTCATCCC CACGCACGAT GAGCAGGGCC GGCCCATCCC CGAGTGGAAG CGCCAGGTGA TGGTGCGCAA GATGCAGCTG AAGATGCAGG AGGAGGAGGA GCAGAGGCGG AAGGAGGAGG AGGAGGAGGC CCGGCTGGCC AGCATGCCCC CCTGGAGGCG GGACCTCCTG CGGAAGAAGC TGGAAGAAGA GAGGGAGCAG AAGCGGAAAG AGGAGGAGCG ACAGAAGCAG GAGGAGCTGC GGC GGAGAA GGAACAGTCA GAGAAGCTGC GGACGCTGGG CTACGATGAG AGCAAGCTGG CGCCCTGGCA GCGACAGGTC ATCCTGAAGA AGGGGGACAT CGCTAAGTAC TAGAGGCCGC AGACTCCTGT CCGCAGCCTC GCAGCTCCGT GGGGCCCTCC GCCCCAGCCC CAGCCAGCCA GGCCCTGGTG GAAAGGCTGG GAGCCGCACA GCCCTCCCC CCTGCGCTGG AAACCCTCCC TGACCCCCAC CCTGGCCCC CGTATCCCCA GCCCTGGCA AACTGGAGT GCACACGCCG CCACGGTTGC CCAGAAAAAG TGCCCAAGCT GCTGACGCAA ACAACAACA ATGCTGCTTA TTTGCATGCC GACTTACATA TATTTGCATG TTCGTTGACT ATCAAAGAGT GCAGAGCTCT CCCAGCCCC GTGGGTGGTG ACTTTGTTTT CCTGCGGGGC TCAGCCCCCT CCAGGATGCA GCCCCCTCCC CCGCACCCCG</p>	
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		<p>GAACCGGCGT CGCTGGCGCA TCCTGGGTGG AGGCAGGCC CGAGCTCGGG GAAGGGGTTT TCCCTTCCTC TCTGACCCAG ATCTGCGCGC GGCCTAGCCC GGGCCTCATT TCTTATCCCC GCCAAGGGTT TCCTCTCAGT CATTGTGTTA CCAGAAACAT GAAAACCTGCC TGTCTGGCCG GGCCGCACTT GTGGCCCCCG GGACCCACC TCTGGCCCCA CCTCCCTCAA GTCTGCGCCC CGTCCCAGC CAGACCCACT CGCTGCCGGG ACCCTTTCAC TGCCCCGGTG GAGTGAATAG AGGATGAGGG GCCCTGACCC TGTGTCTCCA ACTGCTGCAC CCCATCCCCGA CCCTGTCTCC GCCACCTCGC AGCCCCATTA AAGCGCTCTC ATCTGGGCTC CGGTTCACTC A</p>	
<p><i>YWHAQ</i></p>	<p>NM_006826.3</p>	<p>TTGGGCGGTG GACCGCCCCT CGGCCCCGGG GTAGGCTGAC ACGGGAGGGT CCTCAGCTAA AGCCAAAAGC AGATCAAAGT GGTGGGACTC GCGTCGCGGC CGCGGAGACG TGAAGCTCTC GAGGCTCCTC CCGCTGCGGG TCGGCGCTCG CCCTCGCTCT CCTCGCCCTC CGCCCCGGCC CCGGCCCCGC GCCCGCCATG GAGAAGACTG AGCTGATCCA GAAGGCCAAG CTGGCCGAGC AGGCCGAGCG CTACGACGAC ATGGCCACCT GCATGAAGGC AGTGACCGAG CAGGGCGCCG AGCTGTCCA CGAGGAGCGC AACCTGCTCT CCGTGGCCTA CAAGAACGTG GTCGGGGGCC GCAGGTCCGC CTGGAGGGTC ATCTCTAGCA TCGAGCAGAA GACCGACACC TCCGACAAGA AGTTGCAGCT GATTAAGGAC TATCGGGAGA AAGTGGAGTC CGAGCTGAGA TCCATCTGCA CCACGGTGCT GGAATTGTTG GATAAATATT TAATAGCCAA TGCAACTAAT CCAGAGAGTA AGGTCTTCTA TCTGAAAATG AAGGGTGATT ACTTCCGGTA CCTTGCTGAA GTTGCGTGTG GTGATGATCG AAAACAAACG ATAGATAATT CCAAGGAGC TTACCAAGAG GCATTTGATA TAAGCAAGAA AGAGATGCAA CCCACACACC CAATCCGCCT GGGGCTTGCT CTTAACTTTT CTGTATTTTA CTATGAGATT CTTAATAACC CAGAGCTTGC CTGCACGCTG GCTAAAACGG CTTTTGATGA GGCCATTGCT GAACCTGATA CACTGAATGA AGACTCATAC AAAGACAGCA CCCTCATCAT GCAGTTGCTT AGAGACAACC TAACACTTTG GACATCAGAC AGTGCAGGAG AAGAATGTGA TGCGGCAGAA GGGGCTGAAA ACTAAATCCA TACAGGGTGT CATCCTTCTT TCCTTCAAGA AACCTTTTTA CACATCTCCA TTCCTTATTC CACTTGGATT TCCTATAGCA AAGAAACCCA TTCATGTGTA TGGAATCAAC TGTTTATAGT CTTTTCACAC TGCAGCTTTG GGAAAACCTC ATTCCTTGAT TTGTGTTTGT CTTGGCCTTC CTGGTGTGCA GTACTGCTGT AGAAAAGTAT TAATAGCTTC ATTCATATA AACATAAGTA ACTCCCAAAC ACTTATGTAG AGGACTAAAA ATGTATCTGG TATTTAAGTA ATCTGAACCA GTTCTGCAAG TGACTGTGTT TIGTATTACT GTGAAAATAA GAAAATGTAG TTAATTACAA TTTAAAGAGT ATTCCACATA ACTTCTTAAT TTCTACATTC CCTCCCTTAC TCTTCGGGGG TTTCTTTCA GTAAGCAACT TTTCCATGCT CTTAATGTAT TCCTTTTTAG TAGGAATCCG GAAGTATTAG ATTGAATGGA</p>	<p>33</p>

		<p>AAAGCACTTG CCATCTCTGT CTAGGGGTCA CAAATTGAAA TGGCTCCTGT ATCACATACG GAGGTCTTGT GTATCTGTGG CAACAGGGAG TTTCTTATT CACTCTTAT TTGCTGCTGT TTAAGTTGCC AACCTCCCCT CCCAATAAAA ATTCACTTAC ACCTCCTGCC TTGTAGTTC TGGTATTCAC TTTACTATGT GATAGAAGTA GCATGTTGCT GCCAGAATAC AAGCATTGCT TTTGCAAAT TAAAGTGCAT GTCATTTCTT AATACTAG AAAGGGGAAA TAAATTAAG TACACAAGTC CAAGTCTAAA ACTTTAGTAC TTTTCCATGC AGATTTGTGC ACATGTGAGA GGGTGTCCAG TTTGTCTAGT GATTGTTATT TAGAGAGTTG GACCACTATT GTGTGTTGCT AATCATTGAC TGTAGTCCCA AAAAAGCCTT GTGAAAATGT TATGCCCTAT GTAACAGCAG AGTAACATAA AATAAAAAGTA CTTTTATAA ACCATTTACT ATGGCTTTGT AACAATTGCA TACCCATATT TTAAGGGACA GGTGAATTTA CTACTTTCTA AAGTTTATTG ATACTTCCCT TTTATGTAAG ATGTAGTAGT GATACCTATA TTTCCACATT GTGCATTGTG ACACACTTGT CTAGGGATGC CTGGAAGTGT ATAAAATTGG ACTGCATTTT TTAGAGTGTT TTTACTATAGA TCAGTCTCAT GGGCCATCTC TTCCTCAGAT GTAAATGATA TCTGGTTAAG TGTTATATGG AATAAAGTGG ACATTTTAAA ACTAGCAAAG TTAAAAA AAAA AAAAAA AA</p>	
<p>VPS37A</p>	<p>NM_0 011451 52.1</p>	<p>GCAGAGGGGG CGGAGAGCGC CCCCAGGGGGC GGGGCACGCA AGTGACGGCG GCGCGGGTGG TGGAGCGCTG GGCGGCCAGG CTCCCTGGCT GGCCGGTTTG GGCCTCTGGG CCGTGAAGGT GGGACCTCCT GTTCCGGGCC GCAAGTTTCC CTCTCCAGCC GCCCCCGT CGTAGCATGT CCCCAGAAC TCGGGGAGCG CAGGCAGGAC AGGCTTAGAG AAGACGCGGT CCCAGCGCT TGGGCCACGG ACGTCCCACC CCGTCTCTCT GTCGCTGGAG AACCGCCGGG CCGAGCCACT GGGAGAAGCA GGCCAGAGCC TTCCAGGGCC TCCGGCCCGT GGACCCGAGG AGGATGAGCT GGCTTTTTCC CCTGACCAAG AGCGCCTCCT CCTCCGCGGC TGGGTCCCC GGTGGCCTCA CCAGCCTCCA GCAGCAGAAG CAGCGCCTGA TCGAGTCCCT CCGGAACTCA CACTCCAGAT TGCTTCTCC ACAGTTTCTT CAGGAAAAAC CAGTGATCAG TGTTTATCCA CCAATACGAC ATCACTTAAT GGATAAACAA GGAGTGTATG TTACCTCTCC ATTAGTAAAC AATTTTACAA TGCACTCAGA TCTTGGAAAA ATTATTCAGA GTCTGTTGGA TGAGTTTTGG AAGAATCCTC CAGTTTTAGC TCCTACTTCA ACAGCATTTC CTTATCTATA CAGTAACCCA AGTGGGATGT CTCCTTATGC TTCTCAGGGT TTTCCATTTC TTCCTCCATA TCCTCCACAA GAAGCAAACA GGAGTATCAC TTCTTTATCT GTTGCTGACA CTGTTTCTTC TTCAACAACA AGTCATACCA CAGCCAAGCC TGCCGCTCCT TCATTTGGTG TCCTTTCAA TCTGCCATTA CCCATTCCA CAGTGGATGC TTCAATACCG ACAAGCCAAA ATGGTTTTGG GTACAAGATG CCAGATGTCC CTGATGCATT TCCAGAACTC TCAGAACTAA GTGTGTCACA ACTCACAGAT ATGAATGAAC AAGAGGAGGT ATTACTAGAA CAGTTTCTGA CTTTGCCTCA ACTAAAACA ATTATTACCG</p>	<p>34</p>

	<p> ACAAAGATGA CTTAGTAAAA AGTATTGAGG AACTAGCAAG AAAAAATCTC CTTTTGGAGC CCAGCTTGGG AGCCAAAAGA CAAACTGTTT TAGATAAGTA TGAATTACTT ACACAGATGA AGTCCACTTT CGAAAAGAAG ATGCAAAGGC AGCATGAACT TAGTGAGAGC TGTAGTGCAA GTGCCCTTCA GGCAAGATTG AAAGTAGCTG CACATGAAGC TGAGGAAGAA TCTGATAATA TTGCAGAAGA CTTCTTGGAG GGAAAGATGG AAATAGATGA TTTTCTCAGT AGCTTCATGG AAAAGAGAAC AATTTGCCAC TGTAGAAGAG CCAAGGAAGA GAAACTTCAG CAGGCGATAG CAATGCACAG CCAATTTTCAT GCTCCACTAT AGATTTTCTT GGAAACATGA ACTGCCAAGA GAGGAATGGG ACACAAAACC AAACACTGTT TTATATTTAT GGTTTGCAAA CTGGCATTTC ATCAGTGGCT AAATTCACAG ATATCCTATA TAGATTGTAT ACAGAACTGA GACTGATTTT GTACCGATTA GAATGATTGC TATGATCTTT GAGAAATTTT TCTGCACTAT TTGCACTGAA ATGTTTATTT ATTGTTGATA AATTGTATCA TATTTAAGTT CCACTGCTGT TCCTCTTACC TTGATTAAAT GCCTATGCAT GTACTTTTAG CTAGTTTTTA ATATTTTATA AACTTCATT TAAATTTGTA TTTTAACTT GAAGTTCCAT TTCTTTATCA AGGATGGTAT TTAGATTTTT TTCCTCTTAA CCTTTTTTCA AAAACTATTT TCAACTGTGA GGAAACCCTT ATTTTCTTTT CTTTGTGGAT AAAACTTTCA AAAGCAATTT AAGATATTCA TAGTGTTAGG AAACACCAAA CCTGCCTATG TGCCATCTCA CAAAAGAAAC TTTTAATACC TACAATAAAT CAAAAGAATA AACCAGCTGT TCTTATATAT TGTTTTCAATTT TAAAACTAA AGATGCATTT AAGAAGCAAT ACAAGTAAAT ATTTTACCTA ATAGGAAAAA AAAAAGTTGC CTTTCATTTA AACCATTCCA ACAGAAATTC TTATGCTAAT TAAAAACATA TATATATCTG GTAGGTTTGT GGTTGGATAG GTTTTCTAAA TTCCTAATGT TAAAAACAAT CTTTATGTTA ATATACACTA AATCTATACA CAAAAAAGT CAGTGAACCTT TTCTGACCTT TACTGTGAGT TACCTTTTCC TAAGAGGAAA GCTATAGTAA TAAGTAAAT TTAATTTTTA GGCAATCCTG ATTTTTAATG AATTTAATTG AGTGTTCTTG TATACTACAT TGAGCAGTTT GCTTCTATAC CGTGTCACAA AATTCATGTA TTTCTTGAGA AGCCCTAAAA GCTCATAAAG GAAAATGCCG TGAACATATGT AGCTCAGGCT TGGTAAGGTG CCATCTAAAT TACAAAACAA ACTAATGCAT AATTTTGCTT AAATTCATC CCAGTATGAT TGTCTTCCCA ACACCAGCAT ATAGTATAGA TTGTCTGTCT TTTTATATT TTTTAGTTCT TCCTGTACAT GTTTTTGGCA ATAAAGTTAT AGGAAGAACA AAATTATTTT GTTAGAATTA AAACATGCTT AATATTTAGT CTGTTTGTGG AGGGCAGGTA TTCACGTGGA CTGAGATACA ATGTTGGATA CAGAAAATAA CTTTCATTGT CTTCTGACA CTGTGCTAAG GACATGCTGT TAAAGCTTCA AAGTGACCAG ATGAGGAAGG AATAATTAAT TATTACTCCT GATTTGTAGA TAACTGAGGT AAGAGTGTTT CAAATTTATG ATAGTCTTTT GGGTATTCAG AAACCTTTCC TTATACTGCA CTGGCCACCA GAGCTTAATT TTCCAGCAG TTACAGCAAT </p>	
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		<p>GGGAGATAGA ACAGTCTCAA TCTTTTGCCA ACCATCAGGT TCCTAGAAAC CAGGTAGGTG TATCCCATAA CAAGGGAGGA GCATACCACA GCCCCTCATT TGATTAATTC ATTTGATCTA TCTATGTTAT TAAGTACCTA CTAGGAATAA GGCATTGTGG AAATACTATA CAAAGATAAA CATTGTTTAG ATGCTTATCT ACTTTCCTTT TCACCAGAAA AACAGAAAAA AAAGAAACAT TTTCTTACAG AGTAAAAATG TTCTACATAA TCACATGAGT AGTTCATCTC AGTGTTTTTT ATTCITTTAAA GTTGAACAT CCCAGTTTCA TTCTATACCA TTCATTGGAT AACCTTGTTA CAACCCAGTC ATGAAACAGA GCAGTGTGAT CAGTTATCTG CATTFAACAA ATAGACAAAT CAGTTTACAT AAAGGTTATG TATGTCACCC ACGATGAAAA GAATCTGCAT TTGAATATGC CCGTATGAAT GTGGGTTCTG TTTTTGCAAC AGAGATTAAG TGACCATTTT TTCTAATTTT ATGGCTATAT ATTTTCTTCA TAAAAATTGG TCACATCGGA GAAGCAGTGC CACAGGAAAA ATGAAATGCA TGTGAAAGTT TGTATTCTGA TTTTACAAGA TGAGATAGAA ATCAGAATTA AAGAGGAATA CTTAGGAGTT ACTAGGCTAA TCAGTGTACG AATTTGTCAT AGGTAGAGAT TTAAAGGTTA ATATCTTAAA ATAGAAGAAA ATTCTAAATC AATCAATCAG TGAGATATAA ACTAAACAGA CCCACTTCAA AGTTGAAAGA AATTTCTAGG CATAAATTGA GACTAGGAAA TTTATATCAG AATAGAGGGT GCTTGACACA TATATATGCT TAAATTGAAG GACAGCTCAG ATTCATTTTT AGGAGAAGAA AGTAAACTAA TGTGCTCTTA AAGAATAAAA ATTTATTCTA TGGTTTCTGT CTCTGATCAT CACCTTCCAT TCTATAAAAA GCTCAGTTAC TGATTTGCTG GGTTCATGGTC AAAATTCTTA CCTATTTATT TCATATCAAC TTTAAAAAAT AAATTACTTG CATTCTATAT ATTAATAATT GGGAAAGTAAT ATGCCTCAA TCAGTTTTAT ACTGGATTAT TCCCTATGCT TTAACCCT GCTCTCAATA AAACACTTCC TGATTAATGT TTGATTATA GATATTTTAG TCTTGTTGGG GATATTTTAG TCTTGTTGGG TTAGCCATGC TCTGAAGAAT CTGTGAAAGT ACAGTAAAGT TTTAATAAGC AATAAATGTA ACCTTTTATA TAAATCTCAG TGCTAGGTTA ACTTCTAATA AGCAGACGAA CATGTTACAT AAATTATAAT GTCTGTCTTG TAAAAAAGTT GAGGGGACTA AAAGTTTATG ACTCTGATAT GGAAGTTGTC ATATTAATAA ACTACATTTT AAAACATCAA ATATTTATAC TATTTGCTTT TCAAATAAAA GCATAGTGCT GTTTGGCATA</p>	
<p><i>PRRC2</i> <i>B</i></p>	<p>NM_0 13318. 3</p>	<p>GCAGATCGGG AGCGGTGCCG AGAAAAATTT CCTTACTAGA TGACATTTCA TCGCAATGTC CGATCGTTTG GGGCAAATTA CCAAGGGCAA GGATGGGAAA AGCAAGTACT CGACTCTCAG CCTGTTTGAT AAGTATAAAG GAAAATCAGT AGACGCGATT AGATCCTCAG TTATTCCTAG ACATGGCTTA CAGAGTCTTG GGAAAGTTGC TGCAGCCCGG CGCATGCCAC CGCCTGCAAA CCTGCCAAGC TTGAAGTCTG AAAACAAAGG AAACGACCCC AACATCGTGA TAGTACCCAA GGACGGGACG GGATGGGCAA ACAAGCAGGA TCAGCAAGAC CCAAAGAGTT CCAGTGCAC</p>	<p>35</p>

	<p>GGCCTCTCAG CCGCCGGAGT CGCTGCCGCA GCCGGGTTTG CAGAAATCTG TCTCCAATTT GCAGAAACCG ACACAGTCAA TCAGTCAGGA GAATACAAAT TCAGTGCCAG GTGGACCAA GTCATGGGCA CAGCTGAATG GAAAGCCAGT AGGACACGAA GGTGGTTTAA GGGGCTCAAG CCGACTGTTA TCCTTCTCTC CCGAGGAATT TCCGACGCTG AAAGCAGCTG GAGGGCAGGA CAAGGCTGGC AAAGAAAAGG GCGTCTTAGA TCTGTCTGAT GGGCCAGGAC CAAGCCTCCG CCCTCAGAAT GTGACAAGCT GGAGGGAGGG CGGTGGGCGA CACATAATTT CTGCCACGTC TCTGAGCACC TCCCAACTG AGCTGGGCAG CAGGAACTCG AGTACGGGAG ATGGAGCCC CTCCTCGGCA TGTACCAGCG ATTCTAAGGA CCCCTCTCTC CGCCCGGCTC AGCCTGTCCG AAAAGGGGCT TCACAGTTCA TGGGAAATGT ATACCACCCA CCTACATAACC ATGACATGCT TCCTGCTTTT ATGTGTTTCG CGAAGTCATC AGAAAACCAG GGTACAGTGG AACGAGGCTC TTTTCCCCTT CCTCAGCTCC GCCTTGAACC TCGAGTTCCT TTTAGACAGT TCCAGATGAA TGACCAAGAC GGAAAAGAAA ACAGGCTGGG ATGTGCTCGC CCACTCCGCC CACTAAGGCA GCTGGTGGAG CGGGCACCAC GGCCACCAT TATCAATGCG GAAAACCTGA AGGGCCTTGA CGATCTGGAC GCCGATGCCG ATGATGGCTG GGCAGGCCTC CATGAAGAAG TGGACTATT TGAGAACTG AAGTTCAGTG ATGATGAAGA GGAGGAAGAA GTTGTGAAGG ACGGCAGGCC AAAGTGGAAC AGTTGGGACC CTAGGAGGCA GCGGCAGTTG TCAATGAGCT CTGCAGACAG TGCGGACGCT AAGCGGACTC GAGAGGAAGG GAAGGACTGG GCTGAAGCAG TGGGTGCGTC CCGTGTGGTC CGAAAGGCGC CAGACCCTCA GCCACCGCCC AGGAAGCTTC ATGGCTGGGC ACCAGGCCCT GACTACCAGA AGTCATCAAT GGCAGCATG TTCCGGCAAC AGTCCATCGA GGACAAGGAG GACAAGCCCC CACCAAGGCA GAAGTTCATT CAGTCAGAGA TGTCCGAGGC GGTGGAGCGA GCCCGAAAGC GCCGGGAAGA AGAGGAGCGC CGAGCCCAGG AGGAGAGGCT GGCCGCCTGT GCTGCCAAAC TCAAGCAGCT GGACCAGAAG TGTAAGCAGG CACGAAAGGC AGGTGAGGCC CGGAAGCAGG CAGAGAAGGA AGTGCCCTGG TCTCCAAGTG CTGAGAAGGC ATCTCCCAG GAAAACGGCC CTGCTGTCCA CAAAGGCTCC CCAGAATTCC CTGCCAAGA GACCCCCACC ACATTCCCAG AAGAGGCACC CACAGTGTCC CCAGCAGTGG CACAGAGCAA CAGCAGTGAG GAAGAGGCCA GAGAGGCTGG GTCCCCTGCA CAGGAGTTCA AGTATCAGAA GTCCCTTCCT CCCCATTCC AGCGCCAGCA GCAGCAACAA CAGCAGGAGC AGCTGTACAA GATGCAGCAC TGGCAGCCGG TGTACCCCCC GCCGTCCCAC CCCAGCGCA CCTTTTACCC ACACCACCCC CAGATGTTGG GCTTCGATCC CAGGTGGATG ATGATGCCTT CCTACATGGA CCCACGTATC ACGCCACTC GGACCCCGGT GGACTTCTAC CCCTCCGCC TGCATCCCTC AGGACTGATG AAGCCCATGA TGCCCCAGGA GTCCCTCAAT GGGACAGGCT GTCGCTCTGA GGATCAGAAC TGTGTGCCCC CACTCCAAGA AAGAAAAGTG ACCCCCATCG ACTCACCCCC</p>	
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	<p>TGTGTGGAGC CCAGAGGGCT ACATGGCACT GCAGAGCAAG GGCTACCCGC TCCCGCACCC GAAGTCGAGT GACACCTTGG CTATGGACAT GCGTGTGAGG AATGAAAGCT CTTTCTCTGC CTCACTCGGA AGGGCAGGGG GCGTAAGTGC TCAGCGCGAT CTCTTTGAGG AGAGAGGGGA GGAGTACTTG AGTGCTTTTG ACAAGAAGGC CCAAGCAGAC TTTGACAGCT GTATCTCTTC TCAAAGAATA GGCCAGGAGC TTTTGTTC ACCCAAGAA AATGTTGAGG ATGCAGGTGC TCCTGGGGGT CACACCCAAA ACCTCAGGTG TTCCCATTTG GAGCCTGACT TTGTCCCAGA TGAGAAAAG CCAGAGTGTG GCAGTTGGGA TGTTAGCCAC CAGCCAGAGA CCGCTGACAC AGCCCATGGT GTTGAGCGGG AGACACCCCG GGAGGGGACG GCCTTTAACA TCTCCTCCTG GGACAAGAAC GGGAGCCCCA ACAAACAGCC ATCCTCGGAG CCTGAATGGA CTCCCGAGCC CCGGAGCTCC AGCAGCCAGC ACCCGGAGCA GACGGGCAGG ACCCGGAGGT CGGGACCCAT CAAGAAACCA GTCCTGAAAG CCCTCAAGGT GGAAGACAAG GAGAAGGAGC TTGAGAAGAT TAAGCAGGAG CTAGGGGAGG AGAGTACCCG GCTGGCCAAG GAGAAGGAGC AGAGCCCCAC GGCAGAAAAG GATGAGGACG AAGAGAACGA TGCTCTCTG GCCAACTCCT CCACCACCAC TTTGGAGGAC AAAGGCCCTG GCCATGCCAC TTTTGGCCGC GAGGCCACCA AATTTGAAGA GGAGGAGAAA CCTGACAAGG CCTGGGAAGC CAGACCCCCA CGAGAGTCCA GCGATGTTCC CCCCATGAAG AGAAATAACT GGATCTTTAT TGATGAGGAG CAAGCCTTTG GGGTCAGAGG ACAGGCCCGG GGCCGGGGCC GTGGTTTCAG AGAGTTCCT TTTCGTGGTC GGCCTGCTGG CGGAAATGGG AGCGGCCTCT GTGGTGGGGG GGTCTGGGG GCCCGCAGCA TCTACTGCAG CAGTCAGCGC AGCGGCCGTG GCCGGGGCCT GCGAGAGTTT GCGCGGCCAG AGGACTGCC CAGAGCCAAG CCCCAGCGGA GAGTTGCCAG TGAGACCCAT AGCGAGGGCT CAGAGTATGA AGAACTTCCC AAGCGCCGCC GGCAGAGGGG CTCCGAGAAC GGGAATGAAG GCTCGCTCCT GGAGAGGGGAG GAGAGCACCT TGAAGAAGGG CACTGCAGA GATTCTTGGC GGTCCAACAA GGGGTGCTCT GAGGACCACA GCGGTCTAGA TGCCAAGAGC CGAGGCCCTC GGGCCTTTGG GCGAGCCCTC CCTCCCCGGC TGAGCAATTG CGGGTATGGA CGGAGAACCT TCGTCTCCAA AGAGTCACCC CACTGGCAGA GCAAAGTCC AGGCAGCTCT TGGCAGGAAT ATGGCCCTTC CGACACATGC GGATCCCGGC GACCTACAGA CAGAGACTAT GTCCCAGATT CCTACAGACA CCCTGACGCA TTTGGTGGCC GGGGCTTTGA GGACAGCCGC GCGGAGGACA AGAGATCCTT CTTC AAGAT GAACACGTGG CAGATTCTGA AAATGCAGAG AACCGGCCCT TCAGGAGAAG GCGCCCCCA CGCCAAGATA AGCCCCCTCG ATTCCGGCGC CTCCGGCAAG AGCGGGAGTC CCTGGGCCTG TGGGGACCCG AGGAGGAGCC CCACCTGCTG GCAGGTCAGT GGCCAGGCAG GCCAAACTG TGTTCTGGGG ACAAGAGTGG CACTGTGGGC CGCAGGTCCC CTGAGCTCTC CTACCAGAAC TCCTCCGATC ACGCCAATGA GGAGTGGGAG ACGGCCTCCG AAAGCAGCGA</p>	
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	<p>CTTCAGCGAG CGGCGGGAGC GGCGGGAAGG CCCTGGGTCC GAGCCCGACT CCCAGGTGGA TGGTGGCCTG TCGGGGGCTA GTTTGGGTGA GAAGAAGGAG CTGGCCAAGA GGAGCTTCTC CAGTCAGAGA CCCGTGGTTG ACAGACAGAG CCGAAAGCTG GAGCCGGGAG GGTTTGGGGA GAAGCCCGTT AGGCCAGGTG GTGGTGACAC CTCCCCTCGC TATGAGAGCC AACAGAATGG GACGCCTTTG AAAGTGAAAA GATCCCCAGA CGAGGCCTTG CCTGGAGGTC TTAGTGGCTG CAGCAGTGGG AGTGGCCACT CCCCCTATGC CTGGAGCGG GCAGCCCATG CCAAGTGTGA CCTTCCCGAA GCCTCCAGTA AAAAGGCAGA GAAGGAGGCC AAGTTGGCTG CTCCGAGGGC AGGTGAACAG GGAGAGGCCA TGAAACAGTT TGACCTGAAC TATGGAAGTG CCATCATTGA AAATTGCGGG TCCAGCCCCG GGGAGGAGAG TGAGGTGGGT TCTATGGTGG GCGAAGGCTT CATCGAAGTC CTGACCAAGA AGCAGCGCCG CCTGCTGGAG GAAGAGAGAA GAAAGAAGGA GCAGGCCGTG CAGGTGCCTG TCAAAGGTCG AGGCCTTTC TCCCGTATTC CTCCTCGATT TGCAAAAAAG CAGAACAAC TATGTCTGGA GCAAGGTGAC GTGACCGTGC CTGGCAGCAG CCTGGGCACT GAGATCTGGG AGAGCAGCAG CCAGGCTCTC CCTGTGCAGG CCCCAGCCAA CCACTCCTGG AGGAAAGCTG TCACTGCCTT CAGCAGCACC GAGACTGGCT CTGCGGAGCA GGGTTTTAAG AGCAGCCAGG GAGATAGTGG CGTTGACTTG AGTGCCGAGT CTCGGGAGTC GTCTGCGACC TCCTCGCAGC GCAGCTCCCC ATATGGGACT CTGAAGCCAG AGGAGATGAG CGGGCCCCGC CTGGCGGAAC CCAAGGCCGA CAGCCACAAG GAGCAGGCTC CAAAGCCATC TGAGCAGAAG GATTCAGAAC AAGGCTCTGG ACAGAGCAAG GAGCACAGAC CAGGACCCAT CGGCAACGAG CGTTCTCTGA AAAACAGAAA GGGCTCGGAG GGGGCCGAGC GGCTGCAAGG GGCTGTGCTC CCGCCTGTTA ACGGGGTGGG GATTCACGTG GACTCCGTGC TGCCTGTGCC ACCCATTGAA TTTGGAGTCA GTCCAAAAGA CTCCGATTTT AGCTTGCCAC CTGGTTCTGC CTCTGGTCCT ACTGGGAGTC CAGTTGTTAA ACTTCAGGAT GCCTTGGCCA GTAATGCAGG GTTAACACAG AGTATCCCCA TCCTGCGGCG GGACCATCAC ATCCAGAGGG CCATCGGTCT CTCCCCAATG TCCTTCCCCA CCGCCGACCT TACTCTGAAG ATGGAGTCTG CGCGCAAGGC TTGGGAAAAC TCCCCAGTT TGCCGGAGCA GAGCTCTCCA GGCGGCGCTG GCTCAGGCAT CCAGCCTCCA TCCTCTGTGG GTGCCTCCAG CGGGGTCAAC TACAGCTCCT TCGGTGGAGT GTCCATGCCA CCCATGCCTG TGGCCTCTGT AGCACCTTCT GCTTCTATGC CAGGCAGCCA CCTCCCGCCC CTGTACCTGG ATGGCCATGT GTTTGCAAGT CAGCCCCGGC TGGTTCCTCA AACGATACCT CAGCAGCAGA GTTACCAACA GGCCGCCGCT GCCAGCAGA TCCCGATCTC CCTTCACACA TCTCTGCAGG CACAAGCTCA GCTTGGACTG AGGGGTGGGC TTCCTGTGTC CCAGTCCCAG GAGATCTTCA GCTCCTTGCA GCCCTTCAGA TCTCAGGTGT ACATGCACCC CAGCCTGTCA CCGCCAGCA CCATGATCCT CTCTGGGGGC ACAGCCTTGA AGCCTCCATA</p>	
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	<p>CTCGGCGTTC CCAGGCATGC AGCCCTTGGG GATGGTGAAG CCGCAGTCTG GCTCACCTA CCAGCCCATG AGCGGGAACC AAGCCCTGGT CTACGAGGGC CAGCTCAGCC AGGCTGCTGG CCTGGGTGCC TCCCAGATGT TGGACTCCA GCTCCACAG CTGACCATGC CACTGCCTCG GTACGGCTCC GGGCAGCAGC CACTGATCCT GCCCAGTCT ATTCAGCTGC CACCTGGGCA GAGCCTCTCC GTTGGGGCCC CCCGAAGGAT TCCTCCGCCC GGGTCCCAGC CGCCAGTCCT GAACACCAGC AGAGAGCCCT CTCAGATGGA GATGAAAGGC TTCCACTTTG CCGACAGTAA ACAGAATGTC CTTTCAGGAG GCCCCGTGCC ATCGCCACAG ACCTACAGGC CTAGCTCTGC TAGCCCCAGT GGGAAAGCCCT CTGGATCAGC AGTTAACATG GGCTCTGTGC AGGGACACTA CGTGCAACAG GCAAAACAAC GAGTGGATGA GAAACCAGC CTGGGAGCCG TGAAGCTGCA GGAGGCCCCC TCGGCTGCCT CCCAGATGAA GCGAACCAGG GCGATCAAGC CTCGGGCTGT CAAAGTGGAG GAGAGTAAGG CCTGACAGTG CCTGGCTGCC ACCTCGCCTC TCCCTACTGA GGACGGTGCC GCCATGCGGC CTCGACACAG CCGACACTCG GGAGCCTCAC CAGATCCACC GTCCAAATGC GTGGCCAGA CTGAGAGACC TCCCTCCTCT CCTCCTCCGA AAGCTCCGTT GTCAACCAGC TTGCACCCGT GGATATATGG CATTGACCCG CTTGCTTTGA TACGAAACAA AAAAGCAGAC GACTCCTTCA TCCCATCTGC TCCTACCGTG ACTGTGGAGT GACGCCTCCT GTGCAGTGCA GATTTGCCCT CCCTGCCTCC TCCCTGTCCT GCCGCGCAGC CAGGGCGCCT TCTCAGCAGT GCTTCCGGCC CAGCCGCCA TCCTAGGCA CAGTGATTTG GCAGCAGGGT CATTTTACTT TGAGGCTTTT TGTTTTAAAA TGTAGCCAAG GTTTTTACAA AGGGGAAAGG AAAAGAAAAC AAAAACGCAA GCTCCATGTG TATAGCTGAA CTTTTATATG TTCTTGCCA GCCCTCCGC TCCCTTCCAT CTCTAGCCTC TGTCCTGTTT AGTTTGATAC GTCACTGCAG TACCTTAAGA GGTGACTCTT AAGAATGCAT CCCCTCCTGA TTCCTCAGCT GGTTACCCT TGAGGTTATT TGCAAAAAGA AAAGGAGGTT CTTGAGGGCA CCGATTGCGA GCATTCTGGT GCCTGGCTCC CCGCCTGGGA AGCGATGGGG TGCTCAGAGC AGCAGGCAGG TTGGGGGAGG GGGGGGGTCA TAGTTGGGTT CCAGCTCCTG GCTTGATGAG CCCAGGGCGC TTACAGGCAG CCCATGAAGT TGATGACAGT TTTAGCATGA GAATCACACA GGGTCCCTGT CCTGGGCTCC TCTAAAGCCA GTGGATGTGC TGGGCACCAG AGACAAATCA TGGAGATGGC TGCTGGTGGC TCCCAGGTTG GCCCAGATGG GGTGAGCTGA CATAACACAG GCCCATCCA GGCCCCGTGG GCTCTGCTTC TGGGGCTCCA TACCCTGCC TGCAGGGGTG CTGTGTTTTT CACACATTC TTTCCCTGAA GCCTTCTGTA ACCTGTCATT TTCCTTCCTT CCTCTTCCGG AGCCTGCTGC TTTCTCTGGA CCTGTCTCCA CCTCCACAC AGCTCATCGT GAACACCACT TGGTGATGGA GGGAGTGGAC CCGTGTGTGG TCCCAAGTG AGGCCACTGG GAGTTTGCC TTTTCTCCT TTGCTTCACT CCCAGCAGCA GACCAGGTT GTCAGGACAG GAGGGCCTGA GCTAAGCAGT</p>	
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	<p> AGGCATCAGT CTCGTTTGTG TTCAGACGGC GGGGGCAGGT CCAGGGTGAG GCTGGGTGGA GGGCTGACCA AGGTCCAAAG GGCTTGTGTG AGCCATCGTG TGCTGGGCTT GTTTTTAAGT AAGAAACAAG GAAATCACTC CAGATTCTGT CATTCCAAGG AAAGGGAAGG GGACAGTTCA GGTTTCTCAG CTGTTCTTAG GGGTCACTGA GCGTCTACCT CCTCCTCCAG AGGAGGCTGG CTCAGAACAC CTAGAGGAGG GGGCCGGGGA TGCACCCCCC ACCAGAGGCT GCCTTCAGCG TCTCACGGGT GCAGGACAGC GCTCAGGCTT GGGCTCTAAG CTCTGTGTCT AGTGTAGAAC ATGGGGAAGG AGCATCTTAG GAACTGCTGA AGTAACTTCT TACTGCTCTC ACAATTCTAA GGAAGCGGGA GAACGGCCTC CTACCAACAG CGCCACCCC AGAGCTGCCT GGGAAAGGGC AGTTTTACTG AAAGGTGCTT TACTGTTTAC CTGCATCTTT CAGCAGCTCC CCTCCTGCCC TCACCTGGTC TTTTCCCTCT TTATCCCAAG CCTTTATGCT TGAGTCCCTT CCCCAGGGGC TGCCACCCG ACAGTTCAG GCATTCCCTA CCTGAGCTTC TTGTCTGCTT TTCCTTCTCC CACTGCAAGC GGCTGCTTGT GGGGCCTGGG ATGAGCCCTC TCTGTCCCA CCGGCCCTCC TTGCCAAGCC ATTCTGGGT GAGTTCAGGC CTGCGGGAGC CACACATTCA TCTCCACCTG GACACTTGAG CCGCATGGCC AGACCCCTCC CACCTGATGC GGTGGTGGCT GTGATTTGTC AAAAGAAAGC CTTCTGGATG CTGTAAAGAT GTACCCTTCA GGTGAACCTG GTATCAGACC CACAGTACTT GCTGTTTGAG AAAAAATAAA AACAAAAAGG TCACCTGTTT TCCAGCCCTT TTCTCTTACC TGGTATTTCC TTCCTTCTC CTCCCCACC CCAAATAAAA AAACAAAAAA CACTAGAATT TATTTATATG TATTGATGTT GTAGGTCTAG GTGAAAAAAA AAGAAGTAAA TGTTTCACTG CTCTATTTAT ATATAATGTC TGAATTAAT CTGTGCAGGA AAGGCCAGGA AATTGCATGT GAAGTTCGGT GCAGTCACCA CCTGTGTGTG ACCTGAGCTG CAGTCTCTTC GCTGAGATGC AGGTTTTAAA TGAGACTTGG GGGGCTGAGG GCAGGCCTCA GGCCTCCAG CGCCCCAACC CCTCCTTGGT CTAATGAAAT GCAGTCTTA GTGCAGAGAT GTTTTAAGGT GCAATATATC TCTTCTTTC CCGTGGTTTT AGAGCCAAGC TCAAGGTAGT AGGACGTAGG GTCTTATTTT GTTTTCAAAC CCCCATCCTC AGAGCGCAGA TACATGCAGA GGCTTCTGCC AGGATAACCAC GGGGCCTTAG TGGGAACAGG TGGAGACCAG CACTTCCCTT TCCTGCTGCT GAGGTAGGGA TTGGGGGGTC AGAACCCACT CACTTTTGCC TGTTAAAGTT GCCCTCCTGA CGCTGGCAGC TCTGCCTTGG TCACTGGGGA TCGGGCTCGT TGCTCAGCCA CCAGTGGCCT TCGGGTATTG TCCACCATCC ACTAGAGTGG GATGAAGTCC AGAGTGTGGG TATACATCTC AGATGCCCAT CTACCCACTG GGGACTTCAA TGCCAGCTGC ATTTGGTTTG GTTTTCTTAA CTGTTGGCTT CTCCCCACAG CGTTTTTGT TTTTTTTAA ACATTCATAT TGTTTTCAA CTTGGAATTC ATAGACACTC TGGCTCTAGG TTCCTTAAAGG GGGAAAACA AAGATGACTT TATTTACAT TCAAGAAAAT </p>	
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		<p>CAGTTCAGTT CCAAAGCTGT GGTCCCTTCCA GCCACTTCTA GGGACACTGG GGAACCTTGT TAAACGTTGA CATCAGTGCT CTCCAGCCGT GCTGTCACCC TCCTATCTTC TGGATCTGCC TTCGCGATGG TCAGTGACAG CTTCTGGAAG CTGAGCACAC ACAGGTGCAC AGCCATGCTG TGGTCTGGCC TGCTACGGCA GCATGGCAGC TCTGGTGGAG CCTTCTCCCT TGCCATTTGG TTCCCTGTG CCAAGTAGCT GCAGGCTGCC CCTCAAATCT TCATTTGTCC CTTTTCACCTT CCTGCAGAAC AAGCCTGGGT TAGAGGGTCT GCTGGAAATG GCCTTTGAAG ACCAAGGATA CCAGGATGTG TGCACCTCTGT CGTGTCTGT GATGAATGGG AAACGTAGGC TTCCAGAAAG CCAGCTCTCT TCTGAAATGT GACGGACCTA AGCAGGAAGT CATCCAGGAC AGGAGTGGCT CAGTGTGGG GATGGACGCT GTCGCCCAGC CATGCTCCAC CAGGGCCACC AATGTGTAGT TGGCTGGTGG TCTTCGGGCA TGTGAGACCT GCTCTTCACT GTTTCACCC CACTTGGTGG CCTCCAGGAT GGTAGTGGCA CCCTCAGAGC CCCATCTTCA GCATGTTCTG AAGCCTCAGA GTGGAAATTC CTGCTAAGGC TCTGTGTGGA CGCCTTCTC CCGTGATCTA AAGGGGACAC TGTACTCAAG CTTTTGACCT CATGCCTTGT GTAGTAAAA AGGATTTGGG GGTTTTGTGTT GGTTCTGAG AGGGTTGTGT TTTGTHTTTG TTTCTTTTG TTTATGTTTT GGCCTTTCCT CTTTGTCTTT CCATGTAGAC CAGATATTTG AAAGGGCAGA CGATGGCTAG AGGTGTAATG TGCAGCTTGT TTATACGGTA TTTTGGGAAA CTTACCTTGG ATGGGAAATC GAATCGTGGA TTCACCAGGC CGGTGCTGGC AACTCACC TCGCCCTTTC CCTCCGGTTC AGTACCTATT GTTCTCCTT TCAAATATGT GATTGTAATA GCTCTTCCA TATGAAAGAA TTCTCCTTAT TTAAATAAAA AAAGTTTAAA AA</p>	
<p><i>DOPEY</i> 2</p>	<p>NM_0 05128. 3</p>	<p>TCCCACAGTG CCTGGCCCAG AAGCCTTGCT AAATATTTGA ACAGGATTGC CCAATACTTT TCTGCTGTGA GAATGTAAGA TGGATCCAGA AGAGCAGGAG CTCTTAAATG ATTACAGATA CAGAAGCTAC TCTTCAGTGA TTGAAAAGGC TTTGAGAAAT TTTGAGTCCT CGAGTGAATG GGCGGATCTC ATATCTTAC TTGGCAAACCT CAACAAGGCT CTTCAGAGTA ACCTGAGGTA CTCCTTGTTG CCAAGACGGC TCCTCATCAG CAAAAGATTA GCTCAGTGTG TGCACCCTGC CCTGCCCAGT GGTGTCCACT TAAAAGCTCT GGAAACCTAC GAGATTATCT TAAAATCGT GGGGACCAA TGGCTGGCCA AGGACTTGTT TCTGTACAGC TGCGGGTTAT TTCCTCTCCT GGCACACGCG GCGGTGTCCG TGAGGCCGGT GCTGCTCACC CTGTACGAGA AGTACTTCT CCCCTGCAG AAGCTGCTCC TGCCAGTCT GCAGGCCTTC ATCGTGGGCC TGCTGCCCGG CCTTGAAGAG GGCTCCGAGA TCTCCGACAG AACGGATGCT CTGCTCCTGA GACTGTCCGT GGTGGTTGGC AAAGAGGTGT TTTACACCGC CCTCTGGGGG AGCGTCTGG CCAGCCCGTC CATCCGCCTC CCTGCCTCAG TCTTCGTGGT GGGCCACATC AACAGGGATG CCCCCGGCCG</p>	<p>36</p>

	<p>GGAGCAGAAG TACATGCTGG GGACCAATCA CCAACTCAGG GTGAAGTCTT TCGTGCCCTC CCTGTTGGAC TCAAATGTTT TTGTGCAAAG AAATAATCTG GAAATCGTTC TGTTTTTCTT CCCATTTTAT ACCTGTCTGG ATTCCAATGA GAGAGCCATC CCCCTCCTCA GATCTGACAT CGTGCGCATT CTCTCAGCCG CCACCCAGAC CCTACTGAGA AGGGACATGT CCCTGAACAG AAGACTGTAT GCATGGTTAC TAGGCTCAGA CATAAAAGGA AATACCGTTG TGCCAGAATC TGAAATCTCA AATTCTTATG AAGACCAGTC GTCTTATTTT TTIGAAAAAT ACTCCAAGGA TCTTTTAGTT GAGGGTTTGG CTGAGATATT GCATCAGAAG TTCATAGATG CTGACGTGGA GGAACGCCAT CATGCATACC TGAAGCCTTT TCGCGTCCTC ATCAGTCTGC TTGACAAGCC AGAAATAGGG CCTCAAGTGG TTGGGAATTT GTTCTCGAA GTCATCAGGG CCTTTTATTC TFACTGCAGA GATGCCCTTG GCTCTGATCT TAAACTTAGC TACACCAGA GTGGAAATTC GCTGATAAGT GCAATCAAGG AAAACAGAAA TGCCTCTGAG ATTGTCAAAA CGGTAAATTT GCTGATAACT TCTCTAAGCA CAGACTTTCT CTGGGATTAT ATGACAAGGT GTTTTGAGGA ATGCTTTAGA CCAGTGAAGC AGCGTTACAG CGTGAGGAAC AGCGTCAGCC CTCCCCCAC GGTCTCGGAG CTCTGCGCCC TCCTGGTCTT CCTGCTGGAT GTCATTCCTT TGGAACTTTA CTCTGAGGTG CAAACCCAGT ATCTCCCTCA GGTGCTCGGC TGCTGGTGC AGCCTCTTGC TGAGGACATG GAGGCCTTAA GTTTACCTGA ACTCACGCAT GCCTTGAAGA CGTGTTTCAA GGTGCTCAGC AAAGTCCAGA TGCCTCCTTC CTACCTCGAC ACGGAGTCCA CCAGCGGAAC CTCGAGTCCA GTAAAAGGTG AAAACGGCAA AATAATTTTG GAAACAAAGG CAGTGATTCC CGGTGACGAA GATGCTTCGT TTCCCCTCT GAAGTCTGAG GACAGTGGGA TCGGGCTCAG TGCCTCGTCA CCGGAGCTCT CTGAGCACTT GAGGGTTTCT CGAGTTTCTC TGGAAAGGGA CGACGTTTGG AAGAAGGGCG GGAGCATGCA GAGGACGTTT CTTTGCATCC AAGAGCTAAT CGCCAACCTT GCCAGCAAGA ACATTTTTGG AGTACAGCTG ACAGCGTCAG GAGAAGAAAG CAAGTCCGAG GAGCCTGCAG GGAAGAGGGA CAGGGATGGG ACGCAGAGCC TGGCAGCCAA TGATTCCAGC AGGAAGAACT CTTGGGAGCC CAAGCCATC ACTGTGCCTC AGTTCAAGCA GATGCTGTCA GACTTGTTCA CAGCACGAGG GTCTCCATTC AAGACAAAAA GTTCAGAGTC ACCATCGTCT TCGCCAGCA GCCCTGCCAG GAAAAACGGG GGAGAATGGG ATGTTGAGAA GGTGGTCATT GACCTGGGGG GTTCCAGGGA GGAACGCAGG GAGGCCTTTG CCGCCGCTG CCACCTGCTG CTGGATTGTG CCACTTTCC TGTCTACCTG TCCGAGGAAG AGACCGAGCA GCTCTGTGCA ACGCTCTTCC AGCTGCCAGG AGCCGGTGAT TCCAGTTTTT CATCTTGGCT GAAGTCCCTC ATGACTATTT GCTGCTGTGT GACTGACTGC TACCTCAGA ACGTGGCCAT TTCCACTCTG CTGGAAGTGA TAAACCATT CCAGTCCCTG GCGCTTGTC TTGAAGACAA GATGAAACGC TATAAGAGCT CTGGACACAA CCTTTTTTTT GGCAAGCTGC AGATGGTGAC</p>	
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	<p>GGTTCCTCCC ATTGCTCCAG GGATATTGAA AGTCATTGCA GAGAAAACAG ATTTCTATCA GAGGGTGGCT CGTGTGCTTT GGAATCAGCT GAACAAAGAG ACCCGGGAGC ATCACGTAC CTGCGTAGAA TTGTTCTACC GGCTGCACTG CCTGGCCCCT ACGGCCAACA TCTGCGAGGA CATCATCTGC CATGCCCTCC TGGACCCTGA CAAGGGAACA AGGCTGGAAG CTCTGTTTAG ATTTCCGTG ATCTGGCATC TGACAAGAGA GATCCAAGGC AGTCGAGTAA CATCTACAA TCGCTCCTTT GATAGGTCT TGTTTGTGCGT GCTGGACAGC CTGGCCTGCA CGGATGGTGC CATCGGTGCG GCAGCCCAGG GCTGGCTGGT GCGTGCCTC TCCCTCGGGG ACGTGGCTCG CATCCTCGAA CCCGTGCTCC TGCTGCTGCT GCAGCCAAAA ACCCAGAGAA CCTCCATCCA CTGCCTCAAG CAGGAGAACT CGGCCGATGA CTTGACCGT TGGTTTAACA GGAAGAAAAC CTCTTTCAGA GAGGCATGCG CAGTGCCCGA GCCTCAGGAG AGCGGCTCTG AAGAGCACCT GCCTCTGAGC CAGTTCACCA CAGTGGACCG TGAAGCCATT TGGGCCGAAG TGGAGAAGGA GCCCGAGAAG TACCCGCTGC GAGGCGAGCT GAGCGAGGAA GAGCTGCCCT ACTACGTGGA GCTTCCAGAC AGGACGGCCC ACGGCGCCCC GGACAGCAGC GAGCACACCG AGTCTGCAGA TACAAGCTCC TGCCACACGG ACAGCGAGAA CACGTCTCC TTCTCCTCCC CTTCCCACGA CCTGCAGGAG CTGAGCAACG AAGAGAAGTCT CTGTGCACCC ATCCCCATGG GGGGCAGGGC GTACCCCAAG CGCTCGGCC TGCTGGCGGC CTTCAGTCA GAAAGCTTCA AGGCTGGGGC CAAGTTAAGC CTGGTGCGGG TGGACTCGGA CAAGACGCAG GCTTCTGAGT CGTTCTCCAG CGACGAGGAG GCGGACTTGG AGCTCCAGGC CCTCACCACA TCCAGGCTGC TAAAGCAGCA GCGGGAAAGG CAGGAGGCCG TCGAGGCCTT GTTCAAGCAC ATCCTGCTCT ACCTGCAGCC CTACGACTCT CGGCGGGTCC TCTATGCCTT CTCGGTGTG GAGGCTGTGC TAAAACCAA CCCTAAGGAA TTCATCGAGG CTGTGTCCAG GACTAGCATG GATACCAGCT CCACCGCGCA CCTCAACCTC ATCTCCAACC TCCTCGCTCG CCACCAGGAG GCCCTCATTG GCCAGAGTTT CTACGGAAAG CTCCAGACCC AGGTCCCCAA CGTGTGCCCC CACTCTCTGC TCCTGGAGCT GCTCACCTAC CTCTGCTGA GCTTCCTGCG CTCCTACTAC CCTTGCTATT TGAAGGTCTC GCACCGAGAC ATTCTCGGCA ACCGGGACGT GCAGGTCAAA AGTGTCGAGG TTTTGATCAG GATAATGATG CAGCTGGTCT CAGTGGCCAA GTCTTCGGAA GGAAGAACG TGGAGTTCAT CCACAGCTTG CTGCAGAGGT GCAAAGTTCA GGAGTTTGT CTGCTCTCCC TGTCGGCGTC CATGTACACG AGCCAGAAGC GCTACGGGCT GGCCACCGCC CACCACGGCA GGGCCCTGCC AGAGGACAGC CTCTTTGAGG AGAGTCTCAT TAAC TTGGGT CAGGACCAGA TCTGGAGTGA GCACCCGCTG CAGATTGAGC TGCTGAAGCT GCTGCAGGTG CTGATTGTCT TGGAACACCA CCTGGGTCGG GCCCATGAGG AGGCGGAAAA CCAGCCCAGC CTGTCCCAGG AGTGGCAGAG AGCCCTGAAC TTCCAGCAGG CCATCAGCGC CCTGCAGTAC GTGCAGCCCC ACCCCCTCAC</p>	
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	<p>CTCCCAGGGT CTTCTGGTCT CTGCGGTGGT GAGGGGTCTG CAGCCCGCCT ACGGTTACGG CATGCATCCG GCCTGGGTGA GCTTGGTCAC GCATTCCTTG CCCTACTTCG GAAAGTCCCT GGGCTGGACG GTGACACCCT TTGTTGTCCA GATTTGCAAA AACTTGGATG ACTTGGTCAA GCAGTATGAA AGCGAATCTG TGAAGCTCTC TGTCAGCACA ACCTCCAAGA GGGAAAACAT TTCTCCAGAT TATCCACTCA CCCTTCTAGA AGGTCTAACG ACCATTAGTC ATTTTTGTCT TTTGGAACAA GCCAACCAAA ACAAAAAGAC CATGGCTGCA GGTGATCCTG CCAACTTGAG GAATGCCAGA AATGCCATTT TGGAAGAGCT GCCTCGAACT GTTAACACCA TGGCCCTTCT CTGGAATGTT CTCAGAAAGG AGGAGACTCA AAAGAGACCT GTCGATCTCC TAGGGGCCAC GAAGGGATCC TCTTCCGTTT ACTTTAAAAC CACCAAAACC ATAAGACAAA AAATTTTAGA CTTCTTAAAC CCCTTGACGG CCCATCTTGG GGTTTCAGTTG ACAGCGGCTG TTGCGGCAGT GTGGAGCAGA AAGAAAGCCC AGCGTCACAG TAAGATGAAG ATTATCCCAA CGGCAAGTGC ATCCAGCTA ACCCTTGTCG ACTTGGTGTG TGCACTCAGC ACCCTGCAGA CTGACACGCT GCTGCACCTG GTGAAGGAGG TGGTGAAGAG GCCACCCCAA GTCAAAGGGG GTGATGAGAA ATCGCCCCTA GTGGACATTC CTGTGTTGCA GTTTTGCTAT GCTTTTCTCC AAAGGCTCCC AGTACCAGCC TTGCAAGAGA ACTTTTCTTC ACTGTTGGGA GTATTGAAAG AGTCTGTACA GTTGAATCTA GCCCACCTG GGTATTTTCT GCTTCTCAGC ATGCTGAATG ACTTTGTAAC AAGAACTCCC AACCTGGAAA ACAAGAAGGA CCAAAAAGAC CTGCAGGAAA TCACTCAGAA AATCCTAGAA GCTGTGGGGA ACATTGCCGG CTCTTCCTTG GAGCAAACCA GCTGGCTAAG CAGAAACCTG GAAGTGAAGG CCAACCTCA GGCCTCTCTA GAAGAATCTG ATGCTGAGGA GGACCTGTAT GATGCTGCTG CAGCTTCAGC AATGGTGTCT TCATCCGCCCG CGTCGGTGTA CAGCGTGCAA GCCCTCTCTC TCCTGGCAGA GGTACTGGCT TCCCTCCTGG ACATGGTTTA TCGAAGTGAT GAGAAGGAGA AAGCTGTGCC GTTAATCTCC CGTCTGCTTT ACTATGTTTT TCCATACTTA CGCAACCACA GTGCCTACAA TGCTCCCAGC TTCCGGGCTG GCGCTCAGCT GCTGAGCTCC CTGAGTGGCT ATGCCTACAC AAAGCGAGCC TGGAGGAAGG AGGTCCTGGA GCTGTTTCTC GACCCCGCTT TCTTTCAGAT GGATACTTCC TGTGTTTATT GGAAGTCCAT TATTGACCAT CTTTTGACTC ATGAGAAAAC AATGTTTAAAG GATTTAATGA ACATGCAGAG CAGTTCTTTG AAATATTCT CAAGTTTTGA ACAGAAAGCC ATGCTGTTAA AGCGCCAGGC TTTTGCTGTC TTCAGTGGAG AACTTGATCA ATACCACCTT TACCTTCCAC TGATACAAGA ACGCCTGACA GACAATCTCA GAGTTGGACA GACATCCATA GTTGCTGCTC AGATGTTTCT TTTTTTCAGA GTTTTGCTGC TAAGAATATC TCCTCAACAT TTGACTTCAT TGTGGCCAAT AATGGTCTCT GAATTGATTC AGACATTCAC ACAGCTTGAA GAAGATCTAA AAGATGAAGA TGAGTCATIG AGAAGCACCA ACAAAGTAAA CAGAACGAAA GTTTCAGTCC CGGATGCAAA</p>	
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		<p>TGGACCTCA GTGGGGGAGA TACCCAGAG TGAACATC TTGTATTAT CAGCTTGCAA ATTCTTGGAC ACAGCGCTTT CTTTTCCACC TGACAAGATG CCATTATTTT AAATTTATAG GTGGGCATTT ATTCCAGAAG TGGACACAGA GGGCCCTGCC TTCCTGTCGG ATGTAGAGGA GAATCACCAA GAATGCAAAC CCCACACTGT CAGGATTCTA GAACTTCTAA AATTAAAGTT TGGGGAAATC AGTAGCTCTG ATGAGATCAC CATGAAGAGT GAATCCCCGC TTCTGCGCCA ACATTCTGTT TCCAGCATCA GGCAGTTGAT GCCATTCTTC ATGACTCTAA ATGGTGCATT TAAGACCCAG AGACAGCTGC CTGCTGATAG CCCAGGAACT CCATTCTTGG ACTTTCCTGT CACAGATAGC CCAAGGATCT TAAAACAACCT GGAAGAATGC ATCGAATATG ATTTTCTGGA ACATCCAGAA TGTTAACCAT GTGAGAGAGA ATATGTTTAA TCCATGTATT GGTACTTTAC TGAAAACCAG GTTATATTCT AAAGAAGAAA GAAGGCAGGA TAGTGCTTTT GAACAAGCCT ATTTCCATTT TGAAAGTAGA TTTTCAAGGCTA GGTGCGGTGG CTCACACCTG TAATCTCAGC ACTTTGGGAG GCCAAGGCAG GCAGATCACT TGAGGTCAGG AGTTCGAGAC CAGCCTGACC AACATGGTGA GACCCTGTCT CTAATAAAAA TACAAAAATT AGCTGGGTGT GGTGGCGGCG CCTGTAATCC CAGCTACTTG GGAGGCTAAG GCATGAGAAT TGCTTGAACC CAGGAGGTGG AGGCTGCAGT GAGCCGAGAT CACGACACTG CACTCCAGCT GTGTGACAGA ATGAGACCAT CTCCAAAAAA AAAAAAAGT AGATTTTCTA TAATTTACTG TTCAGCAACA GGACACACCT CCCTAAATGC CTTGTAATAT ATTTGAATCT GATTCTGCAT TTCTTCTCA ATTTATGTAA TGAAAATAAA ATTAATATAT CATCTAACAG TAGCACAATA TTTGTAATAT GAAGTAAAGT ATGAAGATAA TGAAGAAGTT GTTTTCTTTG TTGAAGCAGT TATATGGGTC TTTCTCAGTA TATTTCTCTT TTCTCTAAAA GTTTAAACTT ATTAAGAAGAA TGTTATTTTT AACTTTCAA AAAAAAAA</p>	
<p><i>NDUFB</i> 11</p>	<p>NM_019056.6</p>	<p>GCTCTGGCCG GCCCCGGCGA TTGGTCACCG CCCGCTAGGG GACAGCCCTG GCCTCCTCTG ATTGGCAAGC GCTGGCCACC TCCCCACACC CTTTGCGAAC GCTCCCCTAG TGGAGAAAAG GAGTAGCTAT TAGCCAATTC GGCAGGGCCC GCTTTTTAGA AGCTTGATTT CCTTTGAAGA TGAAAGACTA GCGGAAGCTC TGCTCTTTT CCCAGTGGGC GAGGGAAGCTC GGGGCGATTG GCTGGGAAGT GTATCCACCC AAATGTCACC GATTTCTTCC TATGCAGGAA ATGAGCAGAC CCATCAATAA GAAATTTCTC AGCCTGGCCG AAAATGGTTG GCCCCACGAA GCCACGACAA CTGGAGGCAA AGAGGGTTGC TCAACGCCCC GCCTCATTGG AAAACCAAAT CAGATCTGGG ACCTATATAG CGTGGCGGAG GCGGGGCGAT GATTGTCGCG CTCGCACCCA CTGCAGCTGC GCACAGTCGC ATTTCTTTCC CCGCCCCTGA GACCCTGCAG CACCATCTGT CATGGCGGCT GGGCTGTTTG GTTTGAGCGC TCGCCGTCTT TTGGCGGCAG CGGCGACGCG AGGGCTCCCG</p>	<p>37</p>

		<p>GCCGCCCGCG TCCGCTGGGA ATCTAGCTTC TCCAGGACTG TGGTCGCCCC GTCCGCTGTG GCGGGAAAGC GGCCCCCAGA ACCGACCACA CCGTGGCAAG AGGACCCAGA ACCCGAGGAC GAAAACCTTGT ATGAGAAGAA CCCAGACTCC CATGGTTATG ACAAGGACCC CGTTTTGGAC GTCTGGAACA TGC GACTTGT CTTCTTCTTT GCGTCTCCA TCATCCTGGT CCTTGGCAGC ACCTTTGTGG CCTATCTGCC TGA CTACAGG TGCACAGGGT GTCCAAGAGC GTGGGATGGG ATGAAAGAGT GGTCCCGCCG CGAAGCTGAG AGGCTTGTGA AATACCGAGA GGCCAATGGC CTTCCCATCA TGGAATCCAA CTGCTTCGAC CCCAGCAAGA TCCAGCTGCC AGAGGATGAG TGACCAGTTG CTAAGTGGGG CTCAAGAAGC ACCGCCTTCC CCACCCCTG CCTGCCATTC TGACCTCTTC TCAGAGCACC TAATTAAGG GGCTGAAAGT CTGAAAAAAAA AAAAAAAAA</p>	
<p>ND4</p>	<p>NC_01 2920.1</p>	<p>ATGCTAAAATAATCGTCCCAACAATTATATTACTACCACTGAC ATGACTTTCCAAAAACACATAATTTGAATCAACACAACCACCC ACAGCCTAATTATTAGCATCATCCCTCTACTATTTTTTAACCAA TCAACAACAACCTATTTAGCTGTTCCCAACCTTTTCTCCGACC CCCTAACAACCCCTCCTAATACTA ACTACCTGACTCCTACCC CTCACAATCATGGCAAGCCAACGCCACTTATCCAGTGAACCACT ATCACGAAAAAAAACTCTACCTCTCTATACTAATCTCCCTACAAA TCTCCTTAATTATAACATTCACAGCCACAGAACTAATCATATTT TATATCTTCTTCGAAACCACACTTATCCCCACCTTGGCTATCATC ACCCGATGAGGCAACCAGCCAGAACGCCTGAACGCAGGCACAT ACTTCTATTCTACACCCTAGTAGGCTCCCTTCCCTACTCATCG CACTAATTTACACTCACAAACACCCTAGGCTCACTAAACATTCTA CTACTCACTCTCACTGCCCAAGAACTATCAAACCTCTGAGCCAA CAACTAATATGACTAGCTTACACAATAGCTTTTATAGTAAAGA TACCTCTTTACGGACTCCACTTATGACTCCCTAAAGCCCATGTC GAAGCCCCCATCGCTGGGTCAATAGTACTTGCCGCAGTACTCTT AAAAC TAGGCGGCTATGGTATAATACGCCTCACACTCATTCTCA ACCCCTGACAAAACACATAGCCTACCCCTTCTTGTACTATCC CTATGAGGCATAATTATAACAAGCTCCATCTGCCTACGACAAAC AGACCTAAAATCGCTCATTGCATACTCTTCAATCAGCCACATAG CCCTCGTAGTAACAGCCATTCTCATCCAAACCCCTGAAGCTTC ACCGGCGCAGTCATTCTATAATCGCCCACGGGCTTACATCCTC ATTACTATTCTGCCTAGCAAACCTCAAACCTACGAACGCCTCACA GTCGCATCATAATCCTCTCTCAAGGACTTCAAACCTTACTCCCAC TAATAGCTTTTTGATGACTTCTAGCAAGCCTCGCTAACCTCGCC TTACCCCCACTATTAACCTACTGGGAGAACTCTCTGTGCTAGT AACCACGTTCTCCTGATCAAATATCACTCTCCTACTTACAGGAC TCAACATACTAGTCACAGCCCTATACTCCCTCTACATATTTACC ACAACACAATGGGGCTCACTACCCACCACATTAACAACATAA AACCTCATTACACGAGAAAACCCCTCATGTTCATAACCTA TCCCCATTCTCCTCTATCCCTCAACCCCGACATCATTACGGG TTTTCTCTT</p>	<p>38</p>

<p><i>MORF4</i> <i>L1</i></p>	<p>NM_0 012656 03.1</p>	<p>CGGCGTGCCC TGGGGCGGCG CGGGCGCAGG GGC GCGTGCG CGGCGGGCTG TCGTTGGCTG GAGCAGCGGC TGC GCGGGTC GCGGTGCTGT GAGGTCTGCG GCGCTGGCA AATCCGGCCC AGGATGTAGA GCTGGCAGTG CCTGACGGCG CGTCTGACGC GGAGTTGGGT GGGGTAGAGA GTAGGGGGCG GTAGTCGGGG GTGGTGGGAG AAGGAGGAGG CGGCCAATCA CTTATAAATG GCGCCGAAGC AGGACCCGAA GCCTAAATTC CAGGAGGTTG GGATGAATGG GTTCCGGAGA GCAGAGTACT CAAATACGTG GACACCAATT TGCAGAAACA GCGAGAACTT CAAAAAGCCA ATCAGGAGCA GTATGCAGAG GGAAGATGA GAGGGGCTGC CCCAGGAAAG AAGACATCTG GTCTGCAACA GAAAAATGTT GAAGTGAAA CGAAAAGAA CAAACAGAAA ACACCTGGAA ATGGAGATGG TGGCAGTACC AGTGAGACCC CTCAGCCTCC TCGGAAGAAA AGGGCCCGG TAGATCCTAC TGTTGAAAAT GAGGAAACAT TCATGAACAG AGTTGAAGTT AAAGTAAAGA TTCCTGAAGA GCTAAAACCG TGGCTTGTTG ATGACTGGGA CTTAATTACC AGGCAAAAAC AGCTCTTTTA TCTTCCTGCC AAGAAGAATG TGGATTCCAT TCTTGAGGAT TATGCAAATT ACAAGAAATC TCGTGGAAAC ACAGATAATA AGGAGTATGC GGTTAATGAA GTTGTGGCAG GGATAAAAAGA TACTTCAAC GTAATGTTGG GTACCCAGCT ACTCTATAAA TTTGAGAGAC CACAGTATGC TGAAATTCTT GCAGATCATC CCGATGCACC CATGTCCAG GTGTATGGAG CGCCACATCT CCTGAGATTA TTTGTACGAA TTGGAGCAAT GTTGGCTTAT ACACCTCTGG ATGAGAAGAG CCTTGCTTTA TTAACAATT ATCTTCACGA TTTCCTAAAG TACCTGGCAA AGAATTCTGC AACTTTGTTT AGTGCCAGCG ATTATGAAGT GGCTCCTCCT GAGTACCATC GGAAAGCTGT GTGAGAGGCA CTCTCACTCA CTTATGTTG GATCTCCGTA AACACATTTT TGTTCTTAGT CTATCTCTTG TACAAACGAT GTGCTTTGAA GATGTTAGTG TATAACAATT GATGTTTGT TCTGTTTGA TTTTAAACAG AGAAAAATA AAAGGGGGTA ATAGCTCCTT TTTCTTCTT TCTTTTTTTT TTTCATTTCA AAATTGCTGC CAGTGTTTTT AATGATGGAC AACAGAGGGA TATGCTGTAG AGTGTTTTAT TGCCTAGTTG ACAAAGCTGC TTTTGAATGC TGGTGGTTCT ATTCCTTTGA CACTACGCAC TTTTATAATA CATGTTAATG CTATATGACA AAATGCTCTG ATTCCTAGTG CCAAAGGTTT AATTCAGTGT ATATAACTGA ACACACTCAT CCATTTGTGC TTTTGTTTT TTTTATGGTG CTFAAAGTAA AGAGCCCATC CTTTGCAAGT CATCCATGTT GTTACTTAGG CATTTTATCT TGGCTCAAAT TGTTGAAGAA TGGTGGCTTG TTTCATGGTT TTTGTATTIG TGTCTAATGC ACGTTTTAAC ATGATAGACG CAATGCATTG TGTAGCTAGT TTTCTGGAAA AGTCAATCTT TTAGGAATTG TTTTTCAGAT CTTCAATAAA TTTTTCTTT AAATTTCAA GAACAAAAA AAAAAAAA</p>	<p>39</p>
<p><i>MRPL1</i> <i>9</i></p>	<p>NM_0 14763. 3</p>	<p>GTAGTCTTGA CGTGAGCTAG CTGGCATGGC GGCCTGCATT GCAGCGGGGC ACTGGGCTGC AATGGGCCTA GGCCGGAGTT TCCAAGCCGC CAGGACTCTG CTCCCCCGC CGGCCTCTAT</p>	<p>40</p>

	<p>CGCCTGCAGG GTCCACGCGG GGCCTGTCCG GCAGCAGAGC ACTGGGCCTT CCGAGCCCGG TCGGTTCCAA CCGCCGCCGA AACCGGTCAT CGTGGACAAG CACCGCCCGG TGAACCCGA ACGCAGGTTT TTGAGTCTG AATTCATTCC TCGAAGGGGA AGAACAGATC CTCTGAAATT TCAAATAGAA AGAAAAGATA TGTTAGAAAAG GAGAAAAGTA CTCCACATTC CAGAGTTCTA TGTTGGAAGT ATTCTTCGTG TTAACACAGC TGACCCATAT GCCAGTGGAA AAATCAGCCA GTTTCTGGGG ATTTGCATTC AGAGATCAGG AAGAGGACTT GGAGCTACTT TCATCCTTAG GAATGTTATC GAAGGACAAG GTGTGAGAT TTGCTTTGAA CTTTATAATC CTCGGGTCCA GGAGATTCAG GTGGTCAAAT TAGAGAAAACG GCTGGATGAT AGCTTGCTAT ACTTACGAGA TGCCCTTCCT GAATATAGCA CTTTTGATGT GAATATGAAG CCAGTAGTAC AAGAGCCTAA CAAAAAAGTT CCTGTAAATG AGCTGAAAGT AAAAATGAAG CCTAAGCCCT GGTCTAAACG CTGGGAACGT CCAAATTTTA ATATTAAGG AATCAGATTT GATCTTTGTT TAACTGAACA GCAAATGAAA GAAGCTCAGA AGTGGAATCA GCCATGGCTT GAATTTGATA TGATGAGGGA ATATGATACT TCAAAAATTG AAGCTGCAAT ATGGAAGGAA ATTGAAGCGT CGAAAAGGTC TTGATTCTGA GAATGAATTT GGTTAGTTGC AGAAGATACA TTGGCTCTAA GAGGATATAT TTTGAGACCA ATTTAATTTT ATTTATAAGA ACATAGTAAT TAAGTGAACT AAGCATTTCAT TGTTTTATTA ATACTTTTTT TCTAAAATAA AACTTGTAACA CCAGTTTATT ACTCTAAAAA GAGAATTACA CATGCCAAAT GGACCAATGT CCATTTGCTT ATTGGAGGCA AAGCTACAAT AGAAGTCAGA GCATCACCAG AATGGTCTTT AATGAGCATG GAACCTGAGC AAAGGGAATA GGTGGGATGA ATTTTTTTTT TAATTGTGAA ACAATTCATA AGCACAATAT GATTACAGA ATAATAAACA TTCATGTACC CACTATCAGG TTAAGAAATA GAACATTTAT TAATATGTAG GAATGTTAAG AAATAAAACA TTTAATAAGA TCTCAGAAGA CTCCAGTAAA TCTGCAATTG TATCTCTCTC CTTTTAAAT GTAAATATCA TCTTGACTTG TTAATTATTC CTTGCAATTT CTTTTAGTTT ACTGCCAACA CATATATTCT TCAACAATAT ATTTAATTTT GAAAAACCTG AAAAAAAAAA CCTGTTAGCA AGTATAAAGG GGCAGTATTA CTATTATTGC ATGAAGGCTT CAAGGGAAAC GTTACAGTCT TTGGGTCATA GTCTGGCTTC AGCTTCCTCT GAGAGTTTAC AGAGGCCAAT TTTGAGCAAA TTCATGGCTA AGGTTATGAG TGAGTTCTGC TAAACAGAAG GCTCACCACA AGGTATCTGG CAGGATTATA CTGGGTAGCT GGATGTTGCA GAAATGTGGT TAGAGGAAGT AACTGTTTTT TTGATGCTCA CAGCATGATG AATCAAATC TGTATCTTAG GATTAGGTTA AAACAATACC TTTGGTATGA TATGAGTGTT GTTGCTGATC CATGCAGCAT GGATTGGAAA GCTGGGGTAT AAGCACACAT GCTAAAGAAA AACATGTAAT TTGGTCCATA CTCACCTGGA TATACTGTTC CTCAGGTTAA AAAATACAGT ACTATCCTAA ATCTTGAAGG CAACTCTCAG CCTATCCATT GAGTTACCTT CAGATCTGCC CTCTGGTTCC TAGCTGTCTT</p>	
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	<p>GGGACTAACT TCTTTCCTGC GCTCAGCTGT TTTCTGGATT CCATGTTTTT CATTTTATTG AGTACTAACT TGTTTTGTCTG CAGCACATCC TTTGGTAGCT TCTAGAGGAA GTTTGTGTGG AGGTAAAATT TTTGAGACCT TGCATGTCTC ATGTTTGATT GATACTTTAT ACGTTTAGGT AGGAGGTAAT TTTCTTCAG GACTTTAAAA ATATTGTTGC TCCATTTTCT TTGTTTCTAT TGTGTATTG AGAAATCAA TGCCATTTTG ATTTCCCAT CATAAATTC ATGATGATGT GTCTTGGTGT GGGTCTATAT TTATCCATTG TATTGGGTTT TAGGTGAACC CTTCAGATA GTAACTCATT TCTGTCAGTT CTGGGAAACA CTTAGCATTG GTTGATGATT TATTCTCTGC TGCTTTGTTC TCCCAACTAT TATTGGATG TTGGATATCC AGCACTGGGT ATCTATTTTC TTACCTCCCT CCCTTGACCC CAGTCTCTGT TTTTLAGCTC TTTAGCTCAA TCTTCCAACCT CTTTGCTATT GTATTTTAAA ATCTTAAGAC CCCTTCTTGA TTTGTAGAAG TTCCTTTTCT TACAACCAA AAGCCTTTAT CTATGGATTT GTTCACAGAT AAGGGGTATT CAATATAGTG TATTTTTTTT TCATTTAAAA TTGTTTGC GC ATCTATTTCC TCCAAATTTT TTTCTGTATT TATTTTTTGT TGTCTATATT TCAGACTTTT CCAGGATATC TGATAATCTT TGGCTGTCTT CTTATGGTTG AAAGAGGGAC TAAAAAGCTT GGAAAGCCTT TGGGTTGTGG GAAGGGGCTG TCTTTAGGAT TATCTGAATG GGCTTTTTTG GGAGTCCCCT CCTCCACATG AATATTTTGG TTTTGTGAGA TTCCCTAGAA TAGAGGCTTC CAATCTCCTT CCTGGAGGGG TCTGTCCAGG AAGGAGATTG TCTAGGGGTC TGTCAGACAG CAGCTTTCAG CTACTTCCTT GATCTTTTTC ACTAATGATT ATATAGTCAT CTAACTACTG TCAACAAGTA ATAGATATCC TATCCTTCAC TTGTTTAGAT TATTGCTGA GATAACCTCT CAAAAGAACC TCTCAAATA AAAGGTTAAC AAGAGCCTAT ATCTTATATT TTTCTTCTCT TTATCTTGTT AGAAGATAGC TATTA AAACC TGTTCTTTTT CTGTCTTGAT AAACACACTT CAATCTTGGT AGAATGGTAG ATGGGACAGT ATATTTTAGG ACCTAAAGCT CTGCAAATGT ATGATCAGCT TGTAAGTACA GGTGCTCAA AACATGAAA CAATCATGCT TTTTACTCTG TAGGAATATC TTTAAAATTC TTGTGAATTT TTCCCAGAA GTAAAGCAA TCTTCCCCCA GAAATAAAAT TAAATGTGCA TAATCTAAAG CTTTTTTTTT TTATTGTGGT AGGATATATA TATAAAACAT AATTIGCCAT TGTAACACATT TAAAATTTAC AAGTCAGAGG CATTAATTAC ATCACAATGT TGTGAAATTA TTA CTACTAT TTCCAAAATT TTCTCATCAC CCCAACTGA AACTCTGTAA CTGTTGAGCA ATAACCTCAT TCCTGTATCT CTCCCAACC CAGGTAACCT CAAATCTTTC TTTTATCTT TGAGACAAGG TCTCATTCTA TCACTCAGGT AGGAGTGCAG TGGTGTGATC ATAGCTCATT GCAGCCTCAA AATCCTGGGC TCAAGCAATC CTCCTTGAGT AGCTAAGACT ATAGGCACAC ATTA ACTGCG CCTGGCTGAT TTTGTTTTTT GTAGAGATGT GGTCTTGCTA TGTTCCTCAT GCTGGTCTTG AGTTCCTGGC CTCAAGCAGT CCTTAAGATT CATCCATGTT GTGGCATGTG TCAGAATTC</p>	
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	<p>ATTTGTTTT ATGACTAAAT AATATTCCAT TGTATGTATA TACATTTTGT TCATCCATCT TCTGATGAAC ACTGGGATAT GTCTACCTTT TGGCTATTGT GAATAATGCT GCAGTAAACA TTGACATAAC AAGTATGTAT TTGATTGCCT GTTTCTAAGT TCTTTTGGGT ATACATCTTG AGTAGAATTG CTAGATAATG TCATGTTTTA TTTCTCTTGT GATTTCTTCT TCGATCCCCT GGTTGAGTGT GTTAATTTCT ACATGTTTAT GAATTTCCCA CTGTTTTTTT GTTATTGATT TCCAAGTTCA TTCCATTGTG ATTAGAGAAG AACTTAGTA TGATTTTAAAT GTTTTTGAGA ATTGGTGTGT GGCTGATAG ATGGTCTGTC CTGGAGAATG TTCCTCATACT ACTTGAGCAA AATATTTATC ATGCTATTGT TGACTGTAGT TTTCTATATG TCTCTTAGGT CAAGGTGGTT TACAATGTGT TAAGGTTCTC TTTTTTAAA AAAATTTTIG CACAGAGTAT CTTTTCTAT GTGTTCCATG TATTTGTGTC TTTGGAGCTA TAGTCTCTTG TAGACAGCAT ATCACTATCT TGTTTTGTTT TGTTTTTCT GTCCATTCTG CCAATTTCTG CCTTTTGATT GGAAAATTTA ATCCATTTGC ATTTAAAGTA ATTAAGGAAG GACTTTCTTC TACCATTTAA CACTTCTTCT ATATGTCATA TACTTTTTTG GCCCTCATT TCCTCTTTAT GGCCTTCTTT TCTGTTTTTT TGTAGTGAAC TAGTCTGATT CTCTTCCAC TCCCCTTGT GTATATTTGT TAGATGTTTT ATTTGIGGTT GCTATGGGGA TTATAGTTAA CATCCTACAC TTAAAACAAT CTAATTTAAA CTGATACCAA TTTACCTTCA ATAGCATACA AAATCTCTAC TCCTGTAAAG CTCTGCCCT GCCCCCTTA TGTTATTGAT GGCACAAATT GCCTAATAAA TAATTTATAG TTATTTGTAT GAGTTTGTCT TTTAAATCAT TTAGGAAATA AAAAGTGGAG TTAGAAAACA GTATGATAGT AATACTGACT TTTATATTTG TCAATATATT TATCTTATTT TGGATCCTTA TTICATTATA TAGATTTGAG TTAGTGTCTA GTGCCCTTCC ATTTCCGGCC AAAGGATTCC CTTATGCATT TCTTGCAGGG CAAGTCTAAT TGTAATAAAC TCCCTCAGCT TTTGTTTTAT CTGAGAATGT CTTGATTTCT CCCTTATTTT TGATGGATAA TTTTGCCAGA TACATGAATT TTTGGTAAAC GTATTTTTCT TTCAGCACTT TAAATATGTC ATCCCACTAC CTTCTGACTT CATGGTTTCT CATGAGATAT TAGATGTTAT AAAATTTGAG GATTCCTCAT TCTTGATGAG TCAGTCTGT CTTATTGCTT TTCGGATTIG CTCAGCTTTT GTCTTTTGAC AGTTTIGATTA TAACGCGGCT CAGTGTGGGT CTCTGAGTTT ATCCCACTTA GAGTTTGTG AGTTTCTTGG AGTCATAGAT TTATGTCTTT TATCAAATTT TGGACATATT TGGCTATTAT TTCTTCAATT TTTTCACTG CTCTTTTCTT TTCCTTCTGA AATATTCTTA ATGTATATGT TGGTCTGTTT GATGCTGTCT CACCAGTTTC TTAGGCTGTG TTCTTTTTIG TTCCTCAGAC TTGATTATTG CAGTTGCCCT TCTTTTTATT TTTTCAAGT TTGTTGATTC TTCTCCCTGT TCAGATCAAC TGTTGAACTC CTCTAGTGAA TTTATTTTCAG TTAGTGTACT TTTAGCTCC AAGATTTATC TTTGGTTCCT TTTTATAACG TCTGTGTCTT TATTGATATT CTCATTTTGT TCATATGTCT CTCTTCTCT</p>	
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	<p>TTAGTTCCTTT GTCCATGTTT TCCTTTAGCT CTTTGGGCTT ATTTAAGACA ATTGTTTAAA GTCTTTGCAT AGTAAGTCCA ATGTCTGTGT TTCTTCAGGG ATGGTTTTCA TTATTTTGT TTCAATGAGC CATACTTTCC TGTGTCTTTG TATGCTGTCT TTTTGTGTGTT GAAAACGTGA TGTTTGAACA TCATAACGTG GTGGCCCTGA AAATCAGATA TTCCCCCTT CCTGAGAGTT AGTTTTATTT TTATTATTGA AGATTGTAGC AGTCTATTGC TACATGTGCA GTCATTTCCA AACTATTTTT GCAAAGACTG TATTCCTTCT GTGTGTCATC ACTGAAGTCT CTGTTCCCTA GTTTGTGTTT AATAGTTTGA CATAGATTTT CTTGAAAGGA GTTAAACTA GCAGAAAAAT CTCTCTCCCA GTCTTTCCAG TCTTTGTAGA TTGGTTCTGT GCTGGGCTTT TCCATTAATA CTTAGCCAGG CTTGTACTGA GCCTAACAAT CAGGCCCAAA AGCGTAGGGT CTTTGCAGAT CTTGTCTGAG CATGCTTCTT GCTGTGTATG CACGTAGTTT TCTAAATCTC CCTGTATGTG CTGTTGAATA TTCTAATTTT CCAAAGAAAC TCCTTTGCAG CTTTTTCTCA CAGAACATAG ATGGTTTTTT GGATATCTTG ACCATAGTCT TTCGACCCAG GTGTTTGCAGG TTGTTAGTTC ACCTTACACT TTTTCAAGC ATTGCCTACT GCTTACGATG AGTGCTCTGT CAATCCTTTA AGTAGCCCCA GACAGGCTAC CAGAGACTTA AACAAAGAATT TGTAAGTTCT GCTCAGCTTC CTCTAGAAAT GGGGATCAGG GTCCAAGACA GAATGCAGTT GCTGATTTCA AGACTGCTGC AACACCAGGG AGCTTGTGGG GGAAGGGCAA GCAGAAATGT CACAAAGCTT TCTTGCCATT TTAAAGTTGC CTGTTCTTGA CTCAGCATTG GCTTCATTGC TATAAACTTT TACTGTTTT TCAGAGTTCT GATAAAATTG GCTATGCCTG TTCCTGCTTT AAAAAATATA TATATATTT TTAGGGATTG GGGTCTCACT AACTGACCA GGCTGGTCTT GAACTTCTGG CCTCAAGCCA TCCTCTCATT TCAGCTTCCC AAAGTGCTGC AATTACACGC GTGAACCACC ACACCCAGCC CCTGCTTGTT TTTCAATGTG CCTACTCCAC CATGTTGCTC AAGTATGTAT ATTTTCTAAA CTACCTTGTA GTGTTGTGAT GGGAAATAAA TCCCTGAGCC TTTTGAATAA CTCAGAGAGA TCAAAAACCTT AGTTTATCCT ATTCGAAGGA TTAGAAAAAT GATATATCTT TCACTTTTTT AGGGATAGGC TCCTCATTAG AAGGCTCCTA TGTGCCGATG CTGTACAAGA CATTTCAATT CTCTTAATGT TTACAACAAG CTTGTTGCCA AGGCTGATCT TGAACTCCTG GCCTCAAACG ATCCTCCCAG CTCAGTCTCA CAAAGTGTTG GGATGTCTGG CCAACTAATG ACTATCTTAA CTCTTGTGTT TCAATGTTA TGCCTTCTTT TATCTTGACT GATTGTATGA CTATGTCTTC TAGAACAATG TTGAACAGAA ATGGTGAGAG CAGACATCCT TGCTTTAATA TTTCACCATT ATATATGATG TTAGGTATAG ATTTTCTCA CAGATGCCTT TTATCAGATT GAGGAATTTA TATTCCTACT TTGCCGAAAG GTTTTTGTAG TATGAGGGGG TGCTGAATTT TGCAAAACAC TTTTTCGGTA ATAATTGAGA TGATTGGTTC TGCAGTCATC GAGATGTGGA TTTTCTCCTT TATTCTGTTT GTGAGTGATT AACTGGTIG ACTAATGTIA AAACAACCTT ACTTTCCAGG</p>	
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		AATAAACCCCT ATTATCTTTT TTATACA	
<i>PSMC4</i>	NM_1 53001. 2	<p>TGCGGGTACG GACAGCGCAT GAGCTTATGT TGAGGGCGGA GCCCAGACCA GCCCTTCGTC CTATCCTGCC CTTCAGCAC CTCTCAGCCG TAACHTAAAC TACACTTCCC AGAAGCCTCC TCAGCCAGGG ACTTCCGTTG TCGTCAGCGG AAGCGGTGAC AGATCATCCC AGGCCACACA GAGGCCGGCT TGGTCACTAT GGAGGAGATA GGCATCTTGG TGGAGAAGGC TCAGGATGAG ATCCCAGCAC TGTCCGTGTC CCGGCCCCAG ACCGGCCTGT CCTTCCTGGG CCCTGAGCCT GAGGACCTGG AGGACCTGTA CAGCCGCTAC AAGGAGGAGG TGAAGCGAAT CCAAAGCATC CCGCTGGTCA TCGGACAATT TCTGGAGGCT GTGGATCAGA ATACAGCCAT CGTGGGCTCT ACCACAGGCT CCAACTATTA TGTGCGCATC CTGAGCACCA TCGATCGGGA GCTGCTCAAG CCAACGCCT CAGTGGCCCT CCACAAGCAC AGCAATGCAC TGGTGGACGT GCTGCCCCCC GAAGCCGACA GCAGCATCAT GATGCTCACC TCAGACCAGA AGCCAGATGT GATGTACGG GACATCGGAG GCATGGACAT CCAGAAGCAG GAGGTGCGGG AGGCCGTGGA GCTCCCGCTC ACGCATTTCG AGCTTACAA GCAGATCGGC ATCGATCCCC CCGAGGGCGT CCTCATGTAT GGCCACCTG GCTGTGGGAA GACCATGTTG GCAAAGGCGG TGGCACATCA CACAACAGCT GCATTCATCC GGGTCGTGGG CTCGGAGTTT GTACAGAAGT ATCTGGGTGA GGGCCCCCGC ATGGTCCGGG ATGTGTTCCG CCTGGCCAAG GAGAATGCAC CTGCCATCAT CTTCATAGAC GAGATTGATG CCATCGCCAC CAAGAGATTC GATGCTCAGA CAGGGGCCGA CAGGGAGGTT CAGAGGATCC TGCTGGAGCT GCTGAATCAG ATGGATGGAT TTGATCAGAA TGTCAATGTC AAGGTAATCA TGGCCACAAA CAGAGCAGAC ACCCTGGATC CGGCCCTGCT ACGGCCAGGA CGGCTGGACC GTAAAATTGA ATTTCCACTT CCTGACCGCC GCCAGAAGAG ATTGATTTTC TCCACTATCA CTAGCAAGAT GAACCTCTCT GAGGAGGTTG ACTTGGAAGA CTATGTGGCC CGGCCAGATA AGATTTCAGG AGCTGATATT AACTCCATCT GTCAGGAGAG TGGAAATGTTG GCTGTCCGTG AAAACCGCTA CATTGTCTTG GCCAAGGACT TCGAGAAAAGC ATACAAGACT GTCATCAAGA AGGACGAGCA GGAGCATGAG TTTTACAAGT GACCCTTCCC TTCCCTCCAC CACACCACTC AGGGGCTGGG GCTTCTCTCG CACCCCAGC ACCTCTGTCC CAAAACCTCA TTCCCTTTTT TCTTTACCCA GGATTGGTTT CTTCATAAA TAGATAAGAT CGAATCCATT TAATTTCTTC TTAGAAGTTT AACTCCTTTG GAGAATGTGG GCCTTGAATA GGATCCTCTG GGTCCTCTT AATCTGACAG ATGAGCAGAC GAGGTGCATG GCCTGGGTTG CAGCTTGAGA GAACCAAAAT ATTCAAACCA GATGACTTCC AAAATGTGGG GAAAGGGATG GAAAATGAAC CTGAGATGGA GTCCTTAATC ACGGGATAAA GCCCTGTGCA TCTCCCTCAT TTCCTACAGG TAAAAGACAG TAAAGAAATT CAGGTCACAG GCCTTGGGAG TTCATAGGAA GGAGATGTCC AGTGCTGTCC AGTAGAACTT T</p>	41

<p>SF3A1</p>	<p>NM_005877.5</p>	<p>GGTCCCGGAA GTGCGCCAGT CGTACCTTCG CGGCCGCAAC TCGCTCGGCC GCCGCCATCT TGCGAGCTCG TCGTACTGAC CGAGCGGGGA GGCTGTCTTG AGGCGGCACC GCTCACCGAC ACCGAGGCGG ACTGGCAGCC CTGAGCGTCG CAGTCATGCC GGCCGGACCC GTGCAGGCGG TGCCCCCGCC GCCGCCCGTG CCCACGGAGC CCAAACAGCC CACAGAAGAA GAAGCATCTT CAAAGGAGGA TTCTGCACCT TCTAAGCCAG TTGTGGGGAT TATTTACCCT CCTCCAGAGG TCAGAAATAT TGTTGACAAG ACTGCCAGCT TTGTGGCCAG AACCGGGCCT GAATTTGAAG CTAGGATCCG ACAGAACGAG ATCAACAACC CCAAGTTCAA CTTCTGAAC CCAATGACC CTTACCATGC CTACTIONGC CACAAGGTCA GCGAGTTCAA GGAAGGGAAG GCTCAGGAGC CGTCCGCCGC CATCCCCAAG GTCATGCAGC AGCAGCAGCA GACCACCCAG CAGCAGCTGC CCCAGAAGGT CCAAGCCCAA GTAATCCAAG AGACCATCGT GCCCAAAGAG CCTCCTCCTG AGTTTGAGTT CATTGCTGAT CCTCCCTCTA TCTCAGCCTT CGACTTGGAT GTGGTGAAGC TGACGGCTCA GTTTGTGGCC AGGAATGGGC GCCAGTTTCT GACCCAGCTG ATGCAGAAAG AGCAGCGCAA CTACCAGTTT GACTTTCTCC GCCCACAGCA CAGCCTCTTC AACTACTTCA CGAAGCTAGT GGAACAGTAC ACCAAGATCT TGATTCCACC CAAAGGTTTA TTTTCAAAGC TCAAGAAAGA GGCTGAAAAC CCCCAGAGAAG TTTTGGATCA GGTGTGTTAC CGAGTGGAAT GGGCCAAATT CCAGGAACGT GAGAGGAAGA AGGAAGAAGA GGAGAAGGAG AAGGAGCGGG TGGCCTATGC TCAGATCGAC TGGCATGATT TTGTGGTGGT GGAAACAGTG GACTTCCAAC CCAATGAGCA AGGGAAC TTC CCTCCCCCA CCACGCCAGA GGAGCTGGGG GCCCGAATCC TCATTCAGGA GCGCTATGAA AAGTTTGGGG AGAGTGAGGA AGTTGAGATG GAGGTGAGT CTGATGAGGA GGATGACAAA CAGGAGAAGG CGGAGGAGCC TCCTTCCCAG CTGGACCAGG ACACCCAAGT ACAAGATATG GATGAGGGTT CAGATGATGA AGAAGAAGG CAGAAAGTGC CCCCACCCC AGAGACACCC ATGCCTCCAC CTCTGCCCC AACTCCAGAC CAAGTCATTG TCCGCAAGGA TTATGATCCC AAAGCCTCCA AGCCCTTGCC TCCAGCCCCT GCTCCAGATG AGTATCTTGT GTCCCCATT ACTGGGGAGA AGATCCCCGC CAGCAAAATG CAGGAACACA TGCGCATTGG ACTTCTTGAC CCTCGCTGGC TGGAGCAGCG GGATCGCTCC ATCCGTGAGA AGCAGAGCGA TGATGAGGTG TACGCACCAG GTCTGGATAT TGAGAGCAGC TTGAAGCAGT TGGCTGAGCG GCGTACTGAC ATCTTCGGTG TAGAGGAAAC AGCCATTGGT AAGAAGATCG GTGAGGAGGA GATCCAGAAG CCAGAGGAAA AGGTGACCTG GGATGGCCAC TCAGGCAGCA TGGCCCGGAC CCAGCAGGCT GCCCAGGCCA ACATCACCTT CCAGGAGCAG ATTGAGGCCA TTCACAAGGC CAAAGGCCTG GTGCCAGAGG ATGACACTAA AGAGAAGATT GGCCCCAGCA AGCCCAATGA AATCCCTCAA CAGCCACCGC CACCATCTTC AGCCACCAAC ATCCCCAGCT CGGCTCCACC CATCACTTCA GTGCCCGGAC CACCCACAAT GCCACCTCCA GTTCGTA</p>	<p>42</p>
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	<p> CAGTTGTCTC CGCAGTACCC GTCATGCCCC GGCCCCAAT GGCATCTGTG GTCCGGCTGC CCCAGGCTC AGTGATCGCC CCCATGCCGC CCATCATCCA CGCGCCCAGA ATCAACGTGG TGCCCATGCC TCCCTCGGCC CCTCCTATTA TGGCCCCCG CCCACCCCC ATGATTGTGC CAACAGCCTT TGTGCCTGCT CCACCTGTGG CACCTGTCCC AGCTCCAGCC CCAATGCCCC CTGTGCATCC CCCACCTCCC ATGGAAGATG AGCCCACCTC CAAAAAAGT AAGACAGAGG ACAGCCTCAT GCCAGAGGAG GAGTTCCTGC GCAGAAACAA GGGTCCAGTG TCCATCAAAG TCCAGGTGCC CAACATGCAG GATAAGACGG AATGGAAACT GAATGGGCAG GTGCTGGTCT TCACCCTCCC ACTCACGGAC CAGGTCTCTG TCATTAAGGT GAAGATTCAT GAAGCCACAG GCATGCCTGC AGGGAACAG AAGCTACAGT ATGAGGGTAT CTTTCATCAA GATTCCAAC CACTGGCTTA CTACAACATG GCCAATGGCG CAGTCATCCA CCTGGCCCTC AAGGAGAGAG GCGGGAGGAA GAAGTAGACA AGAGGAACCT GCTGTCAAGT CCCTGCCATT TTGCCTCTCC TGTCTCCCAC CCCCTGCCCC AGACCCAGGA GCCCCCCTGA GGCTTTGCCT TGCCTGCATA TTTGTTTCGC TCTTACTCAG TTTGGGAATT CAAATTGTCC TGCAGAGGTT CATTCCCCTG ACCCTTTCCC CACATTGGTA AGAGTAGCTG GGTTTTCTAA GCCACTCTCT GGAATCTCTT TGTGTTAGGG TCTCGATTG AGGACATTCA TTTCTTCAGC AGCCCATTAG CAACTGAGAG CCCAGGGATG TCCTACAGGA TAGTTTCATA GTGACAGGTG GCACTTGGCT AATAGAATAT GGCTGATATT GTCATTAATC ATTTTGTACC TTGACATGGG TTGTCTAATA AAACCTCGGAC CCTTCTTGTG AAATCAGTTA AATAAGACTT GTCTCGGTCA CCTGTGCCCT GTCCAGACTC GAGGCAGTGG TAACACTGCA CAGTGCTATG TGGCTTCTCT TTGAGGATTT TTGGGTTTTG TAACTAAATT CTTGCTGCCC TCATACTTTT TATGTATTAG AATCATATTC GTATTGCCCT TTTAAAACAT TGGGATCCTC CAAAGGCCTG CCCCATGTAT TTAACAGTAA TACAGGAAGC ATGGCAGGCA CCATGCAAAC CAAGGATGGA TGGTGCAGTC CCTGTGTCAG TGGGCGGTGG TTTCTGCTG GCCTGGAATC ACTCATCACC TGATTGATTG GCTCTGTGGT CCTGGGCAGG TGCCCTCATAG GTGTGTGGAT ATGATGACGT TTCTTTAAAA TGTATGTATT TAACAAATAC TTAATTGTAT TAAGGTCATG TACCAAGGAT TTGATAAAGT TTAATAAATT TACTCTCTAC TTTTATCCAT TTTATCCATT TTAACTCATG TAATCCTCAT GTGAGTATTC CTGTTTAAACA CTTGAGTAAA CTGAGGCACA GAGAACATAA GTTGCATGCC ATAGTCACAC ACTGTGAAAG TGAAAAGAGA ATGTGTGCAA AACACGTCAC AGTCCTGGTT TCTGAGTAAA GGCAGGCTGT TATCTTTAGA ATCAAGCTAT CACAGGGAGA TAGGCAATGC TGTGGGTGTT GGAGGAAGGT GAGAGCCTGT TGCTAACAAT TTCCTGGTTT TAAAGCTAAG GCTGATTTTA TTGGGAAGAT CTCACATGTG TGTGGCCCT GAGAGTTCCC AGTGCCTTTT ATTTGCAGTC CTTCCATTTG GACCTCCTAG CTGCCCCATC AGGTCATCTC CAGGGCTCAG AGGGGTGAGA CCATTTCCCA </p>	
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		<p>AGGTACACAGA ACCAGCTCTC TAGTCACCAC CCTGCCTCTC CCTCTCACCC AGAGTCAGTA CCAGTTTTAT GGCTTTATTA CAAACTGCTG GGTCCCTCCC ATTTTCAACT TGATTGATGG GATGTCATCC CTTATCCTGT CTGACATTTG CCTCTGGCCT GGTTGCTAGA AGTTTGCCCC AGGGGCAAGA GTTGAAATTT GGCTTCCTGA GGTGGGCTTT GTGGTTTGC TCCCTAAAGT GAGCCCACTA CTGGTTGCTT GTCCATGGCC AACACCAGAA ATCCCCTGAG CACTACCTGG GTCTCATTCC AAGAAGGAAG AGGGTCAGGA GACCTGGGGA GTCTCATATT CCAAGTTCTT CTTTCTTTCT GGGAGCAGTG GGCAGTTCAT GGTGTTAGGG CACTCACCCC CACAGACTGG CAAACCCTGC AGGACTTCCG TGGCTGAGGC TGTGACCGGA GGCCAGGAAT GCCGTTGGGT GGATTGTGAG TGAATGGGCC CTTTGAGCTG CCTCTAGAG AGCAAATCCA GTTTCCTGGA GCTCCTGAAT GAATATCTGT ACTGGCTCGC TCAGATGCAG AAGCTCCATT GACCATGAGG CTTTGTGAAC ATCAGTGGCC ACAGGCCAG TGTGCTGCTT GGCCTGCAC TAGTTTAGGA CCTGCAGCAT GTAGGTAGCG TCCTAGTGT TATAATACAA AGCTGCTCTG CACAGCTTTT CTGATTCTT TTGCAATCTC CTGAGGATTA TCTGCCCAT TTTTAAAACG AGGTGGAATA CCCAAGGTCA TGTAGCCAGT GAGTGCTCTG GAAAGCCAAA GCAGCTCATC CCTTCTGGG GACCACACTG CTCTGCTCCA CCAGACCACA CTATGAAATA GGAATAAGTG CTCCTGTTGC AGGACTGCTG GGAAAACAGG TGGTGTGGGA CTTAAGTCAC CATAATTTTG AAGACTTGCA TGCAGAGGGC TCCAGGAATT GTAGACATTA AGGAATTTCA CTTTCAGTTC TACCCACTAC TTAAGTACTT GTCATGTACT CTTAGAGGAG GCCAGTAATG ATCAGAACCA TTTTACTTTA AAATTAATAA TATTGTATTA GAGAATATAT TAAATGGTTA TATTGGGTTA TGTTAGGATA TATACTTGAA TGGAAATACA TGTACTATTA GCAATCATAT TICATTTATC CCTGTAATTA GACAAGAAAG CATAATATAG CTCTACTCAT GGGTACACAT ACCAGTGTAT AAGATTTTTA GAAGTTTACT TTTTAAAAAT AAAAGCAAAA TGTAAGATCT TAAAAAATAA AAAAAAATAA</p>	
<p><i>PUMI</i></p>	<p>NM_0 010206 58.1</p>	<p>AGTGGGCCGC CATGTTGTCG GAGTGAAAGG TAAGGGGGAG CGAGAGCGCC AGAGAGAGAA GATCGGGGGG CTGAAATCCA TCTTCATCCT ACCGCTCCGC CCGTGTGGT GGAATGAGCG TTGCATGTGT CTGAAGAGA AAAGCAGTGC TTTGGCAGGA CTCTTTCAGC CCCACCTGA AACATCACCC TCAAGAACCA GCTAATCCCA ACATGCCTGT TGTTTTGACA TCTGGAACAG GGTCGCAAGC GCAGCCACAA CCAGCTGCAA ATCAGGCTCT TGCAGCTGGG ACTCACTCCA GCCCTGTCCC AGGATCTATA GGAGTTGCAG GCCGTTCCCA GGACGACGCT ATGGTGGACT ACTTCTTTCA GAGGCAGCAT GGTGAGCAGC TTGGGGGAGG AGGAAGTGGA GGAGGCGGCT ATAATAATAG CAAACATCGA TGGCCTACTG GGGATAACAT TCATGCAGAA CATCAGGTGC GTTCCATGGA TGAAGTGAAT CATGATTTTC AAGCACTTGC TCTGGAGGGA AGAGCGATGG GAGAGCAGCT CTTGCCAGGT AAAAAGTTTT GGGAAACAGA TGAATCCAGC AAAGATGGAC</p>	<p>43</p>

	<p> CAAAAGGAAT ATTCCTGGGT GATCAATGGC GAGACAGTGC CTGGGGAACA TCAGATCATT CAGTTTCCCA GCCAATCATG GTGCAGAGAA GACCTGGTCA GAGTTTCCAT GTGAACAGTG AGGTCAATTC TGTACTGTCC CCACGATCGG AGAGTGGGGG ACTAGGCGTT AGCATGGTGG AGTATGTGTT GAGCTCATCC CCGGGCGATT CCTGTCTAAG AAAAGGAGGA TTTGGCCCAA GGGATGCAGA CAGTGATGAA AACGACAAAG GTGAAAAGAA GAACAAGGGT ACGTTTGATG GAGATAAGCT AGGAGATTTG AAGGAGGAGG GTGATGTGAT GGACAAGACC AATGGTTTAC CAGTGCAGAA TGGGATTGAT GCAGACGTCA AAGATTTTAG CCGTACCCCT GGTAATTGCC AGAACTCTGC TAATGAAGTG GATCTTCTGG GTCCAAACCA GAATGGTTCT GAGGGCTTAG CCCAGCTGAC CAGCACCAAT GGTGCCAAGC CTGTGGAGGA TTTCTCCAAC ATGGAGTCCC AGAGTGTCCC CTTGGACCCC ATGGAACATG TGGGCATGGA GCCTCTTCAG TTTGATTATT CAGGCACGCA GGTACCTGTG GACTCAGCAG CAGCAACTGT GGGACTTTTT GACTACAATT CTCAACAACA GCTGTTCCAA AGACCTAATG CGCTTGCTGT CCAGCAGTTG ACAGCTGCTC AGCAGCAGCA GTATGCACTG GCAGCTGCTC ATCAGCCGCA CATCGGTTTA GCTCCCGCTG CGTTTGTCCC CAATCCATAC ATCATCAGCG CTGCTCCCCC AGGGACGGAC CCCTACACAG CTGGATTGGC TGCAGCAGCG AACTAGGCC CAGCTGTGGT CCCTCACCAG TATTATGGAG TTA CTCCCTG GGGAGTCTAC CCTGCCAGTC TTTTCCAGCA GCAAGCTGCC GCTGCCGCTG CAGCAACTAA TTCAGCTAAT CAACAGACCA CCCACAGGC TCAGCAAGGA CAGCAGCAGG TTCTCCGTGG AGGAGCCAGC CAACGTCTT TGACCCCAA CCAGAACCAG CAGGGACAGC AAACGGATCC CCTTGTGGCA GCTGCAGCAG TGAATTCTGC CCTTGCATTT GGACAAGGTC TGGCAGCAGG CATGCCAGGT TATCCGGTGT TGGCTCCTGC TGCTTACTAT GACCAAATG GTGCCCTTGT AGTGAATGCA GGC GCGAGAA ATGGTCTTGG AGCTCCTGTT CGACTTGTAG CTCCTGCCCC AGTCATCATT AGTTCCTCAG CTGCACAAGC AGCTGTTGCA GCAGCCGCG CTT CAGCAA TGGAGCAGCT GGTGGTCTTG CTGGAACAAC AAATGGACCA TTTGCGCCTT TAGGAACACA GCAGCCTCAG CCCAGCCCC AGCAGCAGCC CAATAACAAC CTGGCATCCA GTTCTTTCTA CGGCAACAAC TCTCTGAACA GCAATTCACA GAGCAGCTCC CTCTTCTCCC AGGGCTCTGC CCAGCCTGCC AACACATCCT TGGGATTCGG AAGTAGCAGT TCTCTCGGCG CCACCCTGGG ATCCGCCCTT GGAGGGTTTG GAACAGCAGT TGCAAATCCT AACACTGGCA GTGGCTCCCG CCGTGA CTCC CTGACTGGCA GCAGTGACCT TTATAAGAGG ACATCGAGCA GCTTGACCCC CATTGGACAC AGTTTTTATA ACGGCCTTAG CTTTTCTCC TCTCCTGGAC CCGTGGGCAT GCCTCTCCCT AGTCAGGGAC CAGGACATTC ACAGACACCA CCTCCTTCCC TCTCTTACA TGGATCCTCT TCAAGCTTAA ACCTGGGAGG ACTCACGAAT GGCAGTGGAA GATACATCTC TGCTGCTCCA GCGCTGAAG CCAAGTACCG CAGTGCAAGC AGCGCCTCCA </p>	
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	<p>GCCTCTTCAG CCCGAGCAGC ACTCTTTTTCT CTTCCTCTCG TTTGCGATAT GGAATGTCTG ATGTCATGCC TTCTGGCAGG AGCAGGCTTT TGGAAGATTT TCGAAACAAC CGGTACCCCA ATTTACAAC TCGGGAGATT GCTGGACATA TAATGGAATT TTCCAAGAC CAGCATGGGT CCAGATTCAT TCAGCTGAAA CTGGAGCGTG CCACACCAGC TGAGCGCCAG CTTGTCTTCA ATGAAATCCT CCAGGCTGCC TACCAACTCA TGGTGGATGT GTTTGGTAAT TACGTCATTC AGAAGTTCTT TGAATTTGGC AGTCTTGAAC AGAAGCTGGC TTTGGCAGAA CGGATTCGAG GCCACGTCCT GTCATTGGCA CTACAGATGT ATGGCTGCCG TGTTATCCAG AAAGCTCTTG AGTTTATTCC TTCAGACCAG CAGGTAATTA ATGAGATGGT TCGGGAAC TA GATGGCCATG TCTTGAAGTG TGTGAAAGAT CAGAATGGCA ATCACGTGGT TCAGAAATGC ATTGAATGTG TACAGCCCCA GTCTTTGCAA TTTATCATCG ATGCGTTTAA GGGACAGGTA TTTGCCTTAT CCACACATCC TTATGGCTGC CGAGTGATTC AGAGAATCCT GGAGCACTGT CTCCCTGACC AGACACTCCC TATTTTAGAG GAGCTTCACC AGCACACAGA GCAGCTTGTA CAGGATCAAT ATGGAAATTA TGTAATCCAA CATGTACTGG AGCACGGTGC TCCTGAGGAT AAAAGCAAAA TTGTAGCAGA AATCCGAGGC AATGTACTTG TATTGAGTCA GCACAAATTT GCAAGCAATG TTGTGGAGAA GTGTGTTACT CACGCCTCAC GTACGGAGCG CGCTGTGCTC ATCGATGAGG TGTGCACCAT GAACGACGGT CCCCACAGTG CCTTATACAC CATGATGAAG GACCAGTATG CCAAC TACGT GGTCCAGAAG ATGATTGACG TGGCGGAGCC AGGCCAGCGG AAGATCGTCA TGCATAAGAT CCGGCCCCAC ATCGCAACTC TTCGTAAGTA CACCTATGGC AAGCACATTC TGGCCAAGCT GGAGAAGTAC TACATGAAGA ACGGTGTTGA CTTAGGGCCC ATCTGTGGCC CCCCTAATGG TATCATCTGA GGCAGTGTC A CCCGCTGTT CCTCATTCCC GCTGACCTCA CTGGCCCACT GGCAAATCCA ACCAGCAACC AGAAATGTTT TAGTGTAGAG TCTGAGACGG GCAAGTGGTT GCTCCAGGAT TACTCCCTCC TCCAAAAAAG GAATCAAATC CACGAGTGGA AAAGCCTTTG TAAATTTAAT TTTATTACAC ATAACATGTA CTATTTTTTT TAATTGACTA ATTGCCCTGC TGTTTTACTG GTGTATAGGA TACTTGTACA TAGGTAACCA ATGTACATGG GAGGCCACAT ATTTTGTTC CTGTTGTATC TATATTTTAC ATGTGGAAAC TTTCAGGGTG GTTGGTTTAA CAAAAAAAAA AAGCTTTAAA AAAAAAAGAA AAAAAGGAAA AGGTTTTTAG CTCATTGCGC TGGCCGGCAA GTTTTGCAA TAGCTCTTCC CCACCTCCTC ATTTTAGTAA AAAACAAACA AAAACAAAAA AACCTGAGAA GTTTGAATTG TAGTTAAATG ACCCCAAACT GGCATTTAAC ACTGTTTATA AAAAATATAT ATATATATAT ATATATATAT AATGAAAAAG GTTTCAGAGT TGCTAAAGCT TCAGTTTGTG ACATTAAGTT TATGAAATTC TAAAAAATGC CTTTTTTGGG GACTATATTA TGCTGAAGAA GGCTGTTTCG GAGGAGGAGA TGCAGCACC CAGAACGTCT TTTGAGGCTG GGCGGGTGTG ATTTGTTTACT GCCTACTGGA TTTTTTCTA</p>	
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		<p>TTAACATTGA AAGGTAAAAT CTGATTATTT AGCATGAGAA AAAAAAATCC AACTCTGCTT TTGGTCTTGC TTCTATAAAT ATATAGTGTA TACTTGGTGT AGACTTTGCA TATATACAAA TTTGTAGTAT TTTCTTGTTT TGATGTCTAA TCTGTATCTA TAATGTACCC TAGTAGTCGA ACATACTTTT GATTGTACAA TTGTACATTT GTATACCTGT AATGTAAATG TGGAGAAGTT TGAATCAACA TAAACACGTT TTTTGGTAAG AAAAGAGAAT TAGCCAGCCC TGTGCATTCA GTGTATATTC TCACCTTTTA TGGTCGTAGC ATATAGTGTT GTATATTGTA AATTGTAATT TCAACCAGAA GTAAATTTTT TTCTTTTGAA GGAATAAATG TTCTTTATAC AGCCTAGTTA ATGTTTAAAA AGAAAAAAT AGCTTGGTTT TATTTGTCAT CTAGTCTCAA GTATAGCGAG ATTCTTTCTA AATGTTATTC AAGATTGAGT TCTCACTAGT GTTTTTTTAA TCCTAAAAAA GTAATGTTTT GATTTTGTGA CAGTCAAAAG GACGTGCAA AGTCTAGCCT TGCCCGAGCT TTCCTTACAA TCAGAGCCCC TCTCACCTTG TAAAGTGTGA ATCGCCCTTC CCTTTTGTAC AGAAGATGAA CTGTATTTTG CATTTTGTCT ACTTGTAAGT GAATGTAACA TACTGTCAAT TTTCTTGTGTT TGAATATAGA ATTGTAACAC TACACGGTGT ACATTTCCAG AGCCTTGTGT ATATTTCCAA TGAACTTTTT TGCAAGCACA CTTGTAACCA TATGTGTATA ATTAACAAAC CTGTGTATGC TTATGCCTGG GCAACTATTT TTTGTAACTC TTGTGTAGAT TGTCTCTAAA CAATGTGTGA TCTTTATTTT GAAAAATACA GAACTTTTGA ATCTGAAAAA AAAAAAAAAA AAAAAAAAAA AAAAAA</p>	
<p><i>ACTB</i></p>	<p>NM_0 01101. 4</p>	<p>GAGTGAGCGG CGCGGGGCCA ATCAGCGTGC GCCGTTCCGA AAGTTGCCTT TTATGGCTCG AGCGGCCGCG GCGGCGCCCT ATAAAACCCA GCGGCGCGAC GCGCCACCAC CGCCGAGACC GCGTCCGCCC CGCGAGCACA GAGCCTCGCC TTTGCCGATC CGCCGCCCGT CCACACCCGC CGCCAGCTCA CCATGGATGA TGATATCGCC GCGCTCGTGC TCGACAACGG CTCCGGCATG TGCAAGGCCG GCTTCGCGGG CGACGATGCC CCCCGGGCCG TCTTCCCCTC CATCGTGGGG CGCCCCAGGC ACCAGGGCGT GATGGTGGGC ATGGGTCAGA AGGATTCCTA TGTGGGCGAC GAGGCCCAGA GCAAGAGAGG CATCCTCACC CTGAAGTACC CCATCGAGCA CGGCATCGTC ACCAACTGGG ACGACATGGA GAAAATCTGG CACCACACCT TCTACAATGA GCTGCGTGTG GCTCCCGAGG AGCACCCCGT GCTGCTGACC GAGGCCCCCC TGAACCCCAA GGCCAACCGC GAGAAGATGA CCCAGATCAT GTTTGAGACC TTCAACACCC CAGCCATGTA CGTTGCTATC CAGGCTGTGC TATCCCTGTA CGCCTCTGGC CGTACCCTG GCATCGTGAT GGACTCCGGT GACGGGGTCA CCCACACTGT GCCCATCTAC GAGGGGTATG CCTTCCCCA TGCCATCCTG CGTCTGGACC TGGCTGGCCG GGACCTGACT GACTACCTCA TGAAGATCCT CACCGAGCGC GGCTACAGCT TCACCACCAC GGCCGAGCGG GAAATCGTGC GTGACATTAA GGAGAAGCTG</p>	<p>44</p>

		<p>TGCTACGTCG CCCTGGACTT CGAGCAAGAG ATGGCCACGG CTGCTTCCAG CTCCTCCCTG GAGAAGAGCT ACGAGCTGCC TGACGGCCAG GTCATCACCA TTGGCAATGA GCGGTTCCGC TGCCCTGAGG CACTCTTCCA GCCTTCCTTC CTGGGCATGG AGTCCTGTGG CATCCACGAA ACTACCTTCA ACTCCATCAT GAAGTGTGAC GTGGACATCC GCAAAGACCT GTACGCCAAC ACAGTGCTGT CTGGCGGCAC CACCATGTAC CCTGGCATTG CCGACAGGAT GCAGAAGGAG ATCACTGCCC TGGCACCCAG CACAATGAAG ATCAAGATCA TTGCTCCTCC TGAGCGCAAG TACTCCGTGT GGATCGGCGG CTCCATCCTG GCCTCGCTGT CCACCTTCCA GCAGATGTGG ATCAGCAAGC AGGAGTATGA CGAGTCCGGC CCCTCCATCG TCCACCGCAA ATGCTTCTAG GCGGACTATG ACTTAGTTGC GTTACACCCT TTCTTGACAA AACCTAACTT GCGCAGAAAA CAAGATGAGA TTGGCATGGC TTTATTIGTT TTTTTTGTTT TGTTTTGGTT TTTTTTTTTT TTTTGGCTTG ACTCAGGATT TAAAAACTGG AACGGTGAAG GTGACAGCAG TCGGTTGGAG CGAGCATCCC CCAAAGTTCA CAATGTGGCC GAGGACTTTG ATTGCACATT GTTGTTTTTT TAATAGTCAT TCCAAATATG AGATGCGTTG TTACAGGAAG TCCCTTGCCA TCCTAAAAGC CACCCCCTT CTCTCTAAGG AGAATGGCCC AGTCCTCTCC CAAGTCCACA CAGGGGAGGT GATAGCATTG CTTTCGTGTA AATTATGTAA TGCAAAATTT TTTTAATCTT CGCCTTAATA CTTTTTTATT TTGTTTTATT TTGAATGATG AGCCTTCGTG CCCCCCTTC CCCCTTTTTT GTCCCCAAC TTGAGATGTA TGAAGGCTTT TGGTCTCCCT GGGAGTGGGT GGAGGCAGCC AGGGCTTACC TGTACACTGA CTTGAGACCA GTTGAATAAA AGTGACACACC TAAAAAATGA GGAAAAAAAA AAAAAAAAAA</p>	
<p><i>GAPD</i></p>	<p>NM_0 02046. 6</p>	<p>GCTCTCTGCT CCTCCTGTTT GACAGTCAGC CGCATCTTCT TTTGCGTCGC CAGCCGAGCC ACATCGCTCA GACACCATGG GGAAGGTGAA GGTCCGAGTC AACGGATTTG GTCGTATTGG GCGCCTGGTC ACCAGGGCTG CTTTAACTC TGGTAAAGTG GATATTGTTG CCATCAATGA CCCCTTCATT GACCTCAACT ACATGGTTTA CATGTTCCAA TATGATTCCA CCCATGGCAA ATTCCATGGC ACCGTCAAGG CTGAGAACGG GAAGCTTGTC ATCAATGGAA ATCCCATCAC CATCTTCCAG GAGCGAGATC CCTCCAAAAT CAAGTGGGGC GATGCTGGCG CTGAGTACGT CGTGGAGTCC ACTGGCGTCT TCACCACCAT GGAGAAGGCT GGGGCTCATT TGCAGGGGGG AGCCAAAAGG GTCATCATCT CTGCCCCCTC TGCTGATGCC CCCATGTTCC TCATGGGTGT GAACCATGAG AAGTATGACA ACAGCCTCAA GATCATCAGC AATGCCTCCT GCACCACCAA CTGCTTAGCA CCCCTGGCCA AGGTCATCCA TGACAACCTT GGTATCGTGG AAGGACTCAT GACCACAGTC CATGCCATCA CTGCCACCCA GAAGACTGTG GATGGCCCCCT CCGGGAAACT GTGGCGTGAT GGCCGCGGGG CTCTCCAGAA CATCATCCCT GCCTCTACTG GCGCTGCCAA</p>	<p>45</p>

		<p>GGCTGTGGGC AAGGTCATCC CTGAGCTGAA CGGGAAGCTC ACTGGCATGG CCTTCCGTGT CCCCACTGCC AACGTGTCAG TGGTGGACCT GACCTGCCGT CTAGAAAAAC CTGCCAAATA TGATGACATC AAGAAGGTGG TGAAGCAGGC GTCGGAGGGC CCCCTCAAGG GCATCCTGGG CTACACTGAG CACCAGGTGG TCTCCTCTGA CTTCAACAGC GACACCCACT CCTCCACCTT TGACGCTGGG GCTGGCATTG CCCTCAACGA CCACTTTGTC AAGCTCATTT CCTGGTATGA CAACGAATTT GGCTACAGCA ACAGGGTGGT GGACCTCATG GCCACATGG CCTCCAAGGA GTAAGACCCC TGGACCACCA GCCCCAGCAA GAGCACAAGA GGAAGAGAGA GACCCTCACT GCTGGGGAGT CCCTGCCACA CTCAGTCCC CACCACACTG AATCTCCCCT CTCACAGTT GCCATGTAGA CCCCTTGAAG AGGGGAGGGG CCTAGGGAGC CGCACCTTGT CATGTACCAT CAATAAAGTA CCCTGTGCTC AACCAGTTAA AAAAAAAAAA AAAAAAAAAA</p>	
<p><i>GUSB</i></p>	<p>NM_00181.3</p>	<p>GTCCTCAACC AAGATGGCGC GGATGGCTTC AGGCGCATCA CGACACCGGC GCGTCACGCG ACCCGCCCTA CGGGCACCTC CCGCGCTTTT CTFAGCGCCG CAGACGGTGG CCGAGCGGGG GACCGGGAAG CATGGCCCGG GGGTCGGCGG TTGCCTGGGC GGCGCTCGGG CCGTTGTGTG GGGGCTGCGC GCTGGGGCTG CAGGGCGGGA TGCTGTACCC CCAGGAGAGC CCGTCGCGGG AGTGCAAGGA GCTGGACGGC CTCTGGAGCT TCCGCGCCGA CTTCTCTGAC AACCGACGCC GGGGCTTCGA GGAGCAGTGG TACCGGCGGC CGCTGTGGGA GTCAGGCCCC ACCGTGGACA TGCCAGTTCC CTCCAGCTTC AATGACATCA GCCAGGACTG GCGTCTGCGG CATTTTGTGCG GCTGGGTGTG GTACGAACGG GAGGTGATCC TGCCGGAGCG ATGGACCCAG GACCTGCGCA CAAGAGTGGT GCTGAGGATT GGCAGTGCCC ATTCCTATGC CATCGTGTGG GTGAATGGGG TCGACACGCT AGAGCATGAG GGGGGCTACC TCCCCTTCGA GGCCGACATC AGCAACCTGG TCCAGGTGGG GCCCCTGCC TCCCGGCTCC GAATCACTAT CGCCATCAAC AACACACTCA CCCCACCAC CCTGCCACCA GGGACCATCC AATACCTGAC TGACACCTCC AAGTATCCCA AGGGTTACTT TGTCCAGAAC ACATATTTTG ACTTTTTCAA CTACGCTGGA CTGCAGCGGT CTGTACTTCT GTACACGACA CCCACCACCT ACATCGATGA CATCACCGTC ACCACCAGCG TGGAGCAAGA CAGTGGGCTG GTGAATTACC AGATCTCTGT CAAGGGCAGT AACCTGTTC AAGTTGGAAGT GCGTCTTTTG GATGCAGAAA ACAAAGTCGT GCGGAATGGG ACTGGGACCC AGGGCCAAC TAAGGTGCCA GGTGTCAGCC TCTGGTGGCC GTACCTGATG CACGAACGCC CTGCCTATCT GTATTCATTG GAGGTGCAGC TGA CTGCACA GACGTC ACTG GGGCCTGTGT CTGACTTCTA CACTCCCT GTGGGGATCC GCACTGTGGC TGTCACCAAG AGCCAGTTCC TCATCAATGG GAAACCTTTC TATTTCCACG GTGTCAACAA GCATGAGGAT GCGGACATCC GAGGGAAGGG CTTCGACTGG CCGCTGCTGG TGAAGGACTT CAACCTGCTT CGCTGGCTTG GTGCCAACGC TTTCCGTACC AGCCACTACC CCTATGCAGA GGAAGTGATG CAGATGTGTG</p>	<p>46</p>

		<p>ACCGCTATGG GATTGTGGTC ATCGATGAGT GTCCCGGCGT GGGCCTGGCG CTGCCGCAGT TCTTCAACAA CGTTTCTCTG CATCACCACA TGCAGGTGAT GGAAGAAGTG GTGCGTAGGG ACAAGAACCA CCCCgcggTC GTGATGTGGT CTGTGGCCAA CGAGCCTGCG TCCCACCTAG AATCTGCTGG CTACTACTTG AAGATGGTGA TCGCTCACAC CAAATCCTTG GACCCCTCCC GGCCTGTGAC CTTTGTGAGC AACTCTAACT ATGCAGCAGA CAAGGGGGCT CCGTATGTGG ATGTGATCTG TTTGAACAGC TACTACTCTT GGTATCACGA CTACGGGCAC CTGGAGTTGA TTCAGCTGCA GCTGGCCACC CAGTTTGAGA ACTGGTATAA GAAGTATCAG AAGCCATTA TTCAGAGCGA GTATGGAGCA GAAACGATTG CAGGGTTTCA CCAGGATCCA CCTCTGATGT TCACTGAAGA GTACCAGAAA AGTCTGCTAG AGCAGTACCA TCTGGGTCTG GATCAAAAAC GCAGAAAATA CGTGGTTGGA GAGCTCATT GGAATTTTGC CGATTTTCATG ACTGAACAGT CACCGACGAG AGTGCTGGGG AATAAAAAGG GGATCTTCAC TCGGCAGAGA CAACCAAAAA GTGCAGCGTT CCTTTTGCGA GAGAGATACT GGAAGATTGC CAATGAAACC AGGTATCCCC ACTCAGTAGC CAAGTCACAA TGTTTGAAA ACAGCCTGTT TACTTGAGCA AGACTGATAC CACCTGCGTG TCCCTTCCTC CCCGAGTCAG GGCGACTTCC ACAGCAGCAG AACAAGTGCC TCCTGGACTG TTCACGGCAG ACCAGAACGT TTCTGGCCTG GGTTTTGTGG TCATCTATTC TAGCAGGGAA CACTAAAGGT GGAAATAAAA GATTTTCTAT TATGGAAATA AAGAGTTGGC ATGAAAGTGG CTACTGAAAA AAAAAAAAAA AAAAAAAAAA A</p>	
<p>RPLP0</p>	<p>NM_0 01002. 3</p>	<p>GTCTGACGGG CGATGGCGCA GCCAATAGAC AGGAGCGCTA TCCGCGGTTT CTGATTGGCT ACTTTGTTTCG CATTATAAAA GGCACGCGCG GGCgcgAGGC CCTTCTCTCG CCAGGCGTCC TCGTGGAAGT GACATCGTCT TTAACCCCTG CGTGGCAATC CCTGACGCAC CGCCGTGATG CCCAGGGAAG ACAGGGCGAC CTGGAAGTCC AACTACTTCC TTAAGATCAT CCAACTATTG GATGATTATC CGAAATGTTT CATTGTGGGA GCAGACAATG TGGGCTCCAA GCAGATGCAG CAGATCCGCA TGTCCCTTCG CGGGAAGGCT GTGGTGCTGA TGGGCAAGAA CACCATGATG CGCAAGGCCA TCCGAGGGCA CCTGGAAAAC AACCCAGCTC TGGAGAAACT GCTGCCTCAT ATCCGGGGGA ATGTGGGCTT TGTGTTACC AAGGAGGACC TCACTGAGAT CAGGGACATG TTGCTGGCCA ATAAGGTGCC AGCTGCTGCC CGTGCTGGTG CCATTGCCCC ATGTGAAGTC ACTGTGCCAG CCCAGAACAC TGGTCTCGGG CCCGAGAAGA CCTCCTTTTT CCAGGCTTTA GGTATCACCA CTAAAATCTC CAGGGGCACC ATTGAAATCC TGAGTGATGT GCAGCTGATC AAGACTGGAG ACAAAGTGGG AGCCAGCGAA GCCACGCTGC TGAACATGCT CAACATCTCC CCCTTCTCCT TTGGGCTGGT CATCCAGCAG GTGTTTCACA ATGGCAGCAT CTACAACCCT GAAGTGCTTG ATATCACAGA GGAAACTCTG CATTCTCGCT TCCTGGAGGG TGTCGCAAT</p>	<p>47</p>

		<p>GTTGCCAGTG TCTGTCTGCA GATTGGCTAC CCAACTGTTG CATCAGTACC CCATICTATC ATCAACGGGT ACAAACGAGT CCTGGCCTTG TCTGTGGAGA CGGATTACAC CTTCCCACTT GCTGAAAAGG TCAAGGCCTT CTTGGCTGAT CCATCTGCCT TTGTGGCTGC TGCCCCTGTG GCTGCTGCCA CCACAGCTGC TCCTGCTGCT GCTGCAGCCC CAGCTAAGGT TGAAGCCAAG GAAGAGTCGG AGGAGTCGGA CGAGGATATG GGATTTGGTC TCTTTGACTA ATCACC AAAA AGCAACCAAC TTAGCCAGTT TTATTIGCAA AACAAGGAAA TAAAGGCTTA CTTCCTTAAA AAGTAAAAAA AAAAAAAAAA AAAAAAAAAA</p>	
<p><i>TFRC</i></p>	<p>NM_0 03234. 3</p>	<p>AGAGCGTCGG GATATCGGGT GCGGGCTCGG GACGGAGGAC GCGCTAGTGT GAGTGCGGGC TTCTAGA ACT ACACCGACCC TCGTGTCTC CCTTCATCCT GCGGGGCTGG CTGGAGCGGC CGCTCCGGTG CTGTCCAGCA GCCATAGGGA GCCGCACGGG GAGCGGGAAA GCGGTCGCGG CCCCAGGCGG GGCGGCCGGG ATGGAGCGGG GCCGCGAGCC TGTGGGGAAG GGGCTGTGGC GCGCCTCGA GCGGCTGCAG GTTCTTCTGT GTGGCAGTTC AGAATGATGG ATCAAGCTAG ATCAGCATT CTAACCTTGT TTGGTGGAGA ACCATTGTCA TATACCGGT TCAGCCTGGC TCGGCAAGTA GATGGCGATA ACAGTCATGT GGAGATGAAA CTTGCTGTAG ATGAAGAAGA AAATGCTGAC AATAACACAA AGGCCAATGT CACAAAACCA AAAAGGTGTA GTGGAAGTAT CTGCTATGGG ACTATTGCTG TGATCGTCTT TTTCTTGATT GGATTTATGA TTGGCTACTT GGGCTATTGT AAAGGGGTAG AACCAAAAAC TGAGTGTGAG AGACTGGCAG GAACCGAGTC TCCAGTGAGG GAGGAGCCAG GAGAGGACTT CCCTGCAGCA CGTCGCTTAT ATTGGGATGA CCTGAAGAGA AAGTTGTCCG AGAAACTGGA CAGCACAGAC TTCACCGGCA CCATCAAGCT GCTGAATGAA AATTCATATG TCCCTCGTGA GGCTGGATCT CAAAAGATG AAAATCTTGC GTTGTATGTT GAAAATCAAT TTCGTGAATT TAAACTCAGC AAAGTCTGGC GTGATCAACA TTTTGTTAAG ATTCAGGTCA AAGACAGCGC TCAAACTCG GTGATCATAG TTGATAAGAA CGGTAGACTT GTTTACCTGG TGGAGAAATCC TGGGGGTTAT GTGGCGTATA GTAAGGCTGC AACAGTTACT GGTA AACTGG TCCATGCTAA TTTTGGTACT AAAAAAGATT TTGAGGATTT ATACACTCCT GTGAATGGAT CTATAGTGAT TGTCAGAGCA GGGAAAATCA CCTTTGCAGA AAAGGTTGCA AATGCTGAAA GCTTAAATGC AATTGGTGTG TTGATATACA TGGACCAGAC TAAATTTCCC ATTGTAAACG CAGAACTTTC ATTCTTGGGA CATGCTCATC TGGGGACAGG TGACCCTTAC ACACCTGGAT TCCCTTCCTT CAATCACACT CAGTTTCCAC CATCTCGGTC ATCAGGATTG CCTAATATAC CTGTCCAGAC AATCTCCAGA GCTGCTGCAG AAAAGCTGTT TGGAATATG GAAGGAGACT GTCCCTCTGA CTGGAAAACA GACTCTACAT GTAGGATGGT AACCTCAGAA AGCAAGAATG TGAAGCTCAC TGTGAGCAAT GTGCTGAAAG AGATAAAAT</p>	<p>48</p>

	<p>TCTTAACATC TTGGAGTTA TTAAAGGCTT TGTAGAACCA GATCACTATG TTGTAGTTGG GGCCAGAGA GATGCATGGG GCCCTGGAGC TGCAAAATCC GGTGTAGGCA CAGCTCTCCT ATTGAAACTT GCCCAGATGT TCTCAGATAT GGTCTTAAA GATGGGTTT AGCCAGCAG AAGCATTATC TTTGCCAGTT GGAGTGCTGG AGACTTTGGA TCGGTTGGTG CCACTGAATG GCTAGAGGGA TACCTTTCGT CCCTGCATTT AAAGGCTTTC ACTTATATTA ATCTGGATAA AGCGGTTCTT GTTACCAGCA ACTTCAAGGT TTCTGCCAGC CCACTGTTGT ATACGCTTAT TGAGAAAACA ATGCAAAATG TGAAGCATCC GGTTACTGGG CAATTTCTAT ATCAGGACAG CCACTGGGCC AGCAAAGTTG AGAAACTCAC TTTAGACAAT GCTGCTTTC CTTTCCTTGC ATATTCTGGA ATCCCAGCAG TTTCTTTCCTG TTTTTCGCGAG GACACAGATT ATCCTTATTT GGGTACCACC ATGGACACCT ATAAGGAACT GATTGAGAGG ATCCTGAGT TGAACAAAGT GGCACGAGCA GCTGCAGAGG TCGCTGGTCA GTTCGTGATT AACTAACC ATGATGTTGA ATTGAACCTG GACTATGAGA GGTACAACAG CCAACTGCTT TCATTTGTGA GGGATCTGAA CCAATACAGA GCAGACATAA AGGAAATGGG CCTGAGTTTA CAGTGGCTGT ATTCTGCTCG TGGAGACTTC TTCCGIGCTA CTTCCAGACT AACAACAGAT TTCGGGAATG CTGAGAAAAC AGACAGATTT GTCATGAAGA AACTCAATGA TCGTGTGATG AGAGTGGAGT ATCACTTCCT CTCTCCCTAC GTATCTCAA AAGAGTCTCC TTTCCGACAT GTCTTCTGGG GCTCCGGCTC TCACACGCTG CCAGCTTAC TGGAGAACTT GAAACTGCGT AAACAAAATA ACGGTGCTTT TAATGAAACG CTGTTTCAGAA ACCAGTTGGC TCTAGCTACT TGGACTATTC AGGGAGCTGC AAATGCCCTC TCTGGTGACG TTTGGGACAT TGACAATGAG TTTTAAATGT GATACCCATA GCTTCCATGA GAACAGCAGG GTAGTCTGGT TTCTAGACTT GTGCTGATCG TGCTAAATTT TCAGTAGGGC TACAAAACCT GATGTTAAA TTCCATCCCA TCATCTTGGT ACTACTAGAT GTCTTTAGGC AGCAGCTTTT AATACAGGGT AGATAACCTG TACTTCAAGT TAAAGTGAAT AACCCTTAA AAAATGTCCA TGATGGAATA TTCCCCTATC TCTAGAATTT TAAGTGCTTT GTAATGGGAA CTGCCTCTTT CCTGTTGTTG TTAATGAAA TGTCAGAAAC CAGTTATGTG AATGATCTCT CTGAATCCTA AGGGCTGGTC TCTGCTGAAG GTTGTAAGTG GTCGCTTACT TTGAGTGATC CTCCAACCTC ATTTGATGCT AAATAGGAGA TACCAGGTTG AAAGACCTTC TCCAATGAG ATCTAAGCCT TTCCATAAGG AATGTAGCTG GTTTCCTCAT TCCTGAAAGA AACAGTTAAC TTTCAGAAGA GATGGGCTTG TTTTCTTGCC AATGAGGTCT GAAATGGAGG TCCTTCTGCT GGATAAAATG AGGTTCAACT GTTGATTGCA GGAATAAGGC CTTAATATGT TAACCTCAGT GTCATTTATG AAAAGAGGGG ACCAGAAGCC AAAGACTTAG TATATTTTCT TTTCTCTGT CCTTCCCC ATAAGCCTCC ATTTAGTTCT TTGTTATTTT TGTTCCTCC AAAGCACATT GAAAGAGAAC CAGTTTCAGG TGTTTAGTTG CAGACTCAGT TTGTCAGACT</p>	
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	<p>TTAAAGAATA ATATGCTGCC AAATTTTGGC CAAAGTGTTA ATCTTAGGGG AGAGCTTCTGTCCTTTTGG CACTGAGATA TTTATTGTTT ATTTATCAGT GACAGAGTTC ACTATAAATG GTGTTTTTTT AATAGAATAT AATTATCGGA AGCAGTGCCT TCCATAATTA TGACAGTTAT ACTGTCCGGT TTTTTTAAAT AAAAGCAGCA TCTGCTAATA AAACCCAACA GATACTGGAA GTTTTGCATT TATGGTCAAC ACTTAAGGGT TTTAGAAAAC AGCCGTCAGC CAAATGTAAT TGAATAAAGT TGAAGCTAAG ATTTAGAGAT GAATTAATTA TAATTAGGGG TTGCTAAGAA GCGAGCACTG ACCAGATAAG AATGCTGGT TTCCTAAATG CAGTGAATTG TGACCAAGTT ATAAATCAAT GTCACCTAAA GGCTGTGGTA GTACTCCTGC AAAATTTTAT AGCTCAGTTT ATCCAAGGTG TAACTCTAAT TCCCATTTTG CAAAATTTCC AGTACCTTTG TCACAATCCT AACACATTAT CGGGAGCAGT GTCTTCCATA ATGTATAAAG AACAAAGGTAG TTTTACCTA CCACAGTGTG TGTATCGGAG ACAGTGATCT CCATATGTTA CACTAAGGGT GTAAGTAATT ATCGGGAACA GTGTTTCCCA TAATTTTCTT CATGCAATGA CATCTTCAA GCTTGAAGAT CGTTAGTATC TAACATGTAT CCCAACTCCT ATAATTCCT ATCTTTTAGT TTTAGTTGCA GAAACATTTT GTGGTCATTA AGCATTGGGT GGGTAAATTC AACCACTGTA AAATGAAATT ACTACAAAAT TTGAAATTTA GCTTGGGTTT TTGTTACCTT TATGGTTTCT CCAGGTCCTC TACTTAATGA GATAGTAGCA TACATTTATA ATGTTTGCTA TTGACAAGTC ATTTAACTT TATCACATTA TTTGCATGTT ACCTCCTATA AACTTAGTGC GGACAAGTTT TAATCCAGAA TTGACCTTTT GACTTAAAGC AGAGGGACTT TGTATAGAAG GTTTGGGGGC TGTGGGGAAG GAGAGTCCCC TGAAGGCTG ACACGCTGC CTACCCATT GTGGTGATCA ATTAAATGTA GGTATGAATA AGTTCGAAGC TCCGTGAGTG AACCATCATT ATAAACGIGA TGATCAGCTG TTTGTCATAG GGCAGTTGGA AACGGCCTCC TAGGGAAAAG TTCATAGGGT CTCTTCAGGT TCTTAGTGTC ACTTACCTAG ATTTACAGCC TCACTTGAAT GTGTCACTAC TCACAGTCTC TTTAATCTTC AGTTTTATCT TTAATCTCCT CTTTTATCTT GGACTGACAT TTAGCGTAGC TAAGTGAAAA GGTCATAGCT GAGATTCTG GTTCGGGTGT TACGCACACG TACTTAAATG AAAGCATGTG GCATGTTTAT CGTATAACAC AATATGAATA CAGGGCATGC ATTTTGCAGC AGTGAGTCTC TTCAGAAAAC CCTTTTCTAC AGTTAGGGT GAGTTACTTC CTATCAAGCC AGTACGTGCT AACAGGCTCA ATATTCCTGA ATGAAATATC AGACTAGTGA CAAGCTCCTG GTCTTGAGAT GTCTTCTCGT TAAGGAGATG GGCCTTTTGG AGGTAAAGGA TAAAATGAAT GAGTCTGTG ATGATTCCT ATTCTAGAAC TTGCATGACC TTTACTGTGT TAGCTCTTTG AATGTTCTTG AAATTTTAGA CTTTCTTTGT AAACAAATGA TATGTCCTTA TCATTGTATA AAAGCTGTTA TGTGCAACAG TGTGGAGATT CCTTGTCTGA TTTAATAAAA TACTTAAACA CTGAAAAAAA AAAA</p>	
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18S	X0320 5.1	<p>TACCTGGTTG ATCCTGCCAG TAGCATATGC TTGTCTCAA GATTAAGCCA TGCATGTCTA AGTACGCACG GCCGGTACAG TGAAACTGCG AATGGCTCAT TAAATCAGTT ATGGTTCCTT TGGTCGCTCG CTCCTCTCCC ACTTGGATAA CTGTGGTAAT TCTAGAGCTA ATACATGCCG ACGGGCGCTG ACCCCCTTCG CGGGGGGAT GCGTGCATTT ATCAGATCAA AACCAACCCG GTCAGCCCCT CTCCGGCCCC GGCCGGGGGG CGGGCGCCGG CGGCTTTGGT GACTCTAGAT AACCTCGGGC CGATCGCACG CCCCCGTGG CGGCGACGAC CCATTCGAAC GTCTGCCCTA TCAACTTTTCG ATGGTAGTCG CCGTGCCTAC CATGGTGACC ACGGGTGACG GGAATCAGG GTTCGATTCC GGAGAGGGAG CCTGAGAAAC GGCTACCACA TCCAAGGAAG GCAGCAGGCG CGCAAATTAC CCACTCCCGA CCCGGGGAGG TAGTGACGAA AAATAACAAT ACAGGACTCT TTCGAGGCC TGTAATTGGA ATGAGTCCAC TTTAAATCCT TTAACGAGGA TCCATTGGAG GGCAAGTCTG GTGCCAGCAG CCGCGGTAAT TCCAGCTCCA ATAGCGTATA TTAAGTTGC TGCAGTTAAA AAGCTCGTAG TTGGATCTTG GGAGCGGGCG GGCGGTCCGC CGCGAGGCGA GCCACCGCCC GTCCCCGCC CTTGCCTCTC GGCGCCCCCT CGATGCTCTT AGCTGAGTGT CCCGCGGGGC CCGAAGCGTT TACTTTGAAA AAATTAGAGT GTTCAAAGCA GGCCCGAGCC GCCTGGATAC CGCAGCTAGG AATAATGGAA TAGGACCGCG GTTCTATTTT GTTGGTTTTC GGAAGTGGAG CCATGATTAA GAGGGACGGC CGGGGGCATT CGTATTGCGC CGCTAGAGGT GAAATTCTTG GACCGGCGCA AGACGGACCA GAGCGAAAGC ATTTGCCAAG AATGTTTTCA TTAATCAAGA ACGAAAGTCG GAGGTTTCGAA GACGATCAGA TACCGTCGTA GTTCCGACCA TAAACGATGC CGACCGGCGA TCGGGCGGCG TTATTCCCAT GACCCGCCGG GCAGCTTCCG GGAAACCAA GTCTTTGGGT TCCGGGGGGA GTATGGTTGC AAAGCTGAAA CTAAAGGAA TTGACGGAAG GGCACCACCA GGAGTGGAGC CTGCGGCTTA ATTTGACTCA ACACGGGAAA CCTACCCGG CCCGGACACG GACAGGATTG ACAGATTGAT AGCTCTTTCT CGATTCCGTG GGTGGTGGTG CATGGCCGTT CTTAGTTGGT GGAGCGATTT GTCTGGTTAA TTCCGATAAC GAACGAGACT CTGGCATGCT AACTAGTTAC GCGACCCCG AGCGGTCGGC GTCCCCAAC TTCTTAGAGG GACAAGTGGC GTTCAGCCAC CCGAGATTGA GCAATAACAG GTCTGTGATG CCCTTAGATG TCCGGGGCTG CACGCGCGCT ACACTGACTG GCTCAGCGTG TGCCTACCCT ACGCCGGCAG GCGCGGGTAA CCCGTTGAAC CCCATTCGTG ATGGGGATCG GGGATTGCAA TTATTCCCA TGAACGAGGA ATTCCAGTA AGTGCGGGTC ATAAGCTTGC GTTGATTAAG TCCCTGCCCT TTGTACACAC CGCCCGTCGC TACTACCGAT TGGATGGTTT AGTGAGGCC TCGGATCGGC CCCGCCGGG TCGGCCACG GCCCTGGCGG AGCGCTGAGA AGACGGTCGA ACTTGACTAT CTAGAGGAAG TAAAAGTCGT AACAAGGTTT CCGTAGGTGA ACCTGCGGAA GGATCATTA</p>	49
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<p>PPIA</p>	<p>NM_0 21130. 4</p>	<p>GGGGCCGAAC GTGGTATAAA AGGGGCGGGA GGCCAGGCTC GTGCCGTTTT GCAGACGCCA CCGCCGAGGA AAACCGTGTA CTATTAGCCA TGGTCAACCC CACCGTGTTT TTCGACATTG CCGTCGACGG CGAGCCCTTG GGCCGCGTCT CCTTTGAGCT GTTTGCAGAC AAGGTCCCAA AGACAGCAGA AAATTTTCGT GCTCTGAGCA CTGGAGAGAA AGGATTTGGT TATAAGGGT CCTGCTTTCA CAGAATTATT CCAGGGTTTA TGTGTCAGGG TGGTGACTTC ACACGCCATA ATGGCACTGG TGGCAAGTCC ATCTATGGGG AGAAATTTGA AGATGAGAAC TTCATCCTAA AGCATAACGGG TCCTGGCATC TTGTCCATGG CAAATGCTGG ACCCAACACA AATGGTTCCC AGTTTTTCAT CTGCACTGCC AAGACTGAGT GGTGGATGG CAAGCATGTG GTGTTTGGCA AAGTGAAAGA AGGCATGAAT ATTGTGGAGG CCATGGAGCG CTTTGGGTCC AGGAATGGCA AGACCAGCAA GAAGATCACC ATTGCTGACT GTGGACAACG CGAATAAGTT TGACTTGTGT TTTATCTTAA CCACCAGATC ATTCCTTCTG TAGCTCAGGA GAGCACCCCT CCACCCCATG TGCTCGCAGT ATCCTAGAAT CTTTGTGCTC TCGCTGCAGT TCCCTTTGGG TTCCATGTTT TCCTTGTTCC CTCCCATGCC TAGCTGGATT GCAGAGTTAA GTTTATGATT ATGAAATAAA AACTAAATAA CAATTGTCCT CGTTTGAAGT AAGAGTGTG ATGTAGGCTT TATTTAAGC AGTAATGGGT TACTTCTGAA ACATCACTTG TTTGCTTAAT TCTACACAGT ACTTAGATTT TTTTACTTT CCAGTCCCAG GAAGTGCAA TGTTTGTGTA GTGGAATATT GAAAATGTAG GCAGCAACTG GGCATGGTGG CTCACTGTCT GTAATGTATT ACCTGAGGCA GAAGACCACC TGAGGGTAGG AGTCAAGATC AGCCTGGGCA ACATAGTGAG ACGCTGTCTC TACAAAAAAT AATTAGCCTG GCCTGGTGGT GCATGCCTAG TCCTAGCTGA TCTGGAGGCT GACGTGGGAG GATTGCTTGA GCCTAGAGTG AGCTATTATC ATGCCACTGT ACAGCCTGGG TGTTACACAGA TCTTGTGTCT CAAAGGTAGG CAGAGGCAGG AAAAGCAAGG AGCCAGAATT AAGAGGTTGG GTCAGTCTGC AGTGAGTTCA TGCATTTAGA GGTGTTCTTC AAGATGACTA ATGTCAAAAA TTGAGACATC TGTTGCGGTT TTTTTTTTTT TTTTTTCCC TGGAATGCAG TGGCGTGATC TCAGCTCACT GCAGCCTCCG CCTCCTGGGT TCAAGTGATT CTAGTGCCTC AGCCTCCTGA GTAGCTGGGA TAATGGGCGT GTGCCACCAT GCCCAGCTAA TTTTGTATT TTTAGTATAG ATGGGGTTTC ATCATTTTGA CCAGGCTGGT CTCAAACCTCT TGACCTCAGC TGATGCGCCT GCCTTGGCCT CCCAAACTGC TGAGATTACA GATGTGAGCC ACCGCACCCCT ACCTCATTTT CTGTAACAAA GCTAAGCTTG AACACTGTTG ATGTTCTTGA GGAAGCATA TTGGGCTTTA GGCTGTAGGT CAAGTTTATA CATCTTAATT ATGGTGGAAT TCCTATGTAG AGTCTAAAAA GCCAGGTAAT TGGTGCTACA GTCAGTCTCC CTGCAGAGGG TTAAGGCGCA GACTACCTGC AGTGAGGAGG TACTGCTTGT AGCATATAGA GCCTCTCCCT AGCTTTGGTT ATGGAGGCTT TGAGGTTTTG CAAACCTGAC CAATTTAAGC CATAAGATCT GGTCAAAGGG ATACCCTTCC</p>	<p>50</p>
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		<p>CACTAAGGAC TTGGTTTCTC AGGAAATTAT ATGTACAGTG CTTGCTGGCA GTTAGATGTC AGGACAATCT AAGCTGAGAA AACCCCTTCT CTGCCACCT TAACAGACCT CTAGGGTTCT TAACCCAGCA ATCAAGTTTG CCTATCCTAG AGGTGGCGGA TTTGATCATT TGGTGTGTTG GGCAATTTTT GTTTTACTGT CTGGTTCCTT CTGCGTGAAT TACCACCACC ACCACTTGTG CATCTCAGTC TTGTGTGTTG TCTGGTTACG TATCCCTGG GTGATACCAT TCAATGTCTT AATGTACTTG TGGCTCAGAC CTGAGTGCAA GGTGGAAATA AACATCAAAC ATCTTTTCAT TATCCCTA</p>	
<p><i>PGK1</i></p>	<p>NM_00291.3</p>	<p>GAGAGCAGCG GCCGGGAAGG GCGGGTGC GG GAGGCGGGGT GTGGGGCGGT AGTGTGGGCC CTGTTCTG CCGCGCGGTG TTCCGCATTC TGCAAGCCTC CGGAGCGCAC GTCGGCAGTC GGCTCCCTCG TTGACCGAAT CACCGACCTC TCTCCCAGC TGTATTTCCA AAATGTCGCT TTCTAACAAG CTGACGCTGG ACAAGCTGGA CGTTAAAGGG AAGCGGGTCG TTATGAGAGT CGACTTCAAT GTTCTATGA AGAACAACCA GATAACAAAC AACCAGAGGA TTAAGGCTGC TGTCCCAAGC ATCAAATTCT GCTTGGACAA TGGAGCCAAG TCGGTAGTCC TTATGAGCCA CCTAGGCCGG CCTGATGGTG TGCCATGCC TGACAAGTAC TCCTTAGAGC CAGTTGCTGT AGAACTCAA TCTCTGCTGG GCAAGGATGT TCTGTTCTTG AAGGACTGTG TAGGCCAGA AGTGGAGAAA GCCTGTGCCA ACCCAGCTGC TGGGTCTGTC ATCTGCTGG AGAACCTCCG CTTTCATGTG GAGGAAGAAG GGAAGGAAA AGATGCTTCT GGGAACAAGG TTAAAGCCGA GCCAGCCAAA ATAGAAGCTT TCCGAGCTTC ACTTTCCAAG CTAGGGGATG TCTATGTCAA TGATGCTTTT GGCCTGCTC ACAGAGCCCA CAGCTCCATG GTAGGAGTCA ATCTGCCACA GAAGGCTGGT GGGTTTTTGA TGAAGAAGGA GCTGAACTAC TTTGCAAAGG CCTTGGAGAG CCCAGAGCGA CCCTTCCTGG CCATCCTGGG CGGAGCTAAA GTTGCAGACA AGATCCAGCT CATCAATAAT ATGCTGGACA AAGTCAATGA GATGATTATT GGTGGTGGAA TGGCTTTTAC CTTCTTAAG GTGCTCAACA ACATGGAGAT TGGCACTTCT CTGTTTGATG AAGAGGGAGC CAAGATTGTC AAAGACCTAA TGTCCAAAGC TGAGAAGAAT GGTGTGAAGA TTACCTTGCC TGTTGACTTT GTCCTGCTG ACAAGTTTGA TGAGAATGCC AAGACTGGCC AAGCCACTGT GGCTTCTGGC ATACCTGCTG GCTGGATGGG CTTGGACTGT GGTCTGAAA GCAGCAAGAA GTATGCTGAG GCTGTCCTC GGGCTAAGCA GATTGTGTGG AATGGTCTG TGGGGGTATT TGAATGGGAA GCTTTTGCCC GGGGAACCAA AGCTCTCATG GATGAGGTGG TGAAAGCCAC TTCTAGGGGC TGCATACCA TCATAGGTGG TGGAGACACT GCCACTTGCT GTGCCAAATG GAACACGGAG GATAAAGTCA GCCATGTGAG CACTGGGGGT GGTGCCAGTT TGGAGCTCCT GGAAGGTAAG GTCCTTCTG GGGTGGATGC TCTCAGCAAT ATTTAGTACT TTCCTGCCTT</p>	<p>51</p>

		<p>TTAGTTCCTG TGCACAGCCC CTAAGTCAAC TTAGCATT CTGCATCTCC ACTTGGCATT AGCTAAAACC TTCCATGTCA AGATTCAGCT AGTGGCCAAG AGATGCAGTG CCAGGAACCC TTAAACAGTT GCACAGCATC TCAGCTCATC TTCACTGCAC CCTGGATTG CATACTTCT TCAAGATCCC ATTTGAATTT TTTAGTGACT AAACCATTGT GCATTCTAGA GTGCATATAT TTATATTTTG CCTGTAAAA AGAAAGTGAG CAGTGTTAGC TTAGTCTCT TTTGATGTAG GTTATTATGA TTAGCTTTGT CACTGTTTCA CTA CTACAGCA TGGAAACAAG ATGAAATTCC ATTTGTAGGT AGTGAGACAA AATTGATGAT CCATTAAGTA AACAAATAAAA GTGTCCATTG AAACCGTGAT TTTTTTTTT TTCTGTTCAT ACTTTGTTAG GAAGGGTGAG AATAGAATCT TGAGGAACGG ATCAGATGTC TATATTGCTG AATGCAAGAA GTGGGGCAGC AGCAGTGGAG AGATGGGACA ATTAGATAAA TGTCCATTCT TTATCAAGGG CCTACTTTAT GGCAGACATT GTGCTAGTGC TTTTATTCTA ACTTTTATTT TTATCAGTTA CACATGATCA TAATTTAAAA AGTCAAGGCT TATAACAAAA AAGCCCCAGC CCATTCCTCC CATTCAAGAT TCCCCTCCC CAGAGGTGAC CACTTTCAAC TCTTGAGTTT TTCAGGTATA TACCTCCATG TTCTAAGTA ATATGCTTAT ATTGTTCACT TCTTTTTTTT TTATTTTTTA AAGAAATCTA TTTCATACCA TGGAGGAAGG CTCTGTTCCA CATATATTTT CACTTCTTCA TTCTCTCGGT ATAGTTTIGT CACAATTATA GATTAGATCA AAAGTCTACA TAACTAATAC AGCTGAGCTA TGTAGTATGC TATGATTA AAA TTTACTTATG TAAAAAAA AAAAAAAA</p>	
<p><i>RPL13A</i></p>	<p>NM_012423.3</p>	<p>CACTTCTGCC GCCCTGTTT CAAGGGATAA GAAACCTGC GACAAAACCT CCTCCTTTT CAAGCGGCTG CCGAAGATGG CGGAGGTGCA GGTCTGTTG CTGATGGTC GAGGCCATCT CCTGGGCCGC CTGGCGGCA TCGTGGCTAA ACAGGTACTG CTGGGCCGGA AGGTGGTGGT CGTACGCTGT GAAGGCATCA ACATTTCTGG CAATTTCTAC AGAAACAAGT TGAAGTACCT GGCTTTCCTC CGCAAGCGGA TGAACACCAA CCCTTCCGA GGCCCCTACC ACTTCCGGGC CCCCAGCCGC ATCTTCTGGC GGACCGTGCG AGGTATGCTG CCCCACAAAA CCAAGCGAGG CCAGGCCGCT CTGGACCGTC TCAAGGTGTT TGACGGCATC CCACCGCCCT ACGACAAGAA AAAGCGGATG GTGGTTCTTG CTGCCCTCAA GGTCGTGCGT CTGAAGCCTA CAAGAAAGTT TGCTATCTG GGGCGCCTGG CTCACGAGGT TGGCTGGAAG TACCAGGCAG TGACAGCCAC CCTGGAGGAG AAGAGGAAAG AGAAAGCCAA GATCCACTAC CGGAAGAAGA AACAGCTCAT GAGGCTACGG AAACAGGCCG AGAAGAACGT GGAGAAGAAA ATTGACAAAT ACACAGAGGT CCTCAAGACC CACGGACTCC TGGTCTGAGC CCAATAAAGA CTGTTAATTC CTCATGCGTT GCCTGCCCTT CCTCCATTGT TGCCCTGGAA TGTACGGGAC CCAGGGGCAG CAGCAGTCCA GGTGCCACAG GCAGCCCTGG GACATAGGAA GCTGGGAGCA AGGAAAGGGT CTTAGTCACT</p>	<p>52</p>

		GCCTCCCGAA GTTGCTTGAA AGCACTCGGA GAATTGTGCA GGTGTCAATTT ATCTATGACC AATAGGAAGA GCAACCAGTT ACTATGAGTG AAAGGGAGCC AGAAGACTGA TTGGAGGGCC CTATCTTGTG AGTGGGGCAT CTGTTGGACT TTCCACCTGG TCATATACTC TGCAGCTGTT AGAATGTGCA AGCACTTGGG GACAGCATGA GCTTGCTGTT GTACACAGGG TATTTCTAGA AGCAGAAATA GACTGGGAAG ATGCACAACC AAGGGGTTAC AGGCATCGCC CATGCTCCTC ACCTGTATTT TGTAATCAGA AATAAATTGC TTTTAAAGAA AAAAAAAAAA AAAAAA	
<i>B2M</i>	NM_0 04048. 2	AATATAAGTG GAGGCGTCGC GCTGGCGGGC ATTCCTGAAG CTGACAGCAT TCGGGCCGAG ATGTCTCGCT CCGTGGCCTT AGCTGTGCTC GCGCTACTCT CTCTTTCTGG CCTGGAGGCT ATCCAGCGTA CTCCAAGAT TCAGGTTTAC TCACGTCATC CAGCAGAGAA TGGAAAGTCA AATTTCTGTA ATTGCTATGT GTCTGGGTTT CATCCATCCG ACATTGAAGT TGACTTACTG AAGAATGGAG AGAGAATTGA AAAAGTGGAG CATTACAGACT TGCTTTTCAG CAAGGACTGG TCTTTCTATC TCTTGTACTA CACTGAATTC ACCCCCCTG AAAAAGATGA GTATGCCTGC CGTGTGAACC ATGTGACTTT GTCACAGCCC AAGATAGTTA AGTGGGATCG AGACATGTAA GCAGCATCAT GGAGGTTTGA AGATGCCGCA TTTGGATTGG ATGAATTCCA AATTCTGCTT GCTTGCTTTT TAATATTGAT ATGCTTATAC ACTTACACTT TATGCACAAA ATGTAGGGTT ATAATAATGT TAACATGGAC ATGATCTTCT TTATAATTCT ACTTTGAGTG CTGTCTCCAT GTTTGATGTA TCTGAGCAGG TTGCTCCACA GGTAGCTCTA GGAGGGCTGG CAACTTAGAG GTGGGGAGCA GAGAATTCTC TTATCCAACA TCAACATCTT GGTCAGATTT GAACTCTTCA ATCTCTTGCA CTCAAAGCTT GTTAAGATAG TTAAGCGTGC ATAAGTTAAC TTCCAATTTA CATACTCTGC TTAGAATTTG GGGGAAAATT TAGAAATATA ATTGACAGGA TTATTGGAAA TTTGTTATAA TGAATGAAAC ATTTTGTCAT ATAAGATTCA TATTTACTTC TTATACATTT GATAAAGTAA GGCATGGTTG TGGTTAATCT GGTTTATTTT TGTTCCACAA GTTAAATAAA TCATAAAACT TGATGTGTTA TCTCTTA	53
<i>YWHAZ</i>	NM_0 03406. 3	CTTTCTCCTT CCCCTTCTTC CGGGCTCCCG TCCC GGCTCA TCACCCGGCC TGTGGCCCAC TCCCACCGCC AGCTGGAACC CTGGGGACTA CGACGTCCCT CAAACCTTGC TTCTAGGAGA TAAAAAGAAC ATCCAGTCAT GGATAAAAAT GAGCTGGTTC AGAAGGCCAA ACTGGCCGAG CAGGCTGAGC GATATGATGA CATGGCAGCC TGCATGAAGT CTGTAAGTGA GCAAGGAGCT GAATTATCCA ATGAGGAGAG GAATCTTCTC TCAGTTGCTT ATAAAAATGT TGTAGGAGCC CGTAGGTCAT CTTGGAGGGT CGTCTCAAGT ATTGAACAAA AGACGGAAGG TGCTGAGAAA AAACAGCAGA TGGCTCGAGA ATACAGAGAG AAAATTGAGA CGGAGCTAAG AGATATCTGC AATGATGTAC TGTCTTTTT GGAAAAGTTC TTGATCCCCA ATGCTTCACA AGCAGAGAGC	54

	<p>AAAGTCTTCT ATTTGAAAAT GAAAGGAGAT TACTACCGTT ACTTGGCTGA GGTTGCCGCT GGTGATGACA AGAAAGGGAT TGTCGATCAG TCACAACAAG CATACCAAGA AGCTTTTGAA ATCAGCAAAA AGGAAATGCA ACCAACACAT CCTATCAGAC TGGGTCTGGC CCTTAACTTC TCTGTGTTCT ATTATGAGAT TCTGAACTCC CCAGAGAAAAG CCTGCTCTCT TGCAAAGACA GCTTTTGATG AAGCCATTGC TGAAC TTGAT ACATTAAGTG AAGAGTCATA CAAAGACAGC ACGCTAATAA TGCAATTACT GAGAGACAAC TTGACATTGT GGACATCGGA TACCCAAGGA GACGAAGCTG AAGCAGGAGA AGGAGGGGAA AATTAACCGG CCTTCCAAC TTTGTCTGCC TCATTCTAAA ATTTACACAG TAGACCATT GTCATCCATG CTGTCCACA AATAGTTTTT TGTTTACGAT TTATGACAGG TTTATGTTAC TTCTATTTGA ATTTCTATAT TTCCCATGTG GTTTTTATGT TTAATATTAG GGGAGTAGAG CCAGTTAACA TTTAGGGAGT TATCTGTTTT CATCTTGAGG TGGCCAATAT GGGGATGTGG AATTTTTATA CAAGTTATAA GTGTTTGGCA TAGTACTTTT GTTACATTGT GGCTTCAAAA GGGCCAGTGT AAAACTGCTT CCATGTCTAA GCAAAGAAAA CTGCCTACAT ACTGGTTTGT CCTGGCGGGG AATAAAAGGG ATCATTGGTT CCAGTCACAG GTGTAGTAAT TGTGGGTACT TTAAGGTTTG GAGCACTTAC AAGGCTGTGG TAGAATCATA CCCCATGGAT ACCACATATT AAACCATGTA TATCTGTGGA ATACTCAATG TGTACACCTT TGACTACAGC TGCAGAAGTG TTCCTTTAGA CAAAGTTGTG ACCCATTTTA CTCTGGATAA GGGCAGAAAC GGTTACATT CCATTATTTG TAAAGTTACC TGCTGTTAGC TTTTATTATT TTTGCTACAC TCATTTTATT TGTATTTAAA TGTTTTAGGC AACCTAAGAA CAAATGTAAG ATGAAAGATG CAGGAAAAAT GAATTGCTTG GTATTCATTA CTTCATGTAT ATCAAGCACA GCAGTAA AAC AAAAACCCAT GTATTTAACT TTTTTTTAGG ATTTTTGCTT TTGTGATTTT TTTTTTTTTG ATACTTGCCT AACATGCATG TGCTGTAAAA ATAGTTAACA GGGAAATAAC TTGAGATGAT GGCTAGCTTT GTTTAATGTC TTATGAAATT TTCATGAACA ATCCAAGCAT AATTGTTAAG AACACGTGTA TTA AATTCAT GTAAGTGGAA TAAAAGTTTT ATGAATGGAC TTTTCAACTA CTTTCTCTAC AGCTTTTCAT GTAAATTAGT CTTGGTCTG AAACTTCTCT AAAGGAAATT GTACATTTTT TGAAATTTAT TCCTTATTCC CTCTTGGCAG CTAATGGGCT CTTACCAAGT TTAAACACAA AATTATCAT AACAAAAATA CTACTAATAT AACTACTGTT TCCATGTCCC ATGATCCCCT CTCTTCCCTC CCACCCTGAA AAAAATGAGT TCCTATTTTT TCTGGGAGAG GGGGGGATTG ATTAGAAAAA AATGTAGTGT GTTCCATTTA AAATTTTGGC ATATGGCATT TTCTAACTTA GGAAGCCACA ATGTTCTTGG CCCATCATGA CATTGGGTAG CATTAACTGT AAGTTTTGTG CTTCCAAATC ACTTTTTGGT TTTAAGAAT TTCTTGATAC TCTTATAGCC TGCCTTCAAT TTTGATCCTT TATTCCTTCT ATTTGTCAGG TGCACAAGAT TACCTTCCCTG TTTTAGCCTT CTGTCTTGIC ACCAACCATT CTTACTTGGT</p>	
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		<p>GGCCATGTAC TTGGAAAAAG GCCGCATGAT CTTTCTGGCT CCACTCAGTG TCTAAGGCAC CCTGCTTCCT TTGCTTGCAT CCCACAGACT ATTTCCCTCA TCCTATTTAC TGCAGCAAAT CTCTCCTTAG TTGATGAGAC TGTGTTTATC TCCCTTTAAA ACCCTACCTA TCCTGAATGG TCTGTCATTG TCTGCCTTA AAATCCTTCC TCTTTCTTCC TCCTCTATTC TCTAAATAAT GATGGGGCTA AGTTATACCC AAAGCTCACT TTACAAAATA TTTCTCAGT ACTTTGCAGA AAACACCAA CAAAAATGCC ATTTTAAAAA AGGTGTATTT TTTCTTTTAG AATGTAAGCT CCTCAAGAGC AGGGACAATG TTTTCTGTAT GTTCTATTGT GCCTAGTACA CTGTAATGC TCAATAAATA TTGATGATGG GAGGCAGTGA GTCTTGATGA TAAGGGTGAG AAACGAAAT CCCAAACACT GTTTTGTTGC TTGTTTTATT ATGACCTCAG ATTAATTGG GAAATATTGG CCCTTTTGAA TAATTGTCCC AAATATTACA TTCAAATAAA AGTGCAATGG AGAAAAAAA AAA</p>	
<p><i>SDHA</i></p>	<p>NM_0 04168. 3</p>	<p>ACTGCAGCCC CGCTCGACTC CGGCGTGGTG CGCAGGCGCG GTATCCCCC TCCCCGCCA GCTCGACCCC GGTGTGGTGC GCAGGCGCAG TCTGCGCAGG GACTGGCGGG ACTGCGCGGC GGCAACAGCA GACATGTCGG GGGTCCGGGG CCTGTCGCGG CTGCTGAGCG CTCGGCGCCT GGCCTGGCC AAGGCGTGGC CAACAGTGTT GCAAACAGGA ACCCGAGGTT TTCACTTAC TGTTGATGGG AACAAGAGGG CATCTGCTAA AGTTTCAGAT TCCATTTCTG CTCAGTATCC AGTAGTGGAT CATGAATTTG ATGCAGTGGT GGTAGGCGCT GGAGGGGCAG GCTTGCAGC TGCATTTGGC CTTTCTGAGG CAGGGTTTAA TACAGCATGT GTTACCAAGC TGTTTCCTAC CAGGTCACAC ACTGTTGCAG CACAGGGAGG AATCAATGCT GCTCTGGGA ACATGGAGGA GGACAACCTG AGGTGGCATT TCTACGACAC CGTGAAGGGC TCCGACTGGC TGGGGGACCA GGATGCCATC CACTACATGA CGGAGCAGGC CCCC GCCGCC GTGGTCGAGC TAGAAAATTA TGGCATGCCG TTTAGCAGAA CTGAAGATGG GAAGATTTAT CAGCGTGCAT TTGGTGGACA GAGCCTCAAG TTTGGAAAGG GCGGGCAGGC CCATCGGTGC TGCTGTGTGG CTGATCGGAC TGGCCACTCG CTATTGCACA CCTTATATGG AAGGTCTCTG CGATATGATA CCAGCTATTT TGTGGAGTAT TTIGCCTTGG ATCTCCTGAT GGAGAATGGG GAGTGCCGTG GTGTCATCGC ACTGTGCATA GAGGACGGGT CCATCCATCG CATAAGAGCA AAGAACACTG TTGTTGCCAC AGGAGGCTAC GGGCGCACCT ACTTCAGCTG CACGTCTGCC CACACCAGCA CTGGCGACGG CACGGCCATG ATCACCAGGG CAGGCCTTCC TTGCCAGGAC CTAGAGTTTG TTCAGTTCCA CCTACAGGC ATATATGGTG CTGGTTGTCT CATTACGGAA GGATGTCTGTG GAGAGGGAGG CATTCTCATT AACAGTCAAG GCGAAAGGTT TATGGAGCGA TACGCCCTG TCGGAAGGA CCTGGCGTCT AGAGATGTGG TGCTCGGTG CATGACTCTG GAGATCCGAG AAGGAAGAGG</p>	<p>55</p>

		<p>CTGTGGCCCT GAGAAAGATC ACGTCTACCT GCAGCTGCAC CACCTACCTC CAGAGCAGCT GGCCACGCGC CTGCCTGGCA TTTCAGAGAC AGCCATGATC TTCGCTGGCG TGGACGTCAC GAAGGAGCCG ATCCCTGTCC TCCCACCGT GCATTATAAC ATGGGCGGCA TTCCCACCAA CTACAAGGGG CAGGTCCTGA GGCACGTGAA TGGCCAGGAT CAGATTGTGC CCGGCCTGTA CGCCTGTGGG GAGGCCGCCT GTGCCTCGGT ACATGGTGCC AACCGCCTCG GGGCAAACCTC GCTCTTGGAC CTGGTTGTCT TTGGTCGGGC ATGTGCCCTG AGCATCGAAG AGTCATGCAG GCCTGGAGAT AAAGTCCCTC CAATTAACC AAACGCTGGG GAAGAATCTG TCATGAATCT TGACAAATTG AGATTTGCTG ATGGAAGCAT AAGAACATCG GAACTGCGAC TCAGCATGCA GAAGTCAATG CAAAATCATG CTGCCGTGTT CCGTGTGGGA AGCGTGTTC AAGAAGGTTG TGGGAAAATC AGCAAGCTCT ATGGAGACCT AAAGCACCTG AAGACGTTCG ACCGGGGAAT GGTCTGGAAC ACGGACCTGG TGGAGACCCT GGAGCTGCAG AACCTGATGC TGTGTGCGCT GCAGACCATC TACGGAGCAG AGGCACGGAA GGAGTCACGG GGCGCGCATG CCAGGGAAGA CTACAAGGTG CGGATTGATG AGTACGATTA CTCCAAGCCC ATCCAGGGGC AACAGAAGAA GCCCTTTGAG GAGCACTGGA GGAAGCACAC CCTGTCTAT GTGGACGTTG GCACTGGGAA GGTCACTCTG GAATATAGAC CCGTGATCGA CAAAACCTTG AACGAGGCTG ACTGTGCCAC CGTCCCGCCA GCCATTCGCT CCTACTGATG AGACAAGATG TGGTGATGAC AGAATCAGCT TTTGTAATTA TGTATAATAG CTCATGCATG TGTCATGTC ATAACTGTCT TCATACGCTT CTGCACTCTG GGAAGAAGG AGTACATTGA AGGGAGATTG GCACCTAGTG GCTGGGAGCT TGCCAGGAAC CCAGTGGCCA GGGAGCGTGG CACTTACCTT TGCCCTTGC TTCATTCTTG TGAGATGATA AACTGGGCA CAGCTCTTAA ATAAAATATA AATGAACAAA CTTTCTTTTA TTTCCAAATC CATTGAAAT ATTTTACTGT TGTGACTTTA GTCATATTTG TTGACCTAAA AATCAAATGT AATCTTTGTA TTGTGTTACA TCAAATCCA GATATTTTGT ATAGTTTCTT TTTTCTTTTT CTTTTCTTTT TTTTTTTTGA GACAGGATCG GTGCAGTAGT ACAATCACAG CTCACTGCAG CCTCAAACCTC CTGGGCAGCT CAGGTGATCT TCCTGACTCA GCCTTCTGAG TAGTTGGGGC TACAGGTGTG CACCACCATG CCCAGCTCAT TTATTTTGTA ATTGTAGGGA CAGGGTCTCA CTGTGTGGC TAGGCTGGTC TCAAGTGATC CTCCCTCCTT GGCCTCCCAA GGTGCTGGAA TTATAGGTGT GAACAAACCA AAAAAAAAAA AAA</p>	
<p><i>HPRT1</i></p>	<p>NM_00194.2</p>	<p>GGCGGGGCCT GCTTCTCCTC AGCTTCAGGC GGCTGCGACG AGCCCTCAGG CGAACCTCTC GGCTTTCCCG CGCGGGCGCCG CCTCTTGCTG CGCCTCCGCC TCCTCCTCTG CTCCGCCACC GGCTTCTCC TCCTGAGCAG TCAGCCCGCG CGCCGGCCCG CTCCGTTATG GCGACCCGCA GCCCTGGCGT CGTGATTAGT GATGATGAAC CAGGTTATGA CCTIGATTA TTTGCATAC</p>	<p>56</p>

	<p>CTAATCATTG TGCTGAGGAT TTGGAAAGGG TGTTTATTCC TCATGGACTA ATTATGGACA GGAAGTGAACG TCTTGCTCGA GATGTGATGA AGGAGATGGG AGGCCATCAC ATTGTAGCCC TCTGTGTGCT CAAGGGGGGC TATAAATTCT TTGCTGACCT GCTGGATTAC ATCAAAGCAC TGAATAGAAA TAGTGATAGA TCCATTCCCTA TGACTGTAGA TTTTATCAGA CTGAAGAGCT ATTGTAATGA CCAGTCAACA GGGGACATAA AAGTAATTGG TGGAGATGAT CTCTCAACTT TAACTGGAAA GAATGTCTTG ATTGTGGAAG ATATAATTGA CACTGGCAAA ACAATGCAGA CTTTGCTTTC CTTGGTCAGG CAGTATAATC CAAAGATGGT CAAGGTCGCA AGCTTGCTGG TGAAAAGGAC CCCACGAAGT GTTGGATATA AGCCAGACTT TGTTGGATTT GAAATTCCAG ACAAGTTTGT TGTAGGATAT GCCCTTGACT ATAATGAATA CTTCAGGGAT TTGAATCATG TTTGTGTCAT TAGTGAAACT GGAAAAGCAA AATACAAAGC CTAAGATGAG AGTTCAAGTT GAGTTTGGAA ACATCTGGAG TCCTATTGAC ATCGCCAGTA AAATTATCAA TGTTCTAGTT CTGTGGCCAT CTGCTTAGTA GAGCTTTTTG CATGTATCTT CTAAGAATTT TATCTGTTTT GTACTTTAGA AATGTCAGTT GCTGCATTCC TAACTGTTTT ATTTGCACTA TGAGCCTATA GACTATCAGT TCCCTTTGGG CGGATTGTTG TTTAACTTGT AAATGAAAAA ATTCTCTTAA ACCACAGCAC TATTGAGTGA AACATTGAAC TCATATCTGT AAGAAATAAA GAGAAGATAT ATTAGTTTTT TAATTGGTAT TTTAATTTTT ATATATGCAG GAAAGAATAG AAGTGATTGA ATATTGTTAA TTATACCACC GTGTGTTAGA AAAGTAAGAA GCAGTCAATT TTCACATCAA AGACAGCATC TAAGAAGTTT TGTTCTGTCC TGGAATTATT TTAGTAGTGT TTCAGTAATG TTGACTGTAT TTCCAACTT GTTCAAATTA TTACCAGTGA ATCTTTGTCA GCAGTTCCT TTTAAATGCA AATCAATAAA TTCCCAAAAA TTTAAAAAAA AAAAAAAAAA AAAAA</p>	
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Definitions

[0065] The articles “a” and “an” are used in this disclosure to refer to one or more than one (i.e., to at least one) of the grammatical object of the article. By way of example, “an element” means one element or more than one element.

[0066] The term “and/or” is used in this disclosure to mean either “and” or “or” unless indicated otherwise.

[0067] As used herein, the terms “polynucleotide” and “nucleic acid molecule” are used interchangeably to mean a polymeric form of nucleotides of at least 10 bases or base pairs in length, either ribonucleotides or deoxynucleotides or a modified form of either type of

nucleotide, and is meant to include single and double stranded forms of DNA. As used herein, a nucleic acid molecule or nucleic acid sequence that serves as a probe in a microarray analysis preferably comprises a chain of nucleotides, more preferably DNA and/or RNA. In other embodiments, a nucleic acid molecule or nucleic acid sequence comprises other kinds of nucleic acid structures such as for instance a DNA/RNA helix, peptide nucleic acid (PNA), locked nucleic acid (LNA) and/or a ribozyme. Hence, as used herein the term "nucleic acid molecule" also encompasses a chain comprising non-natural nucleotides, modified nucleotides and/or non-nucleotide building blocks which exhibit the same function as natural nucleotides.

[0068] As used herein, the terms "hybridize," "hybridizing", "hybridizes," and the like, used in the context of polynucleotides, are meant to refer to conventional hybridization conditions, such as hybridization in 50% formamide/6XSSC/0.1% SDS/100 µg/ml ssDNA, in which temperatures for hybridization are above 37 degrees and temperatures for washing in 0.1 XSSC/0.1% SDS are above 55 degrees C, and preferably to stringent hybridization conditions.

[0069] As used herein, the term "normalization" or "normalizer" refers to the expression of a differential value in terms of a standard value to adjust for effects which arise from technical variation due to sample handling, sample preparation, and measurement methods rather than biological variation of biomarker concentration in a sample. For example, when measuring the expression of a differentially expressed protein, the absolute value for the expression of the protein can be expressed in terms of an absolute value for the expression of a standard protein that is substantially constant in expression.

[0070] The terms "diagnosis" and "diagnostics" also encompass the terms "prognosis" and "prognostics", respectively, as well as the applications of such procedures over two or more time points to monitor the diagnosis and/or prognosis over time, and statistical modeling based thereupon. Furthermore, the term diagnosis includes: a. prediction (determining if a patient will likely develop aggressive disease (hyperproliferative/invasive)), b. prognosis (predicting whether a patient will likely have a better or worse outcome at a pre-selected time in the future), c. therapy selection, d. therapeutic drug monitoring, and e. relapse monitoring.

[0071] The term "providing" as used herein with regard to a biological sample refers to directly or indirectly obtaining the biological sample from a subject. For example, "providing"

may refer to the act of directly obtaining the biological sample from a subject (e.g., by a blood draw, tissue biopsy, lavage and the like). Likewise, "providing" may refer to the act of indirectly obtaining the biological sample. For example, providing may refer to the act of a laboratory receiving the sample from the party that directly obtained the sample, or to the act of obtaining the sample from an archive.

[0072] "Accuracy" refers to the degree of conformity of a measured or calculated quantity (a test reported value) to its actual (or true) value. Clinical accuracy relates to the proportion of true outcomes (true positives (TP) or true negatives (TN) versus misclassified outcomes (false positives (FP) or false negatives (FN)), and may be stated as a sensitivity, specificity, positive predictive values (PPV) or negative predictive values (NPV), or as a likelihood, odds ratio, among other measures.

[0073] The term "biological sample" as used herein refers to any sample of biological origin potentially containing one or more biomarkers. Examples of biological samples include tissue, organs, or bodily fluids such as whole blood, plasma, serum, tissue, lavage or any other specimen used for detection of disease.

[0074] The term "subject" as used herein refers to a mammal, preferably a human.

[0075] "Treating" or "treatment" as used herein with regard to a condition may refer to preventing the condition, slowing the onset or rate of development of the condition, reducing the risk of developing the condition, preventing or delaying the development of symptoms associated with the condition, reducing or ending symptoms associated with the condition, generating a complete or partial regression of the condition, or some combination thereof.

[0076] Biomarker levels may change due to treatment of the disease. The changes in biomarker levels may be measured by the present disclosure. Changes in biomarker levels may be used to monitor the progression of disease or therapy.

[0077] "Altered", "changed" or "significantly different" refer to a detectable change or difference from a reasonably comparable state, profile, measurement, or the like. Such changes may be all or none. They may be incremental and need not be linear. They may be by orders of magnitude. A change may be an increase or decrease by 1%, 5%, 10%, 20%, 30%, 40%, 50%,

60%, 70%, 80%, 90%, 95%, 99%, 100%, or more, or any value in between 0% and 100%.

Alternatively, the change may be 1-fold, 1.5- fold, 2-fold, 3-fold, 4-fold, 5-fold or more, or any values in between 1-fold and five-fold. The change may be statistically significant with a p value of 0.1, 0.05, 0.001, or 0.0001.

[0078] The term “stable disease” refers to a diagnosis for the presence of a melanoma, however the melanoma has been treated and remains in a stable condition, i.e. one that that is not progressive, as determined by imaging data and/or best clinical judgment.

[0079] The term “progressive disease” refers to a diagnosis for the presence of a highly active state of a melanoma, i.e. one has not been treated and is not stable or has been treated and has not responded to therapy, or has been treated and active disease remains, as determined by imaging data and/or best clinical judgment.

Examples

[0080] The disclosure is further illustrated by the following examples, which are not to be construed as limiting this disclosure in scope or spirit to the specific procedures herein described. It is to be understood that the examples are provided to illustrate certain embodiments and that no limitation to the scope of the disclosure is intended thereby. It is to be further understood that resort may be had to various other embodiments, modifications, and equivalents thereof which may suggest themselves to those skilled in the art without departing from the spirit of the present disclosure and/or scope of the appended claims.

EXAMPLE 1

[0081] Derivation of a 28-marker gene panel

[0082] Raw probe intensities ($n = 6,892,960$ features) from $n = 49$ whole blood samples were used to identify genes that best discriminated between different types of melanoma samples e.g., treated versus untreated, simultaneously. A total of 28 transcripts were identified in an unbiased manner as potential markers of melanoma behavior (**Table 2**).

[0083] An artificial intelligence model of melanoma disease dynamics was built using normalized gene expression of these 28 markers in whole blood from Controls ($n = 90$), Responders/Stable ($n = 68$), and Progressive ($n = 66$) samples. The dataset was randomly split into training ($n = 169$) and testing ($n = 55$) partitions for model creation and validation respectively. Five algorithms (XGB, RF, TreeBag, SVM, NNET) were identified that best predicted the training data. In the test set, each algorithm produced probability scores that predicted the sample. Each probability score reflects the “certainty” of an algorithm that an unknown sample belongs to Control, Responder/Stable or Progressive class. For example, an unknown sample S1 can have the following probability vector {Control = 20, Responder = 50, Progressive = 30}. This sample would be considered a responder, given a score of 50, but there may be potential of progression (probability of 30) (FIG. 1).

[0084] Table 2.

Melanoma Biomarker or Housekeeping Gene		Chromosome location [Cytogenetic band]	Uni Gene ID	RefSeq	Amplification for forward and reverse primers	Assay Location	Exon Boundary
Symbol	Name						
ATL1	atlastin GTPase 1	Chr.14: 50533082 - 50633068 [14q22.1]	Hs.5 849 05	NM_001127 713.1	65	1544	12 - 13

ATP6V0D	ATPase H+ transporting V0 subunit d1	Chr.16: 67438014 - 67481186 [16q22.1]	Hs.1 068 76	NM_0 04691. 4	72	224	1 - 2
C1ORF21	chromosome 1 open reading frame 21	Chr.1: 184387016 - 184629021 [1q25.3]	Hs.4 971 59	NM_0 30806. 3	84	805	5-6
CFLAR	CASP8 and FADD like apoptosis regulator	Chr.2: 201116015 - 201172688 [2q33.1]	Hs.3 907 36	NM_0 01127 183.2	59	874	3-4
CFLAR-AS1	CFLAR antisense RNA 1	Chr.2: 201140289 - 201157792 [2q33.1]	Hs.6 646 13	NR_04 0030.1	103	883	6-7
CHP1	calcineurin like EF-hand protein 1	Chr.15: 41231149 - 41281887 [15q15.1]	Hs.4 062 34	NM_0 07236. 4	120	212	1-2
DDX55	DEAD-box helicase 55	Chr.12: 123602077 - 123620943 [12q24.31]	Hs.2 861 73	NM_0 20936. 2	74	183	1-2
DMD	dystrophin	Chr.X: 31119219 - 33339609 [Xp21.1]	Hs.4 959 12	NM_0 00109. 3	76	9607	63-64
DNAJC9	DnaJ heat shock protein family (Hsp40) member C9	Chr.10: 73241954 - 73247331 [10q22.2]	Hs.4 085 77	NM_0 15190. 4	102	198	1-1
ENOSF1	enolase superfamily member 1	Chr.18: 670012 - 712664 [18p11.32]	Hs.6 585 50	NM_0 01126 123.3	110	X	13-14
FANCL	Fanconi anemia complementation group L	Chr.2: 58159243 - 58241380 [2p16.1]	Hs.6 318 90	NM_0 01114 636.1	138	540	6-7
HJURP	Holliday junction recognition protein	Chr.2: 233836701 - 233854566 [2q37.1]	Hs.5 329 68	NM_0 18410. 4	52	399	4-5
HLA-DOA	major histocompatibility complex, class II, DO alpha	Chr.6: 33004182 - 33009612 [6p21.32]	Hs.6 319 91	NM_0 02119. 3	124	160	1-2

HLA-DRA	major histocompatibility complex, class II, DR alpha	Chr.6: 32439842 - 32445046 [6p21.32]	Hs.5 200 48	NM_019111.4	129	884	4-5
HNRNPA3P1	heterogeneous nuclear ribonucleoprotein A3 pseudogene 1	Chr.10: 43787412 - 43790417 [10q11.21]	Hs.6 329 56	NR_002726.2	99	2455	1-1
IL23A	interleukin 23 subunit alpha	Chr.12: 56334159 - 56340410 [12q13.3]	Hs.3 822 12	NM_016584.2	107	323	1-2
IQGAP1	IQ motif containing GTPase activating protein 1	Chr.15: 90388241 - 90502243 [15q26.1]	Hs.4 305 51	NM_003870.3	69	4562	34-35
LOC494127	NFYC pseudogene	Chr.9: 87083886 - 87085225 [9q21.33]	Hs.6 263 16	NR_036691.1	86	150	1-1
LOC646471	uncharacterized LOC646471	Chr.1: 25819954 - 25823606 [1p36.11]	Hs.7 272 71	NR_024498.1	80	2858	1-1
LOH12CR	loss of heterozygosity, 12, chromosomal region 2 (non-protein coding)	Chr.12: 12355408 - 12357067 [12p13.2]	Hs.6 755 3	NR_024061.1	69	140	1-2
MTRNR2L2_MTO1	CUSTOM PRIMER						
PBXIP1	PBX homeobox interacting protein 1	Chr.1: 154944080 - 154956163 [1q21.3]	Hs.5 058 06	NM_020524.3	60	189	2-3
RNF5	ring finger protein 5	Chr.6: 32178385 - 32180793 [6p21.32]	Hs.7 317 74	NM_006913.3	67	273	1-2
SERTAD2	SERTA domain containing 2	Chr.2: 64631621 - 64751091 [2p14]	Hs.5 915 69	NM_014755.2	91	288	1-2
SLC35G5	solute carrier family 35 member G5	Chr.8: 11330986 - 11332186 [8p23.1]	Hs.4 583 97	NM_054028.1	144	541	1-1
SPATS2L	spermatogenesis associated serine rich 2 like	Chr.2: 200305881 - 200482263 [2q33.1]	Hs.1 203 23	NM_001100422.1	89	1364	10-11

TDRD7	tudor domain containing 7	Chr.9: 97412020 - 97496125 [9q22.33]	Hs.1 938 42	NM_001302 884.1	59	3084	15-16
TOX4 (housekeeping gene)	TOX high mobility group box family member 4	Chr.14: 21477176 - 21499177 [14q11.2]	Hs.5 559 10	NM_001303 523.1	145	447	3-3
TPT1 (housekeeping gene)	tumor protein, translationally-controlled 1	Chr.13: 45333471 - 45341284 [13q14.13]	Hs.3 745 96	NM_001286 272.1	131	377	3-3
TXK	TXK tyrosine kinase	Chr.4: 48066393 - 48135322 [4p12]	Hs.4 796 69	NM_003328. 2	113	1265	11-12
YY2	YY2 transcription factor	Chr.X: 21855987 - 21858727 [Xp22.12]	Hs.6 736 01	NM_206923. 3	110	552	1-1

[0085] Diagnosis: Identification of samples as melanoma

[0086] In the test set 1, the data for the utility of the test to differentiate melanoma from controls are included in **Table 3**. The metrics are included in **FIG. 2**. These are: sensitivity >90%, specificity 100%, PPV 100%, NPV 87%. The overall accuracy is 94%. The tool can therefore differentiate between controls and aggressive and stable melanoma disease.

[0087] Table 3. Confusion matrix showing classification accuracy of the 5-model algorithm that determines whether a sample is a melanoma or a control in blood samples

Predicted/Reference	Melanoma	Control
Control	3	20
Melanoma	30	0
Sensitivity	91%	
Specificity	100%	
Positive Predictive Value	100%	
Negative Predictive Value	87%	
Accuracy	94%	

[0088] The test was evaluated in a second test set (test set 2) that included 74 controls and 92 melanoma patients. The mean Melanomx score in the melanoma group was 73 ± 31 versus 10 ± 8 in the control group (**FIG. 3A**). The receiver operator curve analysis demonstrated the score exhibited an area under the curve (AUC) of 0.96 (**FIG. 3B**) and the metrics were 88-100% (**FIG. 3C**).

[0089] Correlation with surgical removal of melanoma.

[0090] In the surgical series ($n = 46$), removal of melanoma decreased the score from 74 ± 16 to 20 ± 12 (**FIG. 4A**). Evaluation of the post-surgical group identified the score was significantly different between patients with no evidence of disease 13 ± 6 (not different to the control group) and those with residual disease (33 ± 9) (**FIG. 4B**). The Melanomx score can therefore define the extent of surgery and identify those who are surgical cures.

[0091] Evaluation of immune-therapy and targeted (BRAF inhibitor) therapy.

[0092] In the therapy series ($n = 30$), treatment with ipilimumab (immune therapy targeting CTLA-4), nivolumab (immune therapy targeting PD-1) or a BRAF inhibitor (vemurafenib) for 5 months significantly decreased the Melanomx scores from 88 ± 12 to 15 ± 5 ($p < 0.0001$), 35 ± 14 ($p < 0.001$) and 38 ± 21 ($p < 0.0001$), respectively (**FIG. 5**). All patients in the treated groups were stable. This indicates that the Melanomx score can therefore be used to measure the efficacy of both immune and targeted-therapy in melanoma and that a decrease in score correlated with response to therapeutic intervention.

[0093] Confirmation that gene expression is melanoma derived.

[0094] We confirmed that melanoma was the source for the blood-based gene expression assay by evaluating expression in different melanoma cell lines, in tumor tissue and by comparing expression in blood with tumor tissue collected at the same time-point during surgery.

[0095] All 28 genes were highly expressed in all 3 melanoma cell lines. Scores ranged from 94 ± 6 (Hs294T) to 82 ± 5 (CHL) to 42 ± 9 (A375) (**FIG. 6A**).

[0096] Spike-in experiments using these 3 cell lines and normal whole blood demonstrated that gene expression scores were detected when as few as 1 cell was spiked into 1 ml of blood. One single melanoma cell was detectable. Scores ranged from 21 ± 4 (A375) to 30 ± 5 (CHL) to

34±2 (H2294) (**FIG. 6B**). Scores were significantly elevated compared to control blood (no spike-in; $p < 0.0001$).

[0097] We then evaluated gene expression in tumor tissue. All 28 genes were highly expressed in melanoma compared to matched normal tissue and scores were significantly elevated 70±20 versus 4±3 ($p < 0.0001$) (**FIG. 7A**).

[0098] We also compared gene expression in tumor tissue and blood collected at the same time point. Gene expression was highly concordant (Pearson r : 0.74 – 0.83, median: 0.819) identifying gene expression in tumor tissue and blood was concordant (**FIG. 7B**).

[0099] References:

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Equivalents

[00100] While the present invention has been described in conjunction with the specific embodiments set forth above, many alternatives, modifications and other variations thereof will be apparent to those of ordinary skill in the art. All such alternatives, modifications and variations are intended to fall within the spirit and scope of the present invention.

[00101] Reference to any prior art in the specification is not an acknowledgement or suggestion that this prior art forms part of the common general knowledge in any jurisdiction

or that this prior art could reasonably be expected to be combined with any other piece of prior art by a skilled person in the art.

[00102] By way of clarification and for avoidance of doubt, as used herein and except where the context requires otherwise, the term "comprise" and variations of the term, such as "comprising", "comprises" and "comprised", are not intended to exclude further additions, components, integers or steps.

Claims:

1. A method for detecting a melanoma in a subject in need thereof, comprising:
 - determining the expression level of at least 29 RNA biomarkers from a test sample from the subject by contacting the test sample with a plurality of agents specific to detect the expression of the at least 29 RNA biomarkers, wherein the at least 29 RNA biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, YY2, and at least one housekeeping gene;
 - normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of the at least one housekeeping gene, thereby obtaining a normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2;
 - inputting each normalized expression level into an algorithm to generate a score;
 - comparing the score with a predetermined cutoff value; and
 - identifying the presence of a melanoma in the subject when the score is equal to or greater than the predetermined cutoff value or identifying the absence of a melanoma in the subject when the score is below the predetermined cutoff value, wherein the predetermined cutoff value is 20 on a scale of 0 to 100.

2. A method for evaluating the extent of surgical resection in a subject having a melanoma, comprising:
 - determining the expression level of at least 29 RNA biomarkers from a test sample from the subject after the surgical resection by contacting the test sample with a plurality of agents specific to detect the expression of the at least 29 RNA biomarkers, wherein the at least 29 RNA biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1,

RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, YY2, and at least one housekeeping gene;

normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of the at least one housekeeping gene, thereby obtaining a normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2;

inputting each normalized expression level into an algorithm to generate a score;

comparing the score with a predetermined cutoff value; and

identifying that the surgical resection does not remove the entire melanoma when the normalized expression level is equal to or greater than the predetermined cutoff value or identifying that the surgical resection removes the entire melanoma when the normalized expression level is below the predetermined cutoff value, wherein the predetermined cutoff value is 20 on a scale of 0 to 100.

3. The method of claim 2, wherein the method further comprises identifying that the risk of melanoma recurrence is high when the normalized expression level is equal to or greater than the predetermined cutoff value or identifies that the risk of melanoma recurrence is low when the normalized expression level is below the predetermined cutoff value.

4. The method of any one of the preceding claims, wherein the at least one housekeeping gene is selected from the group consisting of ALG9, SEPN, YWHAQ, VPS37A, PRRC2B, DOPEY2, NDUFB11, ND4, MRPL19, PSMC4, SF3A1, PUM1, ACTB, GAPD, GUSB, RPLP0, TFRC, MORF4L1, 18S, PPIA, PGK1, RPL13A, B2M, YWHAZ, SDHA, HPRT1, TOX4, and TPT1.

5. The method of claim 4, wherein the at least one housekeeping gene comprises TOX4 and TPT1.

6. The method of claim 4 or 5, wherein the normalized expression level is obtained by: normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of TOX4, thereby obtaining a first normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2;

normalizing the expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2 to the expression level of TPT1, thereby obtaining a second normalized expression level of each of ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2;

averaging the first normalized expression level and the second normalized expression level to obtain the normalized expression level.

7. A method for determining a response by a subject having a melanoma to a therapy, comprising:

determining a first expression level of at least 28 RNA biomarkers from a first test sample from the subject at a first time point by contacting the first test sample with a plurality of agents specific to detect the expression of the at least 28 RNA biomarkers, wherein the 28 RNA biomarkers comprise ATL1, ATP6V0D, C1ORF21, CFLAR, CFLAR-AS1, CHP1, DDX55, DMD, DNAJC9, ENOSF1, FANCL, HJURP, HLA-DOA, HLA-DRA, HNRNPA3P1, IL23A, IQGAP1, LOC494127, LOC646471, LOH12CR, PBXIP1, RNF5, SERTAD2, SLC35G5, SPATS2L, TDRD7, TXK, and YY2;

determining a second expression level of the at least 28 RNA biomarkers from a second test sample from the subject at a second time point by contacting the second test

sample with a plurality of agents specific to detect the expression of the at least 28 RNA biomarkers, wherein the second time point is after the first time point and after the administration of the therapy to the subject;

comparing the first expression level with the second expression level; and

identifying that the subject is responsive to the therapy when the second expression level is significantly decreased as compared to the first expression level.

8. The method of claim 7, wherein the first time point is
 - (a) prior to the administration of the therapy to the subject; or
 - (b) after the administration of the therapy to the subject.
9. The method of claim 7 or claim 8, wherein the therapy comprises
 - (a) an immunotherapy; and/or
 - (b) a targeted therapy.
10. The method of claim 9, wherein the targeted therapy comprises a BRAF inhibitor.
11. The method of any one of the preceding claims, having
 - (a) a sensitivity of greater than 90%; and/or
 - (b) a specificity of greater than 90%.
12. The method of any one of the preceding claims, wherein the melanoma is progressive.
13. The method of any one of the preceding claims, wherein the expression levels of the RNA biomarkers are detected by forming a complex between the biomarkers and labeled probes or primers.
14. The method of any one of the preceding claims, wherein RNA biomarkers are reverse transcribed to produce cDNA, and the produced cDNA expression level is detected.
15. The method of claim 14, wherein the cDNA is detected by forming a complex between the cDNA and labeled nucleic acids or primers.

16. The method of claim 13 or claim 15, wherein the label is a fluorescent label.
17. The method of any one of the preceding claims, wherein the test sample is blood, serum, plasma, or neoplastic tissue.
18. The method of any one of claims 1-6, wherein the predetermined cutoff value is derived from a plurality of reference samples obtained from subjects free of a neoplastic disease.
19. The method of claim 18, wherein each reference sample is blood, serum, plasma, or non-neoplastic tissue.
20. The method of any one of the preceding claims, wherein the subject in need thereof is a subject diagnosed with a melanoma, a subject having at least one melanoma symptom, or a subject having a predisposition or familial history for developing a melanoma.
21. The method of claim 20, wherein the subject is human.
22. The method of any one of claims 1-6, wherein the algorithm is a classifier built from one or more predictive classification algorithms.
23. The method of claim 22, wherein the one or more predictive classification algorithms is selected from XGB, RF, glmnet, cforest, CART, treebag, knn, nnet, SVM-radial, SVM-linear, NB, NNET, or mlp.
24. The method of claim 23, wherein the one or more predictive classification algorithms is XGB.

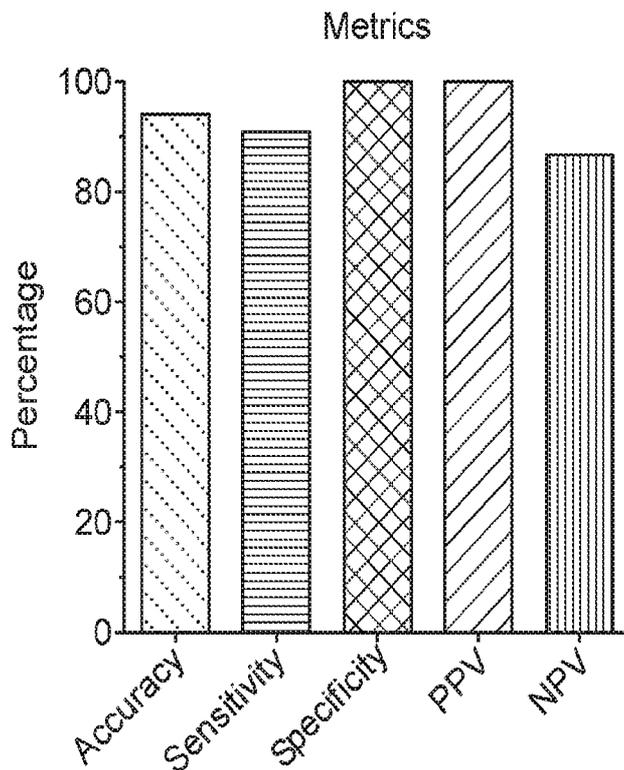
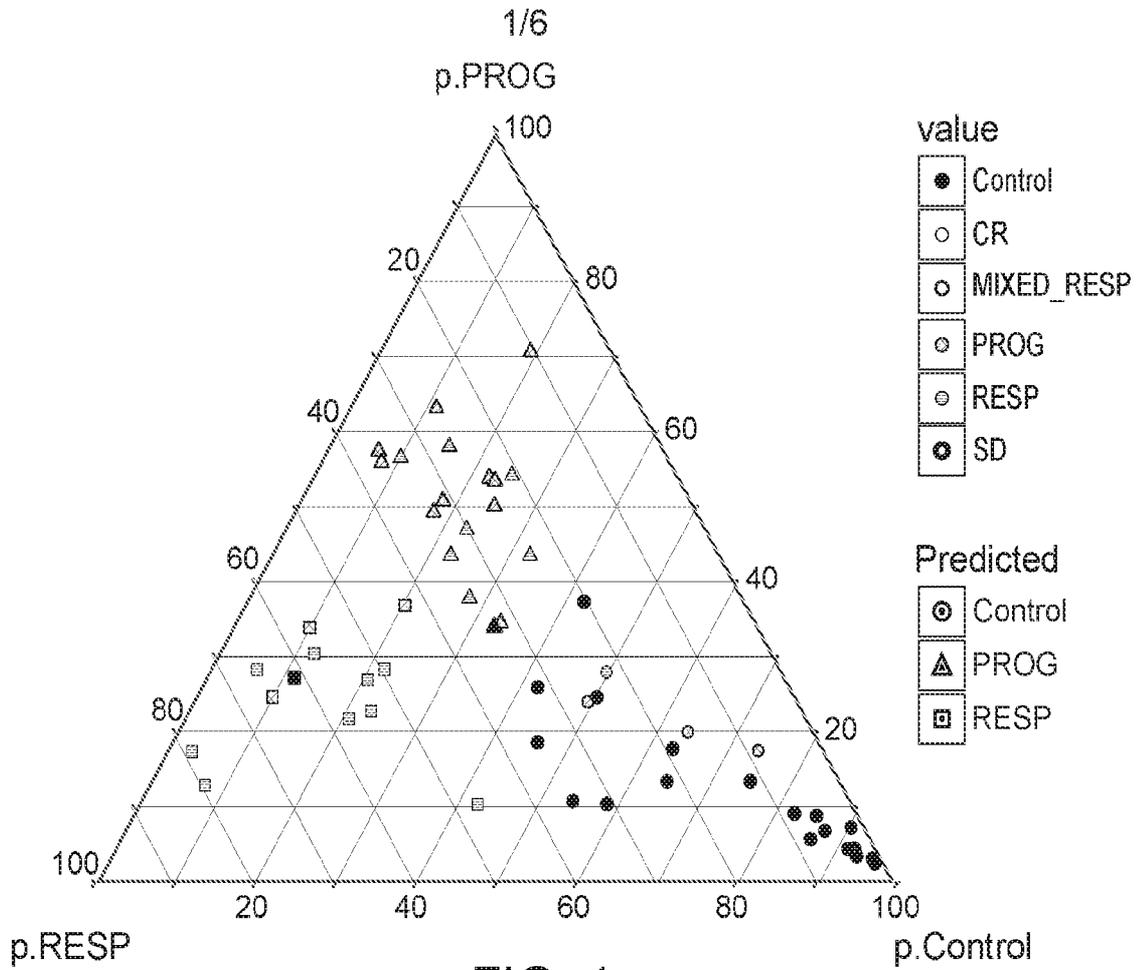


FIG. 2

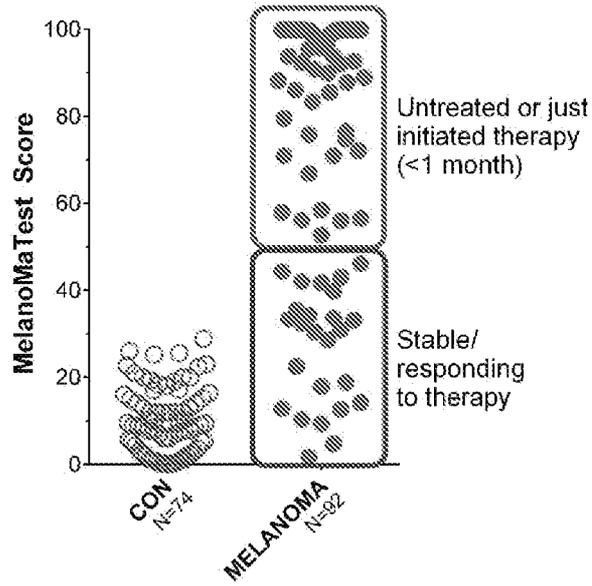


FIG. 3A

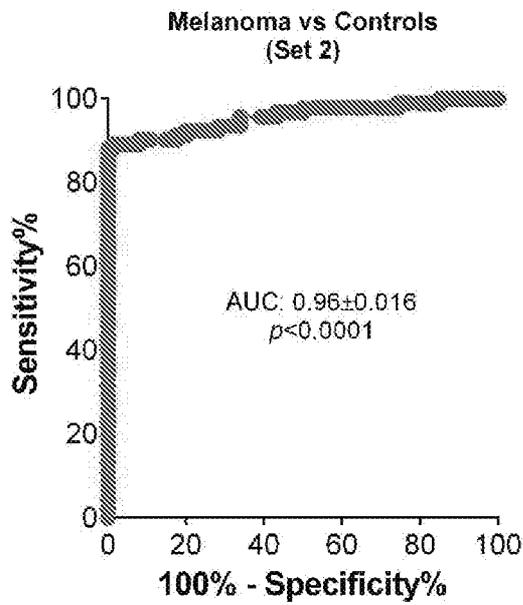


FIG. 3B

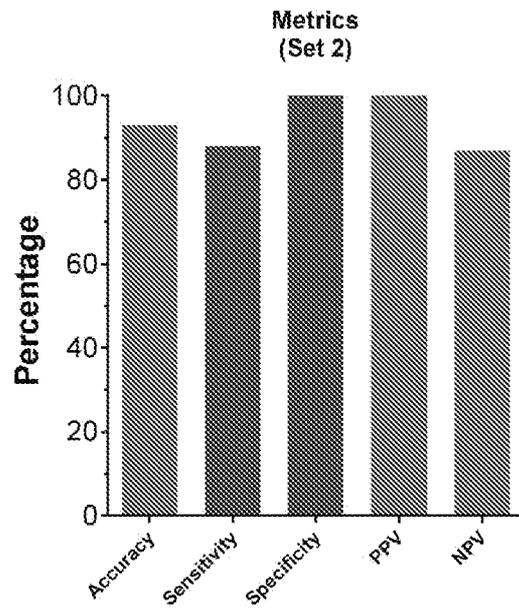


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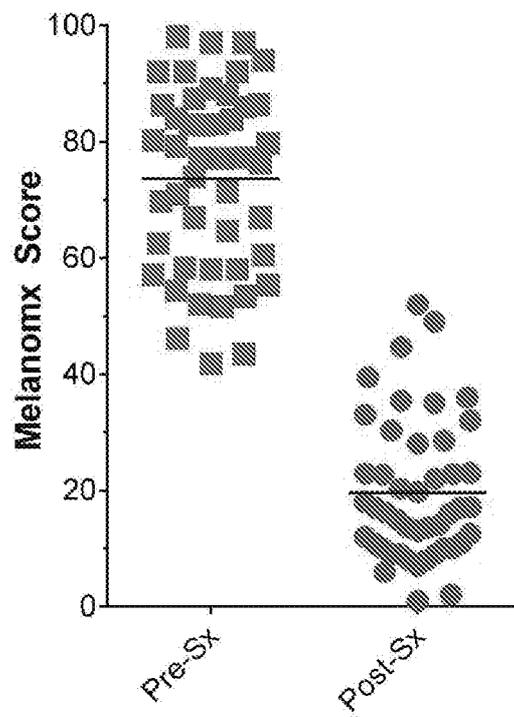


FIG. 4A

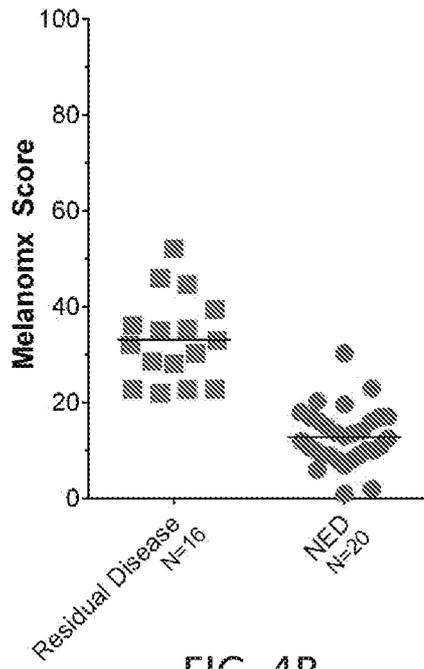


FIG. 4B

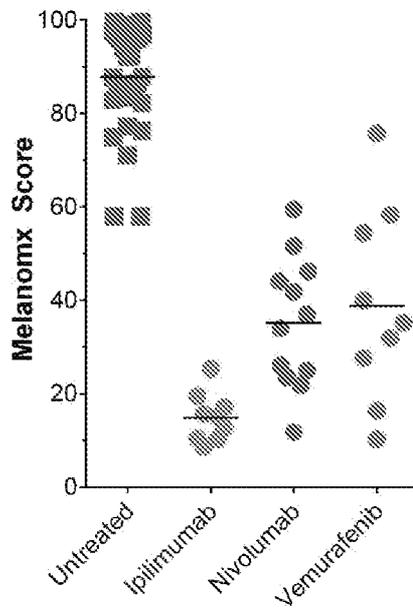


FIG. 5

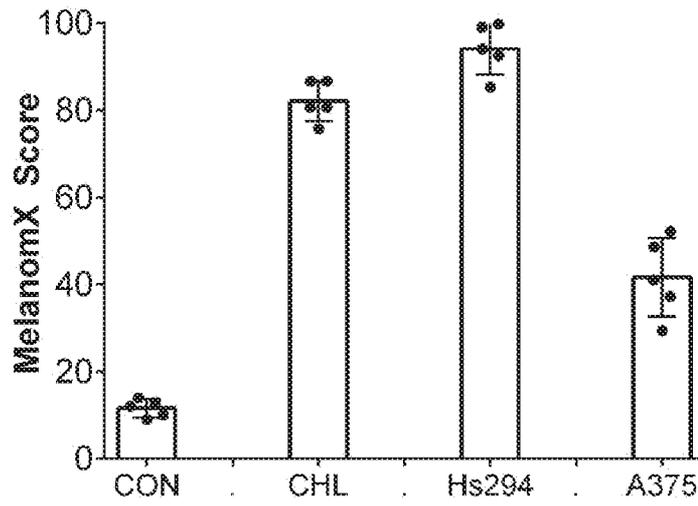


FIG. 6A

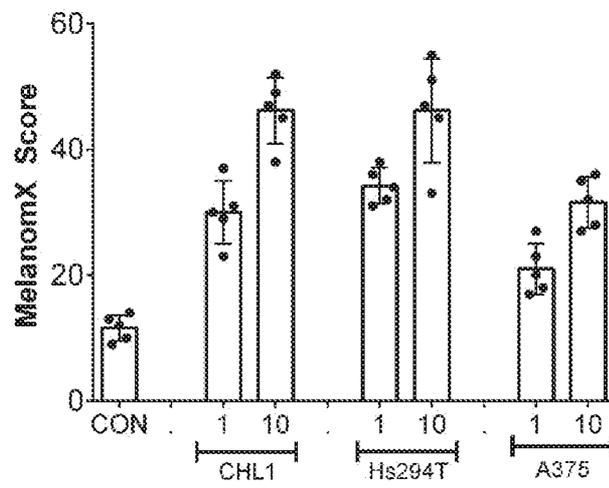


FIG. 6B

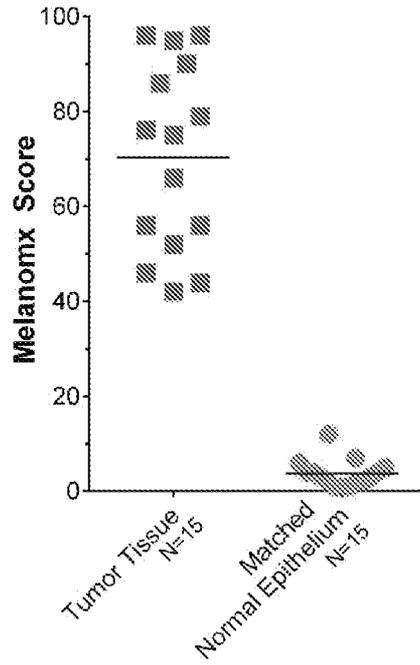


FIG. 7A

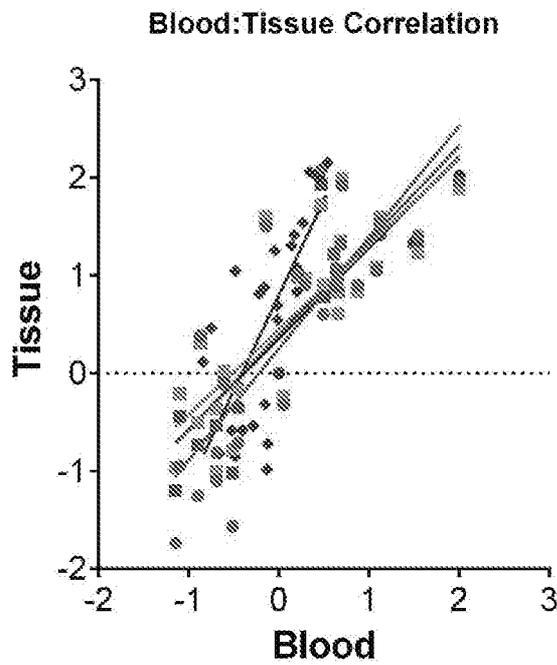


FIG. 7B

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