

(19) United States

(12) Patent Application Publication (10) Pub. No.: US 2025/0263708 A1 **THOMPSON**

Aug. 21, 2025 (43) **Pub. Date:**

(54) COMPOSITION FOR REGULATING PRODUCTION OF INTERFERING RIBONUCLEIC ACID

(71) Applicant: Wyvern Pharmaceuticals Inc., Calgary (CA)

(72) Inventor: Bradley G. THOMPSON, Calgary (CA)

(21) Appl. No.: 18/976,043

(22) Filed: Dec. 10, 2024

Related U.S. Application Data

(62) Division of application No. 18/582,317, filed on Feb. 20, 2024.

Publication Classification

(51) Int. Cl. (2010.01)C12N 15/113 C12N 15/86 (2006.01)

(52) U.S. Cl.

CPC C12N 15/1137 (2013.01); C12N 15/86 (2013.01); C12N 2310/141 (2013.01); C12N 2750/14143 (2013.01)

ABSTRACT (57)

Some embodiments of the present disclosure relate to one or more compositions that upregulate the production of one or more sequences of micro-interfering ribonucleic acid (miRNA). The miRNA may be complimentary to a sequence of target messenger RNA (mRNA) that encodes for a target biomolecule and the miRNA can cause the target mRNA to be degraded or inactivated, decreasing the bioavailability of the target biomolecule within a subject that is administered the one or more compositions. In some embodiments of the present disclosure, the target biomolecule is a complement or a factor. In some embodiments of the present disclosure, the target biomolecule is a complement such as complement C1q, complement C1r, complement C1s, complement C3 or complement C5. In some embodiments of the present disclosure, the target biomolecule is a factor such as Factor B, Factor D or Factor 10.

Specification includes a Sequence Listing.

COMPOSITION FOR REGULATING PRODUCTION OF INTERFERING RIBONUCLEIC ACID

[0001] This application contains a Sequence Listing electronically submitted via Patent Center to the United States Patent and Trademark Office as an XML Document file entitled "A8149442US—Sequence Listing.xml" created on 2024 Feb. 12 and having a size of 76,125 bytes. The information contained in the Sequence Listing is incorporated by reference herein.

TECHNICAL FIELD

[0002] The present disclosure generally relates to compositions for regulating production of interfering ribonucleic acid (RNA). In particular, the present disclosure relates to compositions for regulating gene expression and therefore, the production of interfering RNA that will suppress complement or factor expression.

BACKGROUND

[0003] Bioactive molecules, including complements and factors, are necessary for the homeostatic control of biological systems.

[0004] When bioactive molecules are over-expressed, under-expressed or mis-expressed, homeostasis is lost, and disease is often the result.

[0005] As such, it may be desirable to establish therapies, treatments and/or interventions that address when homeostasis and regulation of bioactive molecules is lost to prevent or treat the resulting disease.

SUMMARY

[0006] Some embodiments of the present disclosure relate to one or more compositions that upregulate the production of one or more sequences of micro-interfering ribonucleic acid (miRNA). The sequences of miRNA may be complimentary to a sequence of target messenger RNA (mRNA) that encodes for translation of a target biomolecule and the miRNA can cause the target mRNA to be degraded or inactivated, thereby causing a decrease in bioavailability of the target biomolecule because it is degraded or inactivated by the miRNA, thereby decreasing the bioavailability of the target biomolecule within a subject that is administered the one or more compositions. In some embodiments of the present disclosure, the target biomolecule is a complement or a factor. In some embodiments of the present disclosure, the target biomolecule is a complement such as complement C1q. In some embodiments of the present disclosure, the target biomolecule is a complement such as complement C1r. In some embodiments of the present disclosure, the target biomolecule is a complement such as complement C1s. In some embodiments of the present disclosure, the target biomolecule is a complement such as complement C3. In some embodiments of the present disclosure, the target biomolecule is a complement such as complement C5. In some embodiments of the present disclosure, the target biomolecule is a factor such as Factor B. In some embodiments of the present disclosure, the target biomolecule is a factor such as Factor D. In some embodiments of the present disclosure, the target biomolecule is a factor such as Factor [0007] In some embodiments of the present disclosure the compositions comprise a plasmid of deoxyribonucleic acid (DNA) that includes one or more insert sequences of nucleic acids that encode for the production of miRNA and a backbone sequence of nucleic acids that facilitates introduction of the one or more insert sequences into one or more of a subject's cells where it is expressed and/or replicated. Expression of the one or more insert sequences by one or more cells of the subject results in an increased production of the miRNA and, therefore, decreased translation or production of the target biomolecule by one or more of the subject's cells.

[0008] Some embodiments of the present disclosure relate to compositions that upregulate the production of miRNA that degrades, or causes degradation of, or inactivates or causes the inactivation of, the mRNA of the target biomolecule

[0009] Some embodiments of the present disclosure relate to a recombinant plasmid (RP). In some embodiments of the present disclosure, the RP comprises a nucleotide sequence of SEQ ID NO. 1 and SEQ ID NO. 2. The RP comprises a nucleotide sequence encoding one or more nucleotide sequences encoding a miRNA sequence that targets the mRNA of complement C1q.

[0010] Some embodiments of the present disclosure relate to a recombinant plasmid. In some embodiments of the present disclosure, the RP comprises a nucleotide sequence of SEQ ID NO. 1 and SEQ ID NO. 3. The RP comprises a nucleotide sequence encoding one or more nucleotide sequences encoding a miRNA sequence that targets the mRNA of complement C1r.

[0011] Some embodiments of the present disclosure relate to a recombinant plasmid. In some embodiments of the present disclosure, the RP comprises a nucleotide sequence of SEQ ID NO. 1 and SEQ ID NO. 4. The RP comprises a nucleotide sequence encoding one or more nucleotide sequences encoding a miRNA sequence that targets the mRNA of complement CIs.

[0012] Some embodiments of the present disclosure relate to a recombinant plasmid. In some embodiments of the present disclosure, the RP comprises a nucleotide sequence of SEQ ID NO. 1 and SEQ ID NO. 5. The RP comprises a nucleotide sequence encoding one or more nucleotide sequences encoding a miRNA sequence that targets the mRNA of complement C3.

[0013] Some embodiments of the present disclosure relate to a recombinant plasmid. In some embodiments of the present disclosure, the RP comprises a nucleotide sequence of SEQ ID NO. 1 and SEQ ID NO. 6. The RP comprises a nucleotide sequence encoding one or more nucleotide sequences encoding a miRNA sequence that targets the mRNA of complement C5.

[0014] Some embodiments of the present disclosure relate to a recombinant plasmid. In some embodiments of the present disclosure, the RP comprises a nucleotide sequence of SEQ ID NO. 1 and SEQ ID NO. 7. The RP comprises a nucleotide sequence encoding one or more nucleotide sequences encoding a miRNA sequence that targets the mRNA of Factor B.

[0015] Some embodiments of the present disclosure relate to a recombinant plasmid. In some embodiments of the present disclosure, the RP comprises a nucleotide sequence of SEQ ID NO. 1 and SEQ ID NO. 8. The RP comprises a

nucleotide sequence encoding one or more nucleotide sequences encoding a miRNA sequence that targets the mRNA of Factor D.

[0016] Some embodiments of the present disclosure relate to a recombinant plasmid. In some embodiments of the present disclosure, the RP comprises a nucleotide sequence of SEQ ID NO. 1 and SEQ ID NO. 9. The RP comprises a nucleotide sequence encoding one or more nucleotide sequences encoding a miRNA sequence that targets the mRNA of Factor 10.

[0017] Some embodiments of the present disclosure relate to a method of making a composition/target cell complex. The method comprising a step of administering a RP comprising SEQ ID NO. 1 and one of SEQ ID NO. 2, SEQ ID NO. 3, SEQ ID NO. 4, SEQ ID NO. 5, SEQ ID NO. 6, SEQ ID NO. 7, SEQ ID NO. 8, or SEQ ID NO. 9 to a target cell for forming the composition/target cell complex, wherein the composition/target cell complex causes the target cell to increase production of one or more sequences of miRNA that decreases production of a target biomolecule.

[0018] Embodiments of the present disclosure relate to at least one approach for inducing endogenous production of one or more sequences of miRNA that target and silence the mRNA of a target biomolecule, for example complement C1q. A first approach utilizes gene vectors containing nucleotide sequences for increasing the endogenous production of one or more sequences of miRNA, which are complete or partial sequences and/or combinations thereof, that target and silence the mRNA of complement C1q, which can be administered to a subject to increase the subject's production of one or more sequences of the miRNA.

[0019] Embodiments of the present disclosure relate to at least one approach for inducing endogenous production of one or more sequences of miRNA that target and silence the mRNA of a target biomolecule, for example complement C1r. A first approach utilizes gene vectors containing nucleotide sequences for increasing the endogenous production of one or more sequences of miRNA, which are complete or partial sequences and/or combinations thereof, that target and silence the mRNA of complement C1r, which can be administered to a subject to increase the subject's production of one or more sequences of the miRNA.

[0020] Embodiments of the present disclosure relate to at least one approach for inducing endogenous production of one or more sequences of miRNA that target and silence the mRNA of a target biomolecule, for example complement C1s. A first approach utilizes gene vectors containing nucleotide sequences for increasing the endogenous production of one or more sequences of miRNA, which are complete or partial sequences and/or combinations thereof, that target and silence the mRNA of complement C1s, which can be administered to a subject to increase the subject's production of one or more sequences of the miRNA.

[0021] Embodiments of the present disclosure relate to at least one approach for inducing endogenous production of one or more sequences of miRNA that target and silence the mRNA of a target biomolecule, for example complement C3. A first approach utilizes gene vectors containing nucleotide sequences for increasing the endogenous production of one or more sequences of miRNA, which are complete or partial sequences and/or combinations thereof, that target and silence the mRNA of complement C3, which can be administered to a subject to increase the subject's production of one or more sequences of the miRNA.

[0022] Embodiments of the present disclosure relate to at least one approach for inducing endogenous production of one or more sequences of miRNA that target and silence the mRNA of a target biomolecule, for example complement C5. A first approach utilizes gene vectors containing nucleotide sequences for increasing the endogenous production of one or more sequences of miRNA, which are complete or partial sequences and/or combinations thereof, that target and silence the mRNA of complement C5, which can be administered to a subject to increase the subject's production of one or more sequences of the miRNA.

[0023] Embodiments of the present disclosure relate to at least one approach for inducing endogenous production of one or more sequences of miRNA that target and silence the mRNA of a target biomolecule, for example Factor B. A first approach utilizes gene vectors containing nucleotide sequences for increasing the endogenous production of one or more sequences of miRNA, which are complete or partial sequences and/or combinations thereof, that target and silence the mRNA of Factor B, which can be administered to a subject to increase the subject's production of one or more sequences of the miRNA.

[0024] Embodiments of the present disclosure relate to at least one approach for inducing endogenous production of one or more sequences of miRNA that target and silence the mRNA of a target biomolecule, for example Factor D. A first approach utilizes gene vectors containing nucleotide sequences for increasing the endogenous production of one or more sequences of miRNA, which are complete or partial sequences and/or combinations thereof, that target and silence the mRNA of Factor D, which can be administered to a subject to increase the subject's production of one or more sequences of the miRNA.

[0025] Embodiments of the present disclosure relate to at least one approach for inducing endogenous production of one or more sequences of miRNA that target and silence the mRNA of a target biomolecule, for example Factor 10. A first approach utilizes gene vectors containing nucleotide sequences for increasing the endogenous production of one or more sequences of miRNA, which are complete or partial sequences and/or combinations thereof, that target and silence the mRNA of Factor 10, which can be administered to a subject to increase the subject's production of one or more sequences of the miRNA.

DETAILED DESCRIPTION

[0026] Unless defined otherwise, all technical and scientific terms used therein have the meanings that would be commonly understood by one of skill in the art in the context of the present description. Although any methods and materials similar or equivalent to those described therein can also be used in the practice or testing of the present disclosure, the preferred compositions, methods and materials are now described. All publications mentioned therein are incorporated therein by reference to disclose and describe the methods and/or materials in connection with which the publications are cited.

[0027] As used therein, the singular forms "a", "an", and "the" include plural references unless the context clearly dictates otherwise. For example, reference to "a composition" includes one or more compositions and reference to "a subject" or "the subject" includes one or more subjects.

[0028] As used therein, the terms "about" or "approximately" refer to within about 25%, preferably within about

20%, preferably within about 15%, preferably within about 10%, preferably within about 5% of a given value or range. It is understood that such a variation is always included in any given value provided therein, whether or not it is specifically referred to.

[0029] As used therein, the term "ameliorate" refers to improve and/or to make better and/or to make more satisfactory.

[0030] As used therein, the term "cell" refers to a single cell as well as a plurality of cells or a population of the same cell type or different cell types. Administering a composition to a cell includes in vivo, in vitro and ex vivo administrations and/or combinations thereof.

[0031] As used therein, the term "complex" refers to an association, either direct or indirect, between one or more particles of a composition and one or more target cells. This association results in a change in the metabolism of the target cell. As used therein, the phrase "change in metabolism" refers to an increase or a decrease in the one or more target cells' production of one or more proteins, and/or any post-translational modifications of one or more proteins.

[0032] As used therein, the term "composition" refers to a substance that, when administered to a subject, causes one or more chemical reactions and/or one or more physical reactions and/or one or more biological reactions in the subject. In some embodiments of the present disclosure, the composition is a plasmid vector.

[0033] As used therein, the term "endogenous" refers to the production and/or modification of a molecule that originates within a subject.

[0034] As used therein, the term "exogenous" refers to a molecule that is within a subject but that did not originate within the subject. As used therein, the terms "production", "producing" and "produce" refer to the synthesis and/or replication of DNA, the transcription of one or more sequences of RNA, the translation of one or more amino acid sequences, the post-translational modifications of an amino acid sequence, and/or the production of one or more regulatory molecules that can influence the production and/or functionality of an effector molecule or an effector cell. For clarity, "production" is also used therein to refer to the functionality of a regulatory molecule, unless the context reasonably indicates otherwise.

[0035] As used therein, the term "subject" refers to any therapeutic target that receives the composition. The subject can be a vertebrate, for example, a mammal including a human. The term "subject" does not denote a particular age or sex. The term "subject" also refers to one or more cells of an organism, an in vitro culture of one or more tissue types, an in vitro culture of one or more cell types, ex vivo preparations, and/or a sample of biological materials such as tissue, and/or biological fluids.

[0036] As used therein, the term "target biomolecule" refers to a complement or factor that is found within a subject. A biomolecule may be endogenous or exogenous to a subject and when bioavailable the biomolecule may inhibit or stimulate a biological process within the subject.

[0037] As used therein, the term "target cell" refers to one or more cells and/or cell types that are deleteriously affected, either directly or indirectly, by a dysregulated biomolecule. The term "target cell" also refers to cells that are not deleteriously affected but that are the cells in which it is desired that the composition interacts.

[0038] As used therein, the term "therapeutically effective amount" refers to the amount of the composition used that is of sufficient quantity to ameliorate, treat and/or inhibit one or more of a disease, disorder or a symptom thereof. The "therapeutically effective amount" will vary depending on the composition used, the route of administration of the composition and the severity of the disease, disorder or symptom thereof. The subject's age, weight and genetic make-up may also influence the amount of the composition that will be a therapeutically effective amount.

[0039] As used therein, the terms "treat", "treatment" and "treating" refer to obtaining a desired pharmacologic and/or physiologic effect. The effect may be prophylactic in terms of completely or partially preventing an occurrence of a disease, disorder or symptom thereof and/or the effect may be therapeutic in providing a partial or complete amelioration or inhibition of a disease, disorder, or symptom thereof. Additionally, the term "treatment" refers to any treatment of a disease, disorder, or symptom thereof in a subject and includes: (a) preventing the disease from occurring in a subject which may be predisposed to the disease but has not yet been diagnosed as having it; (b) inhibiting the disease, i.e., arresting its development; and (c) ameliorating the disease.

[0040] As used therein, the terms "unit dosage form" and "unit dose" refer to a physically discrete unit that is suitable as a unitary dose for patients. Each unit contains a predetermined quantity of the composition and optionally, one or more suitable pharmaceutically acceptable carriers, one or more excipients, one or more additional active ingredients, or combinations thereof. The amount of composition within each unit is a therapeutically effective amount.

[0041] Where a range of values is provided therein, it is understood that each intervening value, to the tenth of the unit of the lower limit unless the context clearly dictates otherwise, between the upper and lower limit of that range and any other stated or intervening value in that stated range, is encompassed within the disclosure. The upper and lower limits of these smaller ranges may independently be included in the smaller ranges, and are also, encompassed within the disclosure, subject to any specifically excluded limit in the stated range. Where the stated range includes one or both of the limits, ranges excluding either or both of those included limits are also included in the disclosure.

[0042] In some embodiments of the present disclosure, a composition is a recombinant plasmid (RP) for introducing genetic material, such as one or more nucleotide sequences, into a target cell for reproduction or transcription of an insert that comprises one or more nucleotide sequences that are carried within the RP. In some embodiments of the present disclosure, the RP is delivered without a carrier, by a viral vector, by a protein coat, or by a lipid vesicle. In some embodiments of the present disclosure, the vector is an adeno-associated virus vector.

[0043] In some embodiments of the present disclosure, the insert comprises one or more nucleotide sequences that encode for production of at least one sequence of miRNA that decreases the production of target biomolecules. The miRNA may, directly or indirectly, bind to and degrade the target mRNA or otherwise inactivate the target mRNA so that less or none of the target-biomolecule protein is produced.

[0044] In some embodiments of the present disclosure, the target biomolecule is the complement C1q.

[0045] In some embodiments of the present disclosure, the target biomolecule is the complement C1r.

[0046] In some embodiments of the present disclosure, the target biomolecule is the complement C1s.

[0047] In some embodiments of the present disclosure, the target biomolecule is the complement C3.

[0048] In some embodiments of the present disclosure, the target biomolecule is the complement C5.

[0049] In some embodiments of the present disclosure, the target biomolecule is Factor B.

 $\cite{[0050]}$ In some embodiments of the present disclosure, the target biomolecule is Factor D.

[0051] In some embodiments of the present disclosure, the target biomolecule is Factor 10.

[0052] In some embodiments of the present disclosure, the insert comprises one or more nucleotide sequences that each encode for one or more miRNA sequences that may be complimentary to and degrade, or cause degradation of, mRNA of the target biomolecule.

[0053] Some embodiments of the present disclosure relate to a composition that can be administered to a subject with a condition that results, directly or indirectly, from the dysregulated production of a biomolecule. When a therapeutically effective amount of the composition is administered to the subject, the subject may change production and/or functionality of one or more biomolecules.

[0054] In some embodiments of the present disclosure, the subject may respond to receiving the therapeutic amount of the composition by changing production and/or functionality of one or more intermediary molecules by changing production of one or more DNA sequences, one or more RNA sequences, and/or one or more proteins that regulate the levels and/or functionality of the one or more intermediary molecules. The one or more intermediary molecules regulate the subject's levels and/or functionality of the one or more biomolecules.

[0055] In some embodiments of the present disclosure, administering a therapeutic amount of the composition to a subject upregulates the production, functionality or both one or more sequences of miRNA that each target the mRNA of one or more target biomolecules. In some embodiments of the present disclosure, there are one, two, three, four, five, or six miRNA sequences that each are complimentary to and degrade, or cause degradation of, one biomolecule, such as the mRNA of complement C1q, complement C1r, complement C1s, complement C3, complement C5, Factor B, Factor D, or Factor 10. In some embodiments of the present disclosure, the composition may comprise multiple copies of the same nucleotide sequence of miRNA.

[0056] In some embodiments of the present disclosure, the composition is an RP that may be used for gene therapy. The gene therapy is useful for increasing the subject's endogenous production of one or more sequences of miRNA that target the mRNA of a target biomolecule. For example, the RP can contain one or more nucleotide sequences that cause increased production of one or more nucleotide sequences that cause an increased production of one or more miRNA

sequences that are each complimentary to and degrade, or cause degradation of, or inactivate, or cause inactivation of, one biomolecule, such as complement C1q, complement C1r, complement Cis, complement C3, complement C5, Factor B, Factor D, or Factor 10.

[0057] In some embodiments of the present disclosure, the delivery vehicle of the RP used for gene therapy may be a vector that comprises a virus that can be enveloped, or not (unenveloped), replication effective or not (replication ineffective), or combinations thereof. In some embodiments of the present disclosure, the vector is a virus that is not enveloped and not replication effective. In some embodiments of the present disclosure, the vector is a virus of the Parvoviridae family. In some embodiments of the present disclosure, the vector is a virus of the present disclosure, the vector is an adeno-associated virus (AAV). In some embodiments of the present disclosure, the vector is a recombinant AAV. In some embodiments of the present disclosure, the vector is a recombinant AAV6.2FF.

[0058] In some embodiments of the present disclosure, the delivery vehicle of the RP used for gene therapy may be a protein coat.

[0059] In some embodiments of the present disclosure, the delivery vehicle of the RP used for gene therapy may be a lipid vesicle.

[0060] The embodiments of the present disclosure also relate to administering a therapeutically effective amount of the composition. In some embodiments of the present disclosure, the therapeutically effective amount of the composition that is administered to a patient is between about 10 and about $1\times10^{16}~\mathrm{TCID}_{50}/\mathrm{kg}$ (50% tissue culture infective dose per kilogram of the patient's body mass). In some embodiments of the present disclosure, the therapeutically effective amount of the composition that is administered to the patient is about 1×10^{13} TCID₅₀/kg. In some embodiments of the present disclosure, the therapeutically effective amount of the composition that is administered to a patient is measured in TPC/kg (total particle count of the composition per kilogram of the patient's body mass). In some embodiments the therapeutically effective amount of the composition is between about 10 and about 1×10^{16} TCP/kg. [0061] Some embodiments of the present disclosure relate to an adeno-associated virus (AAV) genome consisting of a RP that when operable inside a target cell will cause the target cell to produce a miRNA sequence that downregulates production of a biomolecule, with examples being complement C1q, complement C1r, complement C1s, complement C3, complement C5, Factor B, Factor D, or Factor 10. The RP is comprised of AAV2 inverted terminal repeats (ITRs), a composite CASI promoter, a human growth hormone (HGH) signal peptide followed by a miRNA expression cassette containing up to six different miRNAs targeting the mRNA of complement C1q, complement C1r, complement C1s, complement C3, complement C5, Factor B, Factor D, or Factor 10, followed by a Woodchuck Hepatitis Virus post-transcriptional regulatory element (WPRE) and a Simian virus 40 (SV40) polyadenylation (polyA) signal.

 $\tt CCCGTATGGCTTTCATTTTCTCCTCCTTGTATAAATCCTGGTTGCTGTCTCTTTATGAG$ ${\tt ACCCCACTGGTTGGGGCATTGCCACCACCTGTCAGCTCCTTTCCGGGACTTTCGCTT}$ $\tt CAGGGGCTCGGCTGTTGGGCACTGACAATTCCGTGGTGTTGTCGGGGAAATCATCGT$ $\tt CCTTTCCTTGGCTGCTCGCCTGTGTTGCCACCTGGATTCTGCGCGGGACGTCCTTCTG$ CTGCGGCCTCTTCCGCGTCTTCGCCCTCAGACGAGTCGGATCTCCCTTTGGG $\tt CCGCCTCCCCGCCTAAGCTTATCGATACCGTCGAGATCTAACTTGTTTATTGCAGCTT$ ATAATGGTTACAAATAAAGCAATAGCATCACAAATTTCACAAATAAAGCATTTTTTT CACTGCATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTTATCATGTCTGGAT CTCGACCTCGACTAGAGCATGGCTACGTAGATAAGTAGCATGGCGGGTTAATCATTA TCAGTGAGCGAGCGCGCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG CCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAATGGCGATTCCGTTGCAATGGCTG $\tt GCGGTAATATTGTTCTGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTC$ ${\tt AGGCAAGTGATGTTATTACTAATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTG}$ ATGGACAGACTCTTTTACTCGGTGGCCTCACTGATTATAAAAACACTTCTCAGGATT $\tt CTGGCGTACCGTTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTC$ TGATTCTAACGAGGAAAGCACGTTATACGTGCTCGTCAAAGCAACCATAGTACGCG $\tt CCCTGTAGCGGCGCATTAAGCGCGGGGGGGGTGTGGTGGTTACGCGCAGCGTGACCGC$ ${\tt ACGTTCGCCGGCTTTCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGA}$ $\tt TTTAGTGCTTTACGGCACCTCGACCCCAAAAAACTTGATTAGGGTGATGGTTCACGT$ ${\tt AGTGGGCCATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCT}$ TTTTGATTTATAAGGGATTTTGCCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATT TAACAAAATTTAACGCGAATTTTAACAAAATATTAACGTTTACAATTTAAATATTT GCTTATACAATCTTCCTGTTTTTTGGGGCTTTTCTGATTATCAACCGGGGTACATATGA $\tt TTGACATGCTAGTTTTACGATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTC$ AGGCAATGACCTGATAGCCTTTGTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGC $\tt ATGAATTTATCAGCTAGAACGGTTGAATATCATATTGATGGTGATTTGACTGTCTCCC$ $\tt GGCCTTTCTCACCCGTTTGAATCTTTACCTACACATTACTCAGGCATTGCATTTAAAA$ TATATGAGGGTTCTAAAAATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAA ${\tt AAGTATTACAGGGTCATAATGTTTTTGGTACAACCGATTTAGCTTTATGCTCTGAGG}$ CTTTATTGCTTAATTTTGCTAATTCTTTGCCTTGCCTGTATGATTTATTGGATGTTGGA ATTCCTGATGCGGTATTTTCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGG

 $\tt CCAACACCCGCTGACGCGCCCTGACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGA$ CAAGCTGTGACCGTCTCCGGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCACCG AAACGCGCGAGACGAAAGGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATG ATAATAATGGTTTCTTAGACGTCAGGTGGCACTTTTCGGGGAAATGTGCGCGGAACC CCTATTTGTTTATTTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAAC CCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCC GTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTGCCTTCCTGTTTTTGCTCACCCAGAA ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACAT CGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTT GCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGAG TACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATG CGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTC GCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGAC ACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACT ${\tt AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGG}$ ${\tt AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGC}$ $\tt CCTCCCGTATCGTAGTTATCTACACGACGGGGGGGTCAGGCAACTATGGATGAACGA$ ${\tt AATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGTAACTGTCAGAC}$ CAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAATTTAAAAGGA TCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTC GTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTT $\tt TTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAAACCACCGCTACCAGCGGTGGT$ $\tt TTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAGCAG$ AGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAA ${\tt GAACTCTGTAGCACCGCCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGC}$ TGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACCGGA TAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGGAGC GAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACG CTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAG GAGAGCGCACGAGGGAGCTTCCAGGGGGAAACGCCTGGTATCTTTATAGTCCTGTC AGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGG CCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTAC CGCCTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGCGAGT CAGTGAGCGAGGAAGCGGAAGAGCGCCCAATACGCAAACCGCCTCTCCCCGCGCGT

TGGCCGATTCATTAATGCAGCAGCTGCGCGCTCGCTCACTGAGGCCGCCCGGG $\tt CGCAGAGAGGGGAGTGGCCAACTCCATCACTAGGGGTTCCTTGTAGTTAATGATTAAC$ $\tt CCGCCATGCTACTTATCTACGTAGCCATGCTCTAGGACATTGATTATTGACTAGTGG$ AGTTCCGCGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACC CCCGCCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTT ${\tt TCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATC}$ ${\tt AAGTGTATCATATGCCAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCG}$ $\tt CCTGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTACTTGGCAGTACATCTA$ CGTATTAGTCATCGCTATTACCATGGTCGAGGTGAGCCCCACGTTCTGCTTCACTCTC TGCAGCGATGGGGGCGGGGGGGGGGGGGGGGCGCGCCCAGGCGGGGCGGGCGGG GCGAGGGGCGGGCGGGCGAGGCGGAGAGGTGCGGCGGCAGCCAATCAGAGCGG CGCGCTCCGAAAGTTTCCTTTTATGGCGAGGCGGCGGCGGCGGCCGCCCTATAAAA AGCGAAGCGCGCGGGGGGGGGTCGCTGCGCGCTGCCTTCGCCCCGTGCCCCGC $\tt GGACGCTCAGGACAGCGGCCCGCTGCTCATAAGACTCGGCCTTAGAACCCCAGTAT$ $\tt CAGCAGAAGGACATTTTAGGACGGGACTTGGGTGACTCTAGGGCACTGGTTTTCTTT$ $\tt CCAGAGAGCGGAACAGGCGAGGAAAAGTAGTCCCTTCTCGGCGATTCTGCGGAGGG$ ATCTCCGTGGGGCGGTGAACGCCGATGATGCCTCTACTAACCATGTTCATGTTTTCTT $\tt TTTTTTCTACAGGTCCTGGGTGACGAACAGGGTACC$ SEQ ID NO. 2 (miRNA expression cassette No. 2 - complement C1q): $\tt GCCACCATGGCCACCGGCTCTCGCACAAGCCTGCTGCTGCTTTCGGACTGCTGTGC$ $\tt CTGCCTTGGCTCCAGGAGGGCTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGG$ ${\tt AGGCTTGCTGAAGGCTGTATGCTGAGATCTTCGGTTGCACCATGCTCGTTTTGGCCTC}$ TGACTGACGAGCATGGTGACCGAAGATCTCAGGACACAAGGCCTGTTACTAGCACT CACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGACAG ATCTTCGATGTCACCATGCGTTTTGGCCTCTGACTGACGCATGGTGACCGAAGATCT GTCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTCTAGCCTG GAGGCTTGCTGAAGGCTGTATGCTGACAGATCTTCGAAGTCACCATGCGTTTTGGCC TCACATGGAACAAATGGCCTCTCTAGAAT SEQ ID NO. 3 (miRNA expression cassette No. 3 - complement C1r):

GCCACCATGGCCACCGGCTCTCGCACAAGCCTGCTGCTTTCGGACTGCTGTGC
CTGCCTTGGCTCCAGGAGGGCTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGG

3 '

GCCACCATGGCCACCGGCTCTCGCACAAGCCTGCTGGCTTTCGGACTGCTGTGC
CTGCCTTGGCTCCAGGAGGGCTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGG
AGGCTTGCTGAAGGCTGTATGCTGATAGTTCGGGCTGTCAGAATTTCGTTTTGGCCT
CTGACTGACGAAATTCTGAGCCCGAACTATCAGGACACAAGGCCTGTTACTAGCACT
CACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGAGAT
GAATGCCTGATAGCCTTCCGTTTTGGCCTCTGACTGACGGAAGGCTATGGCATTCAT
CTCAGGACCACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTCTAGCCTG
GAGGCTTGCTGAAGGCTGTATGCTGAGATGAATAGCCTTCCGTTTTGGCC
TCTGACTGACGGAAGGCTATGCTGAGATTCATCTCAGGACCAAATAGCCTTTACTAGCAC

SEQ ID NO. 4 (miRNA expression cassette No. 4 - complement C1s):

3 '

TCACATGGAACAAATGGCCTCTCTAGAAT

SEQ ID NO. 5 (miRNA expression cassette No. 5 - complement C3): 5'
GCCACCATGGCCACCGGCTCTCGCACAAGCCTGCTGGCTTTCGGACTGCTGTGC

CTGCCTTGGCTCCAGGAGGGCTCCGCCGCTAGCATCGCTATCCGCTATGTGCTGG

AGGCTTGCTGAAGGCTGTATGCTGTTCAGATCATCTGGGTATCCGGCGTTTTGGCCT

CTGACTGACGCCGGATACCGATGATCTGAACAGGACACAAGGCCTGTTACTAGCAC

TCACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGAATA

ATCAGATGGTGTTGCGATCGTTTTGGCCTCTGACTGACGATCGCAACACCCTGATTA

TTCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTCTAGCCTGG

AGGCTTGCTGAAGGCTGTATGCTGTACTGATGCACAATTTAAACGCCGTTTTTGGCCT

CTGACTGACGGCGTTTAAAGTGCATCAGTACAGGACACAAGGCCTGTTACTAGCACT

CACATGGAACAAATGGCCTCTCTAGAAT

3 '

SEQ ID NO. 6 (miRNA expression cassette No. 6 - complement C5): 5'

3 '

SEQ ID NO. 7 (miRNA expression cassette No. 7 - Factor B): 5'

GCCACCATGGCCACCGGCTCTCGCACAAGCCTGCTGCTGCTTTCGGACTGCTGTGC
CTGCCTTGGCTCCAGGAGGGCTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGG
AGGCTTGCTGAAGGCTGTATGCTGACATAAAGCTATGTCCTGGCAGCGTTTTGGCCT
CTGACTGACGCCAGGATAGCTTTATGTCAGGACACAAGGCCTGTTACTAGCACT
CACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGATACA
GCAGATATCGCGAATTTCGTTTTGGCCTCTGACTGACGAAATTCGCGATCTGCTGTA
TCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTCTAGCCTGG
AGGCTTGCTGAAGGCTGTATGCTGTTTCAGTTTAAAATCAGCGCCACGTTTTGGCCT
CTGACTGACGTGGCGCTGATTAAACTGAAACAGGACACAAGGCCTGTTACTAGCAC
TCACATGGAACAAATGGCCTCTCTAGAAT

3 '

SEQ ID NO. 8 (miRNA expression cassette No. 8 - Factor D): 5'

GCCACCATGGCCACCGGCTCTCGCACAAGCCTGCTGCTGGCTTTCGGACTGCTGTGC
CTGCCTTGGCTCCAGGAGGGCTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGG
AGGCTTGCTGAAGGCTGTATGCTGATACAGGCGTTTGTGCTCGTTTTTGGCCTC
TGACTGACGAACCGAGCAAACGCCTGTATCAGGACACAAGGCCTGTTACTAGCACT
CACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGTCATG
ATCAATATGGTATCCGGCGTTTTGGCCTCTGACTGACGCCGGATACCATTGATCATG
ACAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTCTAGCCTGG
AGGCTTGCTGAAGGCTGTATGCTGTCATGATCAATAAGGTATCCGGCGTTTTGGCCT
CTGACTGACGCCGGATACCATTGATCATGACAGGACACAAGGCCTGTTACTAGCACT
CACATGGAACAAATGGCCTCTCTAGAAT

3 !

SEQ ID NO. 9 (miRNA expression cassette No. 9 - Factor 10): 5'

GCCACCATGGCCACCGGCTCTCGCACAAGCCTGCTGGCTTTCGGACTGCTGTGC
CTGCCTTGGCTCCAGGAGGGCTCCGCCGCTAGCATACCGTCGCTATGTGCTGG
AGGCTTGCTGAAGGCTGTATGCTGATAAACAGGCTTGTTCGCCCAGCGTTTTGGCCT
CTGACTGACGCTGGCGAAAGCCTGTTTATCAGGACACAAGGCCTGTTACTAGCACT

 ${\tt CACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGTCATA}$ GCTGCAATGGTTTCTTCCGTTTTGGCCTCTGACTGACGGAAGAAACCTGCAGCTATG ACAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTCTAGCCTGG ${\tt AGGCTTGCTGAAGGCTGTATGCTGACCACTTCCACAATTCATGCACCGTTTTGGCCT}$ $\tt CTGACTGACGGTGCATGAAGTGGAAGTGGTCAGGACACAAGGCCTGTTACTAGCAC$ TCACATGGAACAAATGGCCTCTCTAGAAT

SEQ ID NO. 10 = SEQ ID NO. 1 + SEQ ID NO. 2

AATCAACCTCTGGATTACAAAATTTGTGAAAGATTGACTGGTATTCTTAACTATGTT GCTCCTTTTACGCTATGTGGATACGCTGCTTTAATGCCTTTGTATCATGCTATTGCTT CCCGTATGGCTTTCATTTTCTCCTCCTTGTATAAATCCTGGTTGCTGTCTCTTTATGAG GAGTTGTGGCCCGTTGTCAGGCAACGTGGCGTGTGTGCACTGTGTTTTGCTGACGCA ACCCCCACTGGTTGGGGCATTGCCACCACCTGTCAGCTCCTTTCCGGGACTTTCGCTT CAGGGGCTCGGCTGTTGGGCACTGACAATTCCGTGGTGTTGTCGGGGAAATCATCGT CCTTTCCTTGGCTGCTCGCCTGTGTTGCCACCTGGATTCTGCGCGGGACGTCCTTCTG $\tt CCGCCTCCCCGCCTAAGCTTATCGATACCGTCGAGATCTAACTTGTTTATTGCAGCTT$ ATAATGGTTACAAATAAAGCAATAGCATCACAAATTTCACAAATAAAGCATTTTTTT CACTGCATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTTATCATGTCTGGAT CTCGACCTCGACTAGAGCATGGCTACGTAGATAAGTAGCATGGCGGGTTAATCATTA ACTACAAGGAACCCCTAGTGATGGAGTTGGCCACTCCCTCTCTGCGCGCCTCGCT TCAGTGAGCGAGCGCGCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG $\verb| CCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAATGGCGATTCCGTTGCAATGGCTG| \\$ $\tt GCGGTAATATTGTTCTGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTC$ AGGCAAGTGATGTTATTACTAATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTG ATGGACAGACTCTTTTACTCGGTGGCCTCACTGATTATAAAAACACTTCTCAGGATT $\tt CTGGCGTACCGTTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTC$ TGATTCTAACGAGGAAAGCACGTTATACGTGCTCGTCAAAGCAACCATAGTACGCG CCCTGTAGCGGCGCATTAAGCGCGGGGGGGTGTGGTGGTTACGCGCAGCGTGACCGC TACACTTGCCAGCGCCCTAGCGCCCGCTCCTTTCGCTTTCTTCCCTTTCTCGCC ACGTTCGCCGGCTTTCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGA TTTAGTGCTTTACGGCACCTCGACCCCAAAAAACTTGATTAGGGTGATGGTTCACGT AGTGGGCCATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCT TTAATAGTGGACTCTTGTTCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTC TTTTGATTTATAAGGGATTTTGCCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATT TAACAAAATTTAACGCGAATTTTAACAAAATATTAACGTTTACAATTTAAATATTT

GCTTATACAATCTTCCTGTTTTTTGGGGCTTTTCTGATTATCAACCGGGGTACATATGA $\tt TTGACATGCTAGTTTTACGATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTC$ ${\tt AGGCAATGACCTGATAGCCTTTGTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGC}$ ${\tt ATGAATTTATCAGCTAGAACGGTTGAATATCATATTGATGGTGATTTGACTGTCTCC}$ $\tt GGCCTTTCTCACCCGTTTGAATCTTTACCTACACATTACTCAGGCATTGCATTTAAAA$ TATATGAGGGTTCTAAAAATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAA ${\tt AAGTATTACAGGGTCATAATGTTTTTGGTACAACCGATTTAGCTTTATGCTCTGAGG}$ CTTTATTGCTTAATTTTGCTAATTCTTTGCCTTGCCTGTATGATTTATTGGATGTTGGA ATTCCTGATGCGGTATTTTCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGG CCAACACCCGCTGACGCCCTGACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGA CAAGCTGTGACCGTCTCCGGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCACCG AAACGCGCGAGACGAAAGGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATG ATAATAATGGTTTCTTAGACGTCAGGTGGCACTTTTCGGGGAAATGTGCGCGGAACC CCTATTTGTTTATTTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAAC CCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCC GTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTGCCTTCCTGTTTTTGCTCACCCAGAA ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACAT $\tt CGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTT$ GCCGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGAG TACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATG $\tt CGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTC$ ACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACT ${\tt AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGG}$ AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGC CCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAACGA AATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGTAACTGTCAGAC CAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAAATTTAAAAGGA TCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTC GTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTT TTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAAACCACCGCTACCAGCGGTGGT TTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAGCAG ${\tt AGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAA}$ GAACTCTGTAGCACCGCCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGC $\tt TGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACCGGA$

GAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACG $\tt CTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAG$ ${\tt GAGAGCGCACGAGGGAGCTTCCAGGGGGAAACGCCTGGTATCTTTATAGTCCTGTC}$ $\tt GGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGGGCGG$ AGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGG $\tt CCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTAC$ CGCCTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGCGAGT CAGTGAGCGAGGAAGCGGCAAGAGCGCCCAATACGCAAACCGCCTCTCCCCGCGCGT CCGCCATGCTACTTATCTACGTAGCCATGCTCTAGGACATTGATTATTGACTAGTGG ${\tt AGTTCCGCGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACC}$ CCCGCCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTT TCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATC ${\tt AAGTGTATCATATGCCAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCG}$ $\tt CCTGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTACTTGGCAGTACATCTA$ $\tt CGTATTAGTCATCGCTATTACCATGGTCGAGGTGAGCCCCACGTTCTGCTTCACTCTC$ GCGAGGGGCGGGGCGAGGCGAGAGGTGCGGCGGCAGCCAATCAGAGCGG $\tt CGCGCTCCGAAAGTTTCCTTTTATGGCGAGGCGGCGGCGGCGGCGGCCCTATAAAA$ GGCGAGCGCTGCCACGTCAGACGAAGGGCGCAGCGAGCGTCCTGATCCTTCCGCCC $\tt GGACGCTCAGGACAGCGGCCCGCTGCTCATAAGACTCGGCCTTAGAACCCCAGTAT$ CAGCAGAAGGACATTTTAGGACGGGACTTGGGTGACTCTAGGGCACTGGTTTTCTTT $\tt CCAGAGAGCGGAACAGGCGAGGAAAAGTAGTCCCTTCTCGGCGATTCTGCGGAGGG$ ATCTCCGTGGGGCGGTGAACGCCGATGATGCCTCTACTAACCATGTTCATGTTTTCTT TTTTTTTCTACAGGTCCTGGGTGACGAACAGGGTACCGCCACCATGGCCACCGGCTC TCGCACAAGCCTGCTGCTTTCGGACTGCTGTGCCTTGGCTCCAGGAGGG CTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGGAGGCTTGCTGAAGGCTGTAT CCGAAGATCTCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCT CTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGACAGATCTTCGATGTCACCATGCG $\tt TTTTGGCCTCTGACTGACGCATGGTGACCGAAGATCTGTCAGGACACAAGGCCTGTT$

 ${\tt ACTAGCACTCACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTAT}$

3 '

SEQ ID NO. 11 = SEQ ID NO. 1 + SEQ ID NO. 3 5'

AATCAACCTCTGGATTACAAAATTTGTGAAAGATTGACTGGTATTCTTAACTATGTT $\tt GCTCCTTTTACGCTATGTGGATACGCTGCTTTAATGCCTTTGTATCATGCTATTGCTT$ $\tt CCCGTATGGCTTTCATTTTCTCCTCCTTGTATAAATCCTGGTTGCTGTCTCTTTATGAG$ GAGTTGTGGCCCGTTGTCAGGCAACGTGGCGTGTGTGCACTGTGTTTGCTGACGCA ${\tt ACCCCCACTGGTTGGGGCATTGCCACCACCTGTCAGCTCCTTTCCGGGACTTTCGCTT}$ CAGGGGCTCGGCTGTTGGGCACTGACAATTCCGTGGTGTTGTCGGGGAAATCATCGT CCTTTCCTTGGCTGCTCGCCTGTGTTGCCACCTGGATTCTGCGCGGGACGTCCTTCTG CTGCGGCCTCTTCCGCGTCTTCGCCCTTCAGACGAGTCGGATCTCCCTTTTGGG CCGCCTCCCCGCCTAAGCTTATCGATACCGTCGAGATCTAACTTGTTTATTGCAGCTT ATAATGGTTACAAATAAAGCAATAGCATCACAAATTTCACAAATAAAGCATTTTTTT ${\tt CACTGCATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTTATCATGTCTGGAT}$ $\tt CTCGACCTCGACTAGAGCATGGCTACGTAGATAAGTAGCATGGCGGGTTAATCATTA$ ${\tt TCAGTGAGCGAGCGCGCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG}$ CCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAATGGCGATTCCGTTGCAATGGCTG $\tt GCGGTAATATTGTTCTGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTC$ AGGCAAGTGATGTTATTACTAATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTG ATGGACAGACTCTTTTACTCGGTGGCCTCACTGATTATAAAAACACTTCTCAGGATT $\tt CTGGCGTACCGTTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTC$ $\tt TGATTCTAACGAGGAAAGCACGTTATACGTGCTCGTCAAAGCAACCATAGTACGCG$ $\tt CCCTGTAGCGGCGCATTAAGCGCGGGGGGGTGTGGTGGTTACGCGCAGCGTGACCGC$ ${\tt TACACTTGCCAGCGCCCTAGCGCCCGCTCCTTTCGCTTTCTCCTTTCCTTTCTCGCC}$ ACGTTCGCCGGCTTTCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGA $\tt TTTAGTGCTTTACGGCACCTCGACCCCAAAAAACTTGATTAGGGTGATGGTTCACGT$ AGTGGGCCATCGCCCTGATAGACGGTTTTTTCGCCCCTTTGACGTTGGAGTCCACGTTCT TTAATAGTGGACTCTTGTTCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTC TTTTGATTTATAAGGGATTTTGCCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATT TAACAAAATTTAACGCGAATTTTAACAAAATATTAACGTTTACAATTTAAATATTT GCTTATACAATCTTCCTGTTTTTTGGGGCTTTTCTGATTATCAACCGGGGTACATATGA TTGACATGCTAGTTTTACGATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTC

continued AGGCAATGACCTGATAGCCTTTGTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGC ${\tt ATGAATTTATCAGCTAGAACGGTTGAATATCATATTGATGGTGATTTGACTGTCTCC}$ GGCCTTTCTCACCCGTTTGAATCTTTACCTACACATTACTCAGGCATTGCATTTAAAA TATATGAGGGTTCTAAAAATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAA AAGTATTACAGGGTCATAATGTTTTTGGTACAACCGATTTAGCTTTATGCTCTGAGG CTTTATTGCTTAATTTTGCTAATTCTTTGCCTTGCCTGTATGATTTATTGGATGTTGGA ATTCCTGATGCGGTATTTTCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGG CCAACACCCGCTGACGCCCTGACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGA CAAGCTGTGACCGTCTCCGGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCACCG AAACGCGCGAGACGAAAGGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATG ATAATAATGGTTTCTTAGACGTCAGGTGGCACTTTTCGGGGGAAATGTGCGCGGAACC CCTATTTGTTTATTTTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAACCCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCC GTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTGCCTTCCTGTTTTTTGCTCACCCAGAA ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACAT CGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTT ${\tt TACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATG}$ $\tt CGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTC$ GCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGAC ACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACT ${\tt AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGG}$ ${\tt AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGC}$ CCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAACGA ${\tt AATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGTAACTGTCAGAC}$ CAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAAATTTAAAAGGA TCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTC GTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTT TTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAACCACCGCTACCAGCGGTGGT TTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTTCCGAAGGTAACTGGCTTCAGCAG AGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAA GAACTCTGTAGCACCGCCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGC TGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACCGGA TAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGGAGC

GAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACG

continued $\tt CTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAG$ ${\tt GAGAGCGCACGAGGGAGCTTCCAGGGGGAAACGCCTGGTATCTTTATAGTCCTGTC}$ GGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGGCGG ${\tt AGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGG}$ CGCCTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGCGAGT CAGTGAGCGAGGAAGCGGAAGAGCGCCCAATACGCAAACCGCCTCTCCCCGCGCGT TGGCCGATTCATTAATGCAGCAGCTGCGCGCTCGCTCACTGAGGCCGCCCGGG CGCAGAGAGGGAGTGGCCAACTCCATCACTAGGGGTTCCTTGTAGTTAATGATTAAC CCGCCATGCTACTTATCTACGTAGCCATGCTCTAGGACATTGATTATTGACTAGTGG AGTTCCGCGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACC CCCGCCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTT TCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATC AAGTGTATCATATGCCAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCG CCTGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTACTTGGCAGTACATCTA CGTATTAGTCATCGCTATTACCATGGTCGAGGTGAGCCCCACGTTCTGCTTCACTCTC GGCGAGCGCTGCCACGTCAGACGAAGGGCGCAGCGAGCGTCCTGATCCTTCCGCCC $\tt GGACGCTCAGGACAGCGGCCCGCTGCTCATAAGACTCGGCCTTAGAACCCCAGTAT$ ${\tt CAGCAGAAGGACATTTTAGGACGGGACTTGGGTGACTCTAGGGCACTGGTTTTCTTT}$ CCAGAGAGCGGAACAGGCGAGGAAAAGTAGTCCCTTCTCGGCGATTCTGCGGAGGG ATCTCCGTGGGGCGGTGAACGCCGATGATGCCTCTACTAACCATGTTCATGTTTTCTT TTTTTTTCTACAGGTCCTGGGTGACGACAGGGTACCGCCACCATGGCCACCGCTC CTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGGAGGCTTGCTGAAGGCTGTAT TGCCATAACTCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCT CTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGTGATCATACGGATGCAATGCACCG $\tt TTTTGGCCTCTGACTGACGGTGCATTGCCCGTATGATCACAGGACACAAGGCCTGTT$ ACTAGCACTCACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTAT

- continued
AGCTATAACTCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCT
CTCTAGAAT

3 '

SEQ ID NO. 12 = SEQ ID NO. 1 + SEQ ID NO. 4 5'

 ${\tt AATCAACCTCTGGATTACAAAATTTGTGAAAGATTGACTGGTATTCTTAACTATGTT}$ GCTCCTTTTACGCTATGTGGATACGCTGCTTTAATGCCTTTGTATCATGCTATTGCTT $\tt CCCGTATGGCTTTCATTTTCTCCTCCTTGTATAAATCCTGGTTGCTGTCTCTTTATGAG$ GAGTTGTGGCCCGTTGTCAGGCAACGTGGCGTGGTGTGCACTGTGTTTGCTGACGCA ${\tt ACCCCCACTGGTTGGGGCATTGCCACCACCTGTCAGCTCCTTTCCGGGACTTTCGCTT}$ CAGGGGCTCGGCTGTTGGGCACTGACAATTCCGTGGTGTTGTCGGGGAAATCATCGT CCTTTCCTTGGCTGCTCGCCTGTGTTGCCACCTGGATTCTGCGCGGGACGTCCTTCTG CTGCGGCCTCTTCCGCGTCTTCGCCCTCAGACGAGTCGGATCTCCCTTTGGG CCGCCTCCCCGCCTAAGCTTATCGATACCGTCGAGATCTAACTTGTTTATTGCAGCTT ATAATGGTTACAAATAAAGCAATAGCATCACAAATTTCACAAATAAAGCATTTTTTT CACTGCATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTTATCATGTCTGGAT CTCGACCTCGACTAGAGCATGGCTACGTAGATAAGTAGCATGGCGGGTTAATCATTA TCAGTGAGCGAGCGCGCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG CCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAATGGCGATTCCGTTGCAATGGCTG $\tt GCGGTAATATTGTTCTGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTC$ $\tt AGGCAAGTGATGTTATTACTAATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTG$ ${\tt ATGGACAGACTCTTTTACTCGGTGGCCTCACTGATTATAAAAACACTTCTCAGGATT}$ $\tt CTGGCGTACCGTTCCTGTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTC$ $\tt TGATTCTAACGAGGAAAGCACGTTATACGTGCTCGTCAAAGCAACCATAGTACGCG$ $\tt CCCTGTAGCGGCGCATTAAGCGCGGGGGGGTGTGGTTACGCGCAGCGTGACCGC$ ${\tt TACACTTGCCAGCGCCCTAGCGCCCGCTCCTTTCGCTTTCTCCTTTCCTTTCTCGCC}$ ACGTTCGCCGGCTTTCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGA TTTAGTGCTTTACGGCACCTCGACCCCAAAAAACTTGATTAGGGTGATGGTTCACGT AGTGGGCCATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCT TTAATAGTGGACTCTTGTTCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTC TTTTGATTTATAAGGGATTTTGCCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATT TAACAAAATTTAACGCGAATTTTAACAAAATATTAACGTTTACAATTTAAATATTT GCTTATACAATCTTCCTGTTTTTTGGGGCTTTTTCTGATTATCAACCGGGGTACATATGATTGACATGCTAGTTTTACGATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTC AGGCAATGACCTGATAGCCTTTGTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGC

 $\tt ATGAATTTATCAGCTAGAACGGTTGAATATCATATTGATGGTGATTTGACTGTCTCC$

GGCCTTTCTCACCCGTTTGAATCTTTACCTACACATTACTCAGGCATTGCATTTAAAA TATATGAGGGTTCTAAAAATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAA ${\tt AAGTATTACAGGGTCATAATGTTTTTGGTACAACCGATTTAGCTTTATGCTCTGAGG}$ CTTTATTGCTTAATTTTGCTAATTCTTTGCCTTGCCTGTATGATTTATTGGATGTTGGA ATTCCTGATGCGGTATTTTCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGG $\tt CCAACACCCGCTGACGGGCCTGACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGA$ CAAGCTGTGACCGTCTCCGGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCACCG AAACGCGCGAGACGAAAGGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATG ATAATAATGGTTTCTTAGACGTCAGGTGGCACTTTTCGGGGGAAATGTGCGCGGAACC CCTATTTGTTTATTTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAAC CCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCC GTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTGCCTTCCTGTTTTTTGCTCACCCAGAA ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACAT $\tt CGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTT$ TCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGAC ${\tt TACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATG}$ $\tt CGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTC$ GCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGAC ACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACT ${\tt AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGTTTATTGCTGATAAATCTGG}$ ${\tt AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGC}$ $\tt CCTCCCGTATCGTAGTTATCTACACGACGGGGGGGTCAGGCAACTATGGATGAACGA$ ${\tt AATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGTAACTGTCAGAC}$ CAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAATTTAAAAGGA TCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTC GTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTT TTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAAACCACCGCTACCAGCGGTGGT TTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAGCAG AGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAA GAACTCTGTAGCACCGCCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGC TGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACCGGA TAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGGAGC ${\tt GAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACG}$ CTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAG

GGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGGGCGG ${\tt AGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGG}$ $\tt CCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTAC$ $\tt CGCCTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGCGAGT$ ${\tt CAGTGAGCGAGGAAGCGGCAAGCGCCCAATACGCAAACCGCCTCTCCCCGCGCGTT}$ $\tt TGGCCGATTCATTAATGCAGCAGCTGCGCGCTCGCTCACTGAGGCCGCCCGGG$ $\tt CGCAGAGAGGGGAGTGGCCAACTCCATCACTAGGGGTTCCTTGTAGTTAATGATTAAC$ CCGCCATGCTACTTATCTACGTAGCCATGCTCTAGGACATTGATTATTGACTAGTGG AGTTCCGCGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACC CCCGCCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTT TCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATC AAGTGTATCATATGCCAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCG CCTGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTACTTGGCAGTACATCTA CGTATTAGTCATCGCTATTACCATGGTCGAGGTGAGCCCCACGTTCTGCTTCACTCTC $\tt GGACGCTCAGGACAGCGGCCCGCTGCTCATAAGACTCGGCCTTAGAACCCCAGTAT$ $\tt CAGCAGAAGGACATTTTAGGACGGGACTTGGGTGACTCTAGGGCACTGGTTTTCTTT$ $\tt CCAGAGAGCGGAACAGGCGAGGGAAAAGTAGTCCCTTCTCGGCGATTCTGCGGAGGG$ ATCTCCGTGGGGCGGTGAACGCCGATGATGCCTCTACTAACCATGTTCATGTTTTCTT $\tt TTTTTTCTACAGGTCCTGGGTGACGAACAGGGTACCGCCACCATGGCCACCGGCTC$ TCGCACAAGCCTGCTGCTTCCGGACTGCTGTGCCTGCCTTGGCTCCAGGAGGG CTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGGAGGCTTGCTGAAGGCTGTAT CCCGAACTATCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCT CTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGAGATGAATGCCTGATAGCCTTCCG TTTTGGCCTCTGACTGACGGAAGGCTATGGCATTCATCTCAGGACACAAGGCCTGTT ACTAGCACTCACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTAT $\tt GCATTCATCTCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTC$ TCTAGAAT

SEQ ID NO. 13 = SEQ ID NO. 1 + SEQ ID NO' 5 5'

 ${\tt AATCAACCTCTGGATTACAAAATTTGTGAAAGATTGACTGGTATTCTTAACTATGTT}$ ${\tt GCTCCTTTTACGCTATGTGGATACGCTGCTTTAATGCCTTTGTATCATGCTATTGCTT}$ $\tt CCCGTATGGCTTTCATTTTCTCCTCCTTGTATAAATCCTGGTTGCTGTCTCTTTATGAG$ GAGTTGTGGCCCGTTGTCAGGCAACGTGGCGTGGTGTGCACTGTGTTTGCTGACGCA ${\tt ACCCCCACTGGTTGGGGCATTGCCACCACCTGTCAGCTCCTTTCCGGGACTTTCGCTT}$ CAGGGGCTCGGCTGTTGGGCACTGACAATTCCGTGGTGTTTGTCGGGGAAATCATCGT CTGCGGCCTCTTCCGCGTCTTCGCCCTTCAGACGAGTCGGATCTCCCTTTTGGGCCGCCTCCCCCCCTAAGCTTATCGATACCGTCGAGATCTAACTTGTTTATTGCAGCTTATAATGGTTACAAATAAAGCAATAGCATCACAAATTTCACAAATAAAGCATTTTTTT CACTGCATTCTAGTTGTGGTTTGTCCAAACTCATCATGTATCTTATCATGTCTGGAT CTCGACCTCGACTAGAGCATGGCTACGTAGATAAGTAGCATGGCGGGTTAATCATTA ${\tt TCAGTGAGCGAGCGGCGCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG}$ $\tt CCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAATGGCGATTCCGTTGCAATGGCTG$ $\tt GCGGTAATATTGTTCTGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTC$ AGGCAAGTGATGTTATTACTAATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTG ATGGACAGACTCTTTTACTCGGTGGCCTCACTGATTATAAAAACACTTCTCAGGATT $\tt CTGGCGTACCGTTCCTGTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTC$ TGATTCTAACGAGGAAAGCACGTTATACGTGCTCGTCAAAGCAACCATAGTACGCG ACGTTCGCCGGCTTTCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGA $\tt TTTAGTGCTTTACGGCACCTCGACCCCAAAAAACTTGATTAGGGTGATGGTTCACGT$ ${\tt AGTGGGCCATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCT}$ TTAATAGTGGACTCTTGTTCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTC TTTTGATTTATAAGGGATTTTGCCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATT TAACAAAATTTAACGCGAATTTTAACAAAATATTAACGTTTACAATTTAAATATTT GCTTATACAATCTTCCTGTTTTTTGGGGCTTTTCTGATTATCAACCGGGGTACATATGA TTGACATGCTAGTTTTACGATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTC AGGCAATGACCTGATAGCCTTTGTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGC ATGAATTTATCAGCTAGAACGGTTGAATATCATATTGATGGTGATTTGACTGTCTCC GGCCTTTCTCACCCGTTTGAATCTTTACCTACACATTACTCAGGCATTGCATTTAAAA TATATGAGGGTTCTAAAAATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAA

continued AAGTATTACAGGGTCATAATGTTTTTGGTACAACCGATTTAGCTTTATGCTCTGAGG ATTCCTGATGCGGTATTTTCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGG $\tt CCAACACCCGCTGACGCCCCTGACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGA$ ${\tt CAAGCTGTGACCGTCTCCGGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCACCG}$ AAACGCGCGAGACGAAAGGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATG ATAATAATGGTTTCTTAGACGTCAGGTGGCACTTTTCGGGGAAATGTGCGCGGAACC CCTATTTGTTTATTTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAAC CCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCC GTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTGCCTTCCTGTTTTTTGCTCACCCAGAA ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACAT CGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTTCGCCCCGAAGAACGTTT TCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGAC GCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGAG TACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATG $\tt CGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTC$ $\tt GCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGAC$ ${\tt AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGG}$ AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGC CCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAACGA AATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGTAACTGTCAGAC CAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAATTTAAAAGGA ${\tt TCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTC}$ GTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTT $\tt TTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAAAACCACCGCTACCAGCGGTGGT$ TTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAGCAG AGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAA GAACTCTGTAGCACCGCCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGC TGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACCGGA TAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGGAGC GAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACG CTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAG GAGAGCGCACGAGGGAGCTTCCAGGGGGAAACGCCTGGTATCTTTATAGTCCTGTC AGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGG

continued $\tt CCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTAC$ $\tt CGCCTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGCGAGT$ CAGTGAGCGAGGAAGCGGCAAGAGCGCCCAATACGCAAACCGCCTCTCCCCGCGCGT $\tt TGGCCGATTCATTAATGCAGCAGCTGCGCGCTCGCTCACTGAGGCCGCCCGGG$ CGCAGAGAGGGGAGTGGCCAACTCCATCACTAGGGGTTCCTTGTAGTTAATGATTAAC $\tt CCGCCATGCTACTTATCTACGTAGCCATGCTCTAGGACATTGATTATTGACTAGTGG$ AGTTCCGCGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACC $\tt CCCGCCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTT$ TCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATC AAGTGTATCATATGCCAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCG $\tt CCTGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTACTTGGCAGTACATCTA$ $\tt CGTATTAGTCATCGCTATTACCATGGTCGAGGTGAGCCCCACGTTCTGCTTCACTCTC$ TGCAGCGATGGGGGGGGGGGGGGGGGGGGGGGCGCGCCAGGCGGGGCGGGCGGG CGCGCTCCGAAAGTTTCCTTTTATGGCGAGGCGGCGGCGGCGGCGGCCCTATAAAA $\tt GGCGAGCGTGCCACGTCAGACGAAGGGCGCAGCGAGCGTCCTGATCCTTCCGCCC$ GGACGCTCAGGACAGCGGCCCGCTGCTCATAAGACTCGGCCTTAGAACCCCAGTAT ${\tt CAGCAGAAGGACATTTTAGGACGGGACTTGGGTGACTCTAGGGCACTGGTTTTCTTT}$ CCAGAGAGCGGAACAGGCGAGGAAAAGTAGTCCCTTCTCGGCGATTCTGCGGAGGG ATCTCCGTGGGGCGGTGAACGCCGATGATGCCTCTACTAACCATGTTCATGTTTTCTT TTTTTTTCTACAGGTCCTGGGTGACGAACAGGGTACCGCCACCATGGCCACCGGCTC TCGCACAAGCCTGCTGCTTCCGGACTGCTGTGCCTGCCTTGGCTCCAGGAGGG $\tt CTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGAAGGCTTGCTGAAGGCTGTAT$ ${\tt ATGATCTGAACAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCT}$ $\tt CTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGAATAATCAGATGGTGTTGCGATCG$ $\tt TTTTGGCCTCTGACTGACGATCGCAACACCCTGATTATTCAGGACACAAGGCCTGTT$ ACTAGCACTCACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTAT $\tt TGCATCAGTACAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCT$ CTCTAGAAT

-continued SEQ ID NO. 14 = SEQ ID NO. 1 + SEQ ID NO. 6

AATCAACCTCTGGATTACAAAATTTGTGAAAGATTGACTGGTATTCTTAACTATGTT GCTCCTTTTACGCTATGTGGATACGCTGCTTTAATGCCTTTGTATCATGCTATTGCTT $\tt CCCGTATGGCTTTCATTTTCTCCTCCTTGTATAAATCCTGGTTGCTGTCTCTTTATGAG$ ${\tt GAGTTGTGGCCCGTTGTCAGGCAACGTGGCGTGTGTGCACTGTTTTGCTGACGCA}$ ${\tt ACCCCACTGGTTGGGGCATTGCCACCACCTGTCAGCTCCTTTCCGGGACTTTCGCTT}$ CAGGGGCTCGGCTGTTGGGCACTGACAATTCCGTGGTGTTGTCGGGGAAATCATCGT $\tt CCTTTCCTTGGCTGCTCGCCTGTGTTGCCACCTGGATTCTGCGCGGGACGTCCTTCTG$ CTGCGGCCTCTTCCGCGTCTTCGCCTTCGCGAGCGAGTCGGATCTCCCTTTGGG CCGCCTCCCCGCCTAAGCTTATCGATACCGTCGAGATCTAACTTGTTTATTGCAGCTT ATAATGGTTACAAATAAAGCAATAGCATCACAAATTTCACAAATAAAGCATTTTTTT CACTGCATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTTATCATGTCTGGAT CTCGACCTCGACTAGAGCATGGCTACGTAGATAAGTAGCATGGGGGTTAATCATTA TCAGTGAGCGAGCGCGCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG $\tt CCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAATGGCGATTCCGTTGCAATGGCTG$ $\tt GCGGTAATATTGTTCTGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTC$ AGGCAAGTGATGTTATTACTAATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTG ATGGACAGACTCTTTTACTCGGTGGCCTCACTGATTATAAAAACACTTCTCAGGATT CTGGCGTACCGTTCCTGTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTC CCCTGTAGCGGCGCATTAAGCGCGGGGGGTGTGGTGGTTACGCGCAGCGTGACCGC ${\tt TACACTTGCCAGCGCCCTAGCGCCCGCTCCTTTCGCTTTCTCCTTTCCTTTCTCGCC}$ ACGTTCGCCGGCTTTCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGA $\tt TTTAGTGCTTTACGGCACCTCGACCCCAAAAAACTTGATTAGGGTGATGGTTCACGT$ AGTGGGCCATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCT TTAATAGTGGACTCTTGTTCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTC TTTTGATTTATAAGGGATTTTGCCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATT TAACAAAATTTAACGCGAATTTTAACAAAATATTAACGTTTACAATTTAAATATTT GCTTATACAATCTTCCTGTTTTTTGGGGCTTTTCTGATTATCAACCGGGGTACATATGA TTGACATGCTAGTTTTACGATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTC AGGCAATGACCTGATAGCCTTTGTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGC ATGA ATTTATCAGCTAGAACGGTTGAATATCATATTGATGGTGATTTGACTGTCTCC GGCCTTTCTCACCCGTTTGAATCTTTACCTACACATTACTCAGGCATTGCATTTAAAA TATATGAGGGTTCTAAAAATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAA ${\tt AAGTATTACAGGGTCATAATGTTTTTGGTACAACCGATTTAGCTTTATGCTCTGAGG}$

CTTTATTGCTTAATTTTGCTAATTCTTTGCCTTGCCTGTATGATTTATTGGATGTTGGA ${\tt ATTCCTGATGCGGTATTTTCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGG}$ $\tt CCAACACCCGCTGACGCGCCTTGACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGA$ ${\tt CAAGCTGTGACCGTCTCCGGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCACCG}$ AAACGCGCGAGACGAAAGGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATG ATAATAATGGTTTCTTAGACGTCAGGTGGCACTTTTCGGGGAAATGTGCGCGGAACC CCTATTTGTTTATTTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAAC CCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCC GTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTTGCCCTTCCTGTTTTTTGCTCACCCAGAA ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACAT CGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTT TCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGAC GCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGAG TACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATG $\tt CGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTC$ $\tt GCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGAC$ ${\tt ACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACT}$ ${\tt AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGG}$ ${\tt AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGC}$ CCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAACGA ${\tt CAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAATTTAAAAGGA}$ ${\tt TCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTC}$ $\tt GTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTT$ $\tt TTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAAAACCACCGCTACCAGCGGTGGT$ TTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAGCAG AGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAA GAACTCTGTAGCACCGCCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGC TGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACCGGA TAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGGAGC GAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACG CTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAG GAGAGCGCACGAGGGAGCTTCCAGGGGGAAACGCCTGGTATCTTTATAGTCCTGTC GGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGGGCGG AGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGG $\tt CCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTAC$

CGCCTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGCGAGT ${\tt CAGTGAGCGAGGAAGCGGCAAGCGCCCAATACGCAAACCGCCTCTCCCCGCGCGTT}$ $\tt CGCAGAGAGGGAGTGGCCAACTCCATCACTAGGGGTTCCTTGTAGTTAATGATTAAC$ CCGCCATGCTACTTATCTACGTAGCCATGCTCTAGGACATTGATTATTGACTAGTGG AGTTCCGCGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACC $\tt CCCGCCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTT$ ${\tt TCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATC}$ AAGTGTATCATATGCCAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCG CCTGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTACTTGGCAGTACATCTA $\tt CGTATTAGTCATCGCTATTACCATGGTCGAGGTGAGCCCCACGTTCTGCTTCACTCACTCAC$ $\tt CGCGCTCCGAAAGTTTCCTTTTATGGCGAGGCGGCGGCGGCGGCGGCCCTATAAAA$ GGCGAGCGCTGCCACGTCAGACGAAGGGCGCAGCGAGCGTCCTGATCCTTCCGCCC $\tt GGACGCTCAGGACAGCGGCCCGCTGCTCATAAGACTCGGCCTTAGAACCCCAGTAT$ CAGCAGAAGGACATTTTAGGACGGGACTTGGGTGACTCTAGGGCACTGGTTTTCTTT CCAGAGAGCGGAACAGGCGAGGAAAAGTAGTCCCTTCTCGGCGATTCTGCGGAGGG ATCTCCGTGGGGCGGTGAACGCCGATGATGCCTCTACTAACCATGTTCATGTTTTCTT TTTTTTTCTACAGGTCCTGGGTGACGAACAGGGTACCATGGCCACCGGCTCTCGCAC ${\tt AAGCCTGCTGGCTTTCGGACTGCTGTGCCTTGGCTCCAGGAGGGCTCCGC}$ $\tt GTAATCGGCTGATGCGTTTTGGCCTCTGACTGACGCAAACGCATGCCGATT$ ACCTCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTCTAGCCT GGAGGCTTGCTGAAGGCTGTATGCTGTTATACACGGTATGCCTTTCAGCGTTTTGGC $\tt CTCTGACTGACGCTGAAAGGCACCGTGTATAACAGGACACAAGGCCTGTTACTAGC$ ACTCACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGAT TGATCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTCTCTAGA

-continued SEQ ID NO. 15 = SEQ ID NO. 1 + SEQ ID NO. 7

AATCAACCTCTGGATTACAAAATTTGTGAAAGATTGACTGGTATTCTTAACTATGTT $\tt GCTCCTTTTACGCTATGTGGATACGCTGCTTTAATGCCTTTGTATCATGCTATTGCTT$ $\tt CCCGTATGGCTTTCATTTTCTCCTCCTTGTATAAATCCTGGTTGCTGTCTCTTTATGAG$ ${\tt GAGTTGTGGCCCGTTGTCAGGCAACGTGGCGTGTGTGCACTGTTTTGCTGACGCA}$ ${\tt ACCCCACTGGTTGGGGCATTGCCACCACCTGTCAGCTCCTTTCCGGGACTTTCGCTT}$ CAGGGGCTCGGCTGTTGGGCACTGACAATTCCGTGGTGTTGTCGGGGAAATCATCGT $\tt CCTTTCCTTGGCTGCTCGCCTGTGTTGCCACCTGGATTCTGCGCGGGACGTCCTTCTG$ CTGCGGCCTCTTCCGCGTCTTCGCCTTCGCGAGCGAGTCGGATCTCCCTTTGGG CCGCCTCCCCGCCTAAGCTTATCGATACCGTCGAGATCTAACTTGTTTATTGCAGCTT ATAATGGTTACAAATAAAGCAATAGCATCACAAATTTCACAAATAAAGCATTTTTTT CACTGCATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTTATCATGTCTGGAT CTCGACCTCGACTAGAGCATGGCTACGTAGATAAGTAGCATGGCGGGTTAATCATTA TCAGTGAGCGAGCGCGCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG $\tt CCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAATGGCGATTCCGTTGCAATGGCTG$ $\tt GCGGTAATATTGTTCTGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTC$ AGGCAAGTGATGTTATTACTAATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTG ATGGACAGACTCTTTTACTCGGTGGCCTCACTGATTATAAAAACACTTCTCAGGATT CTGGCGTACCGTTCCTGTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTC CCCTGTAGCGGCGCATTAAGCGCGGGGGGTGTGGTGGTTACGCGCAGCGTGACCGC ${\tt TACACTTGCCAGCGCCCTAGCGCCCGCTCCTTTCGCTTTCTCCTTTCCTTTCTCGCC}$ ACGTTCGCCGGCTTTCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGA $\tt TTTAGTGCTTTACGGCACCTCGACCCCAAAAAACTTGATTAGGGTGATGGTTCACGT$ AGTGGGCCATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCT TTAATAGTGGACTCTTGTTCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTC TTTTGATTTATAAGGGATTTTGCCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATT TAACAAAATTTAACGCGAATTTTAACAAAATATTAACGTTTACAATTTAAATATTT GCTTATACAATCTTCCTGTTTTTTGGGGCTTTTCTGATTATCAACCGGGGTACATATGA TTGACATGCTAGTTTTACGATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTC AGGCAATGACCTGATAGCCTTTGTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGC ATGA ATTTATCAGCTAGAACGGTTGAATATCATATTGATGGTGATTTGACTGTCTCC GGCCTTTCTCACCCGTTTGAATCTTTACCTACACATTACTCAGGCATTGCATTTAAAA TATATGAGGGTTCTAAAAATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAA ${\tt AAGTATTACAGGGTCATAATGTTTTTGGTACAACCGATTTAGCTTTATGCTCTGAGG}$

CTTTATTGCTTAATTTTGCTAATTCTTTGCCTTGCCTGTATGATTTATTGGATGTTGGA ${\tt ATTCCTGATGCGGTATTTTCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGG}$ $\tt CCAACACCCGCTGACGCGCCTTGACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGA$ ${\tt CAAGCTGTGACCGTCTCCGGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCACCG}$ AAACGCGCGAGACGAAAGGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATG ATAATAATGGTTTCTTAGACGTCAGGTGGCACTTTTCGGGGAAATGTGCGCGGAACC CCTATTTGTTTATTTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAAC CCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCC GTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTTGCCCTTCCTGTTTTTTGCTCACCCAGAA ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACAT CGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTT TCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGAC GCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGAG TACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATG $\tt CGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTC$ $\tt GCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGAC$ ${\tt ACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACT}$ ${\tt AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGG}$ ${\tt AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGC}$ CCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAACGA ${\tt CAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAATTTAAAAGGA}$ ${\tt TCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTC}$ $\tt GTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTT$ $\tt TTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAAAACCACCGCTACCAGCGGTGGT$ TTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAGCAG AGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAA GAACTCTGTAGCACCGCCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGC TGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACCGGA TAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGGAGC GAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACG CTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAG GAGAGCGCACGAGGGAGCTTCCAGGGGGAAACGCCTGGTATCTTTATAGTCCTGTC GGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGGGCGG AGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGG $\tt CCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTAC$

CGCCTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGCGAGT ${\tt CAGTGAGCGAGGAAGCGGCAAGCGCCCAATACGCAAACCGCCTCTCCCCGCGCGTT}$ $\tt CGCAGAGAGGGAGTGGCCAACTCCATCACTAGGGGTTCCTTGTAGTTAATGATTAAC$ CCGCCATGCTACTTATCTACGTAGCCATGCTCTAGGACATTGATTATTGACTAGTGG AGTTCCGCGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACC $\tt CCCGCCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTT$ ${\tt TCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATC}$ AAGTGTATCATATGCCAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCG CCTGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTACTTGGCAGTACATCTA $\tt CGTATTAGTCATCGCTATTACCATGGTCGAGGTGAGCCCCACGTTCTGCTTCACTCACTCAC$ $\tt CGCGCTCCGAAAGTTTCCTTTTATGGCGAGGCGGCGGCGGCGGCGGCGCCCTATAAAA$ GGCGAGCGCTGCCACGTCAGACGAAGGGCGCAGCGAGCGTCCTGATCCTTCCGCCC $\tt GGACGCTCAGGACAGCGGCCCGCTGCTCATAAGACTCGGCCTTAGAACCCCAGTAT$ CAGCAGAAGGACATTTTAGGACGGGACTTGGGTGACTCTAGGGCACTGGTTTTCTTT CCAGAGAGCGGAACAGGCGAGGAAAAGTAGTCCCTTCTCGGCGATTCTGCGGAGGG ATCTCCGTGGGGCGGTGAACGCCGATGATGCCTCTACTAACCATGTTCATGTTTTCTT TTTTTTTCTACAGGTCCTGGGTGACGAACAGGGTACCGCCACCATGGCCACCGGCTC $\tt CTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGAGGCTTGCTGAAGGCTGTAT$ ${\tt AGCTTTATGTCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTC}$ TAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGATACAGCAGATATCGCGAATTTCGT $\tt TTTGGCCTCTGACTGACGAAATTCGCGATCTGCTGTATCAGGACACAAGGCCTGTTA$ CTAGCACTCACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTATG AACTGAAACAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTCT CTAGAAT

-continued SEQ ID NO. 16 = SEQ ID NO. 1 + SEQ ID NO. 8

AATCAACCTCTGGATTACAAAATTTGTGAAAGATTGACTGGTATTCTTAACTATGTT $\tt GCTCCTTTTACGCTATGTGGATACGCTGCTTTAATGCCTTTGTATCATGCTATTGCTT$ $\tt CCCGTATGGCTTTCATTTTCTCCTCCTTGTATAAATCCTGGTTGCTGTCTCTTTATGAG$ ${\tt GAGTTGTGGCCCGTTGTCAGGCAACGTGGCGTGTGTGCACTGTTTTGCTGACGCA}$ ${\tt ACCCCACTGGTTGGGGCATTGCCACCACCTGTCAGCTCCTTTCCGGGACTTTCGCTT}$ CAGGGGCTCGGCTGTTGGGCACTGACAATTCCGTGGTGTTGTCGGGGAAATCATCGT $\tt CCTTTCCTTGGCTGCTCGCCTGTGTTGCCACCTGGATTCTGCGCGGGACGTCCTTCTG$ CTGCGGCCTCTTCCGCGTCTTCGCCTTCGCGAGCGAGTCGGATCTCCCTTTGGG CCGCCTCCCCGCCTAAGCTTATCGATACCGTCGAGATCTAACTTGTTTATTGCAGCTT ATAATGGTTACAAATAAAGCAATAGCATCACAAATTTCACAAATAAAGCATTTTTTT CACTGCATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTTATCATGTCTGGAT CTCGACCTCGACTAGAGCATGGCTACGTAGATAAGTAGCATGGCGGGTTAATCATTA TCAGTGAGCGAGCGCGCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG $\tt CCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAATGGCGATTCCGTTGCAATGGCTG$ $\tt GCGGTAATATTGTTCTGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTC$ AGGCAAGTGATGTTATTACTAATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTG ATGGACAGACTCTTTTACTCGGTGGCCTCACTGATTATAAAAACACTTCTCAGGATT CTGGCGTACCGTTCCTGTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTC CCCTGTAGCGGCGCATTAAGCGCGGGGGGTGTGGTGGTTACGCGCAGCGTGACCGC ${\tt TACACTTGCCAGCGCCCTAGCGCCCGCTCCTTTCGCTTTCTCCTTTCCTTTCTCGCC}$ ACGTTCGCCGGCTTTCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGA $\tt TTTAGTGCTTTACGGCACCTCGACCCCAAAAAACTTGATTAGGGTGATGGTTCACGT$ AGTGGGCCATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCT TTAATAGTGGACTCTTGTTCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTC TTTTGATTTATAAGGGATTTTGCCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATT TAACAAAATTTAACGCGAATTTTAACAAAATATTAACGTTTACAATTTAAATATTT GCTTATACAATCTTCCTGTTTTTTGGGGCTTTTCTGATTATCAACCGGGGTACATATGA TTGACATGCTAGTTTTACGATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTC AGGCAATGACCTGATAGCCTTTGTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGC ATGA ATTTATCAGCTAGAACGGTTGAATATCATATTGATGGTGATTTGACTGTCTCC GGCCTTTCTCACCCGTTTGAATCTTTACCTACACATTACTCAGGCATTGCATTTAAAA TATATGAGGGTTCTAAAAATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAA ${\tt AAGTATTACAGGGTCATAATGTTTTTGGTACAACCGATTTAGCTTTATGCTCTGAGG}$

CTTTATTGCTTAATTTTGCTAATTCTTTGCCTTGCCTGTATGATTTATTGGATGTTGGA ATTCCTGATGCGGTATTTTCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGG $\tt CCAACACCCGCTGACGCGCCTTGACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGA$ ${\tt CAAGCTGTGACCGTCTCCGGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCACCG}$ AAACGCGCGAGACGAAAGGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATG ATAATAATGGTTTCTTAGACGTCAGGTGGCACTTTTCGGGGAAATGTGCGCGGAACC CCTATTTGTTTATTTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAAC CCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCC GTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTTGCCCTTCCTGTTTTTTGCTCACCCAGAA ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACAT CGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTT TCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGAC GCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGAG TACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATG $\tt CGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTC$ $\tt GCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGAC$ ${\tt ACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACT}$ ${\tt AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGG}$ ${\tt AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGC}$ CCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAACGA ${\tt CAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAATTTAAAAGGA}$ ${\tt TCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTC}$ $\tt GTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTT$ $\tt TTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAAAACCACCGCTACCAGCGGTGGT$ TTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAGCAG AGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAA GAACTCTGTAGCACCGCCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGC TGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACCGGA TAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGGAGC GAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACG CTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAG GAGAGCGCACGAGGGAGCTTCCAGGGGGAAACGCCTGGTATCTTTATAGTCCTGTC GGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGGGCGG AGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGG $\tt CCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTAC$

CGCCTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGCGAGT ${\tt CAGTGAGCGAGGAAGCGGCAAGCGCCCAATACGCAAACCGCCTCTCCCCGCGCGTT}$ $\tt CGCAGAGAGGGAGTGGCCAACTCCATCACTAGGGGTTCCTTGTAGTTAATGATTAAC$ CCGCCATGCTACTTATCTACGTAGCCATGCTCTAGGACATTGATTATTGACTAGTGG AGTTCCGCGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACC $\tt CCCGCCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTT$ ${\tt TCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATC}$ AAGTGTATCATATGCCAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCG CCTGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTACTTGGCAGTACATCTA $\tt CGTATTAGTCATCGCTATTACCATGGTCGAGGTGAGCCCCACGTTCTGCTTCACTCACTCAC$ $\tt CGCGCTCCGAAAGTTTCCTTTTATGGCGAGGCGGCGGCGGCGGCGGCGCCCTATAAAA$ GGCGAGCGCTGCCACGTCAGACGAAGGGCGCAGCGAGCGTCCTGATCCTTCCGCCC $\tt GGACGCTCAGGACAGCGGCCCGCTGCTCATAAGACTCGGCCTTAGAACCCCAGTAT$ CAGCAGAAGGACATTTTAGGACGGGACTTGGGTGACTCTAGGGCACTGGTTTTCTTT CCAGAGAGCGGAACAGGCGAGGAAAAGTAGTCCCTTCTCGGCGATTCTGCGGAGGG ATCTCCGTGGGGCGGTGAACGCCGATGATGCCTCTACTAACCATGTTCATGTTTTCTT TTTTTTTCTACAGGTCCTGGGTGACGAACAGGGTACCGCCACCATGGCCACCGGCTC CTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGGAGGCTTGCTGAAGGCTGTAT ${\tt ACGCCTGTATCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCT}$ CTAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGTCATGATCAATATGGTATCCGGCG $\tt TTTTGGCCTCTGACTGACGCCGGATACCATTGATCATGACAGGACACAAGGCCTGTT$ ACTAGCACTCACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTAT $\tt TTGATCATGACAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTC$ TCTAGAAT

-continued SEQ ID NO. 17 = SEQ ID NO. 1 + SEQ ID NO. 9

AATCAACCTCTGGATTACAAAATTTGTGAAAGATTGACTGGTATTCTTAACTATGTT GCTCCTTTTACGCTATGTGGATACGCTGCTTTAATGCCTTTGTATCATGCTATTGCTT $\tt CCCGTATGGCTTTCATTTTCTCCTCCTTGTATAAATCCTGGTTGCTGTCTCTTTATGAG$ ${\tt GAGTTGTGGCCCGTTGTCAGGCAACGTGGCGTGTGTGCACTGTTTTGCTGACGCA}$ ${\tt ACCCCACTGGTTGGGGCATTGCCACCACCTGTCAGCTCCTTTCCGGGACTTTCGCTT}$ CAGGGGCTCGGCTGTTGGGCACTGACAATTCCGTGGTGTTGTCGGGGAAATCATCGT $\tt CCTTTCCTTGGCTGCTCGCCTGTGTTGCCACCTGGATTCTGCGCGGGACGTCCTTCTG$ CTGCGGCCTCTTCCGCGTCTTCGCCTTCGCGAGCGAGTCGGATCTCCCTTTGGG CCGCCTCCCCGCCTAAGCTTATCGATACCGTCGAGATCTAACTTGTTTATTGCAGCTT ATAATGGTTACAAATAAAGCAATAGCATCACAAATTTCACAAATAAAGCATTTTTTT CACTGCATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTTATCATGTCTGGAT CTCGACCTCGACTAGAGCATGGCTACGTAGATAAGTAGCATGGCGGGTTAATCATTA TCAGTGAGCGAGCGCGCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG $\tt CCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAATGGCGATTCCGTTGCAATGGCTG$ $\tt GCGGTAATATTGTTCTGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTC$ AGGCAAGTGATGTTATTACTAATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTG ATGGACAGACTCTTTTACTCGGTGGCCTCACTGATTATAAAAACACTTCTCAGGATT CTGGCGTACCGTTCCTGTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTC TGATTCTAACGAGGAAAGCACGTTATACGTGCTCGTCAAAGCAACCATAGTACGCG CCCTGTAGCGGCGCATTAAGCGCGGGGGGTGTGGTGGTTACGCGCAGCGTGACCGC ${\tt TACACTTGCCAGCGCCCTAGCGCCCGCTCCTTTCGCTTTCTCCTTTCCTTTCTCGCC}$ ACGTTCGCCGGCTTTCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGA $\tt TTTAGTGCTTTACGGCACCTCGACCCCAAAAAACTTGATTAGGGTGATGGTTCACGT$ AGTGGGCCATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCT TTAATAGTGGACTCTTGTTCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTC TTTTGATTTATAAGGGATTTTGCCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATT TAACAAAATTTAACGCGAATTTTAACAAAATATTAACGTTTACAATTTAAATATTT GCTTATACAATCTTCCTGTTTTTGGGGCTTTTCTGATTATCAACCGGGGTACATATGA TTGACATGCTAGTTTTACGATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTC AGGCAATGACCTGATAGCCTTTGTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGC ATGA ATTTATCAGCTAGAACGGTTGAATATCATATTGATGGTGATTTGACTGTCTCC GGCCTTTCTCACCCGTTTGAATCTTTACCTACACATTACTCAGGCATTGCATTTAAAA TATATGAGGGTTCTAAAAATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAA

 ${\tt AAGTATTACAGGGTCATAATGTTTTTGGTACAACCGATTTAGCTTTATGCTCTGAGG}$

CTTTATTGCTTAATTTTGCTAATTCTTTGCCTTGCCTGTATGATTTATTGGATGTTGGA ${\tt ATTCCTGATGCGGTATTTTCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGG}$ $\tt CCAACACCCGCTGACGCGCCTTGACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGA$ ${\tt CAAGCTGTGACCGTCTCCGGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCACCG}$ AAACGCGCGAGACGAAAGGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATG ATAATAATGGTTTCTTAGACGTCAGGTGGCACTTTTCGGGGAAATGTGCGCGGAACC CCTATTTGTTTATTTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAAC CCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCC GTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTTGCCCTTCCTGTTTTTTGCTCACCCAGAA ACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACAT CGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTTT TCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGAC GCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGAG TACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATG $\tt CGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTC$ $\tt GCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGAC$ ${\tt ACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACT}$ ${\tt AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGG}$ ${\tt AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGC}$ CCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAACGA ${\tt CAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAATTTAAAAGGA}$ ${\tt TCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTC}$ $\tt GTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTT$ $\tt TTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAAAACCACCGCTACCAGCGGTGGT$ TTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAGCAG AGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAA GAACTCTGTAGCACCGCCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGC TGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACCGGA TAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGGAGC GAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACG CTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAG GAGAGCGCACGAGGGAGCTTCCAGGGGGAAACGCCTGGTATCTTTATAGTCCTGTC GGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGGGCGG AGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGG $\tt CCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTAC$

CGCCTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGCGAGT ${\tt CAGTGAGCGAGGAAGCGGCAAGCGCCCAATACGCAAACCGCCTCTCCCCGCGCGTT}$ $\tt CGCAGAGAGGGAGTGGCCAACTCCATCACTAGGGGTTCCTTGTAGTTAATGATTAAC$ CCGCCATGCTACTTATCTACGTAGCCATGCTCTAGGACATTGATTATTGACTAGTGG AGTTCCGCGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACC $\tt CCCGCCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTT$ ${\tt TCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATC}$ AAGTGTATCATATGCCAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCG CCTGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTACTTGGCAGTACATCTA $\tt CGTATTAGTCATCGCTATTACCATGGTCGAGGTGAGCCCCACGTTCTGCTTCACTCACTCAC$ $\tt CGCGCTCCGAAAGTTTCCTTTTATGGCGAGGCGGCGGCGGCGGCGGCGCCCTATAAAA$ GGCGAGCGCTGCCACGTCAGACGAAGGGCGCAGCGAGCGTCCTGATCCTTCCGCCC $\tt GGACGCTCAGGACAGCGGCCCGCTGCTCATAAGACTCGGCCTTAGAACCCCAGTAT$ CAGCAGAAGGACATTTTAGGACGGGACTTGGGTGACTCTAGGGCACTGGTTTTCTTT CCAGAGAGCGGAACAGGCGAGGAAAAGTAGTCCCTTCTCGGCGATTCTGCGGAGGG ATCTCCGTGGGGCGGTGAACGCCGATGATGCCTCTACTAACCATGTTCATGTTTTCTT $\tt TTTTTTCTACAGGTCCTGGGTGACGAACAGGGTACCGCCACCATGGCCACCGGCTC$ $\tt CTCCGCCGCTAGCATCGATACCGTCGCTATGTGCTGGAGGCTTGCTGAAGGCTGTAT$ $\tt GCCTGTTTATCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTC$ TAGCCTGGAGGCTTGCTGAAGGCTGTATGCTGTCATAGCTGCAATGGTTTCTTCCGTT $\tt TTGGCCTCTGACTGACGGAAGAACCTGCAGCTATGACAGGACACAAGGCCTGTTA$ $\tt CTAGCACTCACATGGAACAAATGGCCTCTAGCCTGGAGGCTTGCTGAAGGCTGTATG$ GGAAGTGGTCAGGACACAAGGCCTGTTACTAGCACTCACATGGAACAAATGGCCTC TCTAGAAT

[0062] As will be appreciated by those skilled in the art, because the recombinant plasmid is a circular vector, the one or more sequences of the miRNA expression cassettes may be connected at the 3' end of SEQ ID NO. 1, as shown in SEQ ID NO. 10, SEQ ID NO. 11, SEQ ID NO. 12, SEQ ID NO. 13, SEQ ID NO. 14, SEQ ID NO. 15, SEQ ID NO. 16 and SEQ ID NO. 17, or at the 5' end of SEQ ID NO. 1.

[0063] As will be appreciated by those skilled in the art, a perfect match of nucleotides with each of the miRNA expression cassette sequences is not necessary in order to have the desired result of decreased bioavailability of the target biomolecule as a result of the target cell producing the miRNA sequence that will bind to and degrade the mRNA of the target biomolecule. In some embodiments of the present disclosure, about 80% to about 100% nucleotide sequence matching with each of the miRNA expression cassettes causes the desired result. In some embodiments of the present disclosure, about 85% to about 100% nucleotide sequence matching with each of the miRNA expression cassettes causes the desired result. In some embodiments of the present disclosure, about 90% to about 100% nucleotide sequence matching with each of the miRNA expression cassettes causes the desired result. In some embodiments of the present disclosure, about 95% to about 100% nucleotide sequence matching with each of the miRNA expression cassettes causes the desired result.

Sequence total quantity: 17

Example 1—Expression Cassette

[0064] Expression cassettes for expressing miRNA were synthesized. The synthesized miRNA expression cassettes were cloned into the pAVA-00200 plasmid backbone containing the CASI promoter, multiple cloning site (MCS), Woodchuck Hepatitis Virus post-transcriptional regulatory element (WPRE), and Simian virus 40 (SV40) polyadenylation (polyA) sequence, all flanked by the AAV2 inverted terminal repeats (ITR). pAVA-00200 was cut with the restriction enzymes KpnI and XbaI in the MCS and separated on a 1% agarose gel. The band of interest was excised and purified using a gel extraction kit. Each miRNA expression cassette was amplified by polymerase chain reaction (PCR) using Taq polymerase and the PCR products were gel purified and the bands on interest were also excised and purified using a gel extraction kit. These PCR products contained the miRNA expression cassettes in addition to 15 base pair 5' and 3' overhangs that aligned with the ends of the linearized pAVA-00200 backbone. Using in-fusion cloning, the amplified miRNA expression cassettes were integrated with the pAVA-00200 backbone via homologous recombination. The resulting RP contained the following: 5' ITR, CASI promoter, miRNA expression cassette, WPRE, SV40 polyA and ITR 3'.

SEQUENCE LISTING

```
SEQ ID NO: 1
                       moltype = DNA length = 5799
FEATURE
                       Location/Qualifiers
                       1..5799
source
                       mol type = other DNA
                       organism = synthetic construct
SEQUENCE: 1
aatcaacctc tggattacaa aatttgtgaa agattgactg gtattcttaa ctatgttgct
ccttttacgc tatgtggata cgctgcttta atgcctttgt atcatgctat tgcttcccgt
atggetttea titteteete ettgtataaa teetggttge tgtetettta tgaggagttg
                                                                   180
tggcccgttg tcaggcaacg tggcgtggtg tgcactgtgt ttgctgacgc aacccccact
ggttggggca ttgccaccac ctgtcagctc ctttccggga ctttcgcttt ccccctccct
                                                                    300
attgccacgg cggaactcat cgccgcctgc cttgcccgct gctggacagg ggctcggctg
ttgggcactg acaattccgt ggtgttgtcg gggaaatcat cgtcctttcc ttggctgctc
                                                                    420
geotgtgttg ccacctggat tetgegeggg aegteettet getaegteee tteggeeete
aatccagegg accttectte eegeggeetg etgeeggete tgeggeetet teegegtett
                                                                    540
egecttegee eteagaegag teggatetee etttgggeeg eeteeegge taagettate
                                                                    600
gataccqtcq agatctaact tqtttattqc agcttataat qqttacaaat aaaqcaataq
                                                                    660
catcacaaat ttcacaaata aaqcattttt ttcactqcat tctaqttqtq qtttqtccaa
                                                                    720
                                                                    780
acticational quatitatic atquirtiggat ctcqacctcq actaqaqcat qqctacqtaq
ataagtagca tggcgggtta atcattaact acaaggaacc cctagtgatg gagttggcca
                                                                    840
ctccctctct qcqcqctcqc tcqctcactq aqqccqqqcq accaaaqqtc
                                                                    900
                                                       acccaacacc
                                                                    960
egggetttge eegggegge teagtgageg agegagegeg cagetggegt aatagegaag
aggeograe eqategoeet teccaacaqt tgegcageet gaatggegaa tggcgattee
                                                                    1020
                                                                    1080
gttgcaatgg ctggcggtaa tattgttctg gatattacca gcaaggccga tagtttgagt
tottotacto aggoaagtga tgttattact aatcaaagaa gtattgcgac aacggttaat
                                                                    1140
ttgcgtgatg gacagactct tttactcggt ggcctcactg attataaaaa cacttctcag
                                                                    1200
gattetggeg taccettect gtetaaaate cetttaateg geeteetgtt tageteecge
                                                                    1260
totgattota acgaggaaag cacgttatac gtgctcgtca aagcaaccat agtacgcgcc
                                                                   1320
ctgtagcggc gcattaagcg cggcgggtgt ggtggttacg cgcagcgtga ccgctacact
                                                                    1380
tgccagcgcc ctagcgcccg ctcctttcgc tttcttccct tcctttctcq ccacgttcgc
                                                                    1440
eggettteee egteaagete taaategggg geteeettta gggtteegat ttagtgettt
                                                                    1500
acggcacctc gaccccaaaa aacttgatta gggtgatggt tcacgtagtg ggccatcgcc
                                                                    1560
ctgatagacg gtttttcgcc ctttgacgtt ggagtccacg ttctttaata gtggactctt
                                                                    1620
gttccaaact ggaacaacac tcaaccctat ctcggtctat tcttttgatt tataagggat
                                                                   1680
tttgccgatt tcggcctatt ggttaaaaaa tgagctgatt taacaaaaat ttaacgcgaa
                                                                   1740
ttttaacaaa atattaacgt ttacaattta aatatttgct tatacaatct tcctgttttt
                                                                   1800
ggggcttttc tgattatcaa ccggggtaca tatgattgac atgctagttt tacgattacc
                                                                   1860
gttcatcgat tetettgttt getecagaet eteaggeaat gaeetgatag eetttgtaga
                                                                   1920
gacctctcaa aaatagctac cctctccggc atgaatttat cagctagaac ggttgaatat
                                                                   1980
catattgatg gtgatttgac tgtctccggc ctttctcacc cgtttgaatc tttacctaca
cattactcag gcattgcatt taaaatatat gagggttcta aaaattttta tccttgcgtt
```

```
gaaataaagg cttctcccgc aaaagtatta cagggtcata atgtttttgg tacaaccgat
ttagetttat getetgagge tttattgett aattttgeta attetttgee ttgeetgtat
                                                                  2220
gatttattgg atgttggaat teetgatgeg gtattttete ettaegeate tgtgeggtat
                                                                  2280
ttcacaccgc atatggtgca ctctcagtac aatctgctct gatgccgcat agttaagcca
                                                                  2340
geocegacae eegecaacae eegetgaege geoctgaegg gettgtetge teeeggeate
                                                                  2400
cgcttacaga caagctgtga ccgtctccgg gagctgcatg tgtcagaggt tttcaccgtc
                                                                  2460
atcaccgaaa cgcgcgagac gaaagggcct cgtgatacgc ctatttttat aggttaatgt
                                                                  2520
catgataata atggtttctt agacgtcagg tggcactttt cggggaaatg tgcgcggaac
                                                                  2580
ccctatttgt ttatttttct aaatacattc aaatatgtat ccgctcatga gacaataacc
                                                                  2640
ctgataaatg cttcaataat attgaaaaag gaagagtatg agtattcaac atttccgtgt
egecettatt ceettttttg eggcattttg eetteetgtt tttgeteace eagaaaeget
                                                                  2760
ggtgaaagta aaagatgctg aagatcagtt gggtgcacga gtgggttaca tcgaactgga
                                                                  2820
tctcaacagc ggtaagatcc ttgagagttt tcgccccgaa gaacgttttc caatgatgag
                                                                  2880
cacttttaaa gttctgctat gtggcgcggt attatcccgt attgacgccg ggcaagagca
actoggtogo ogcatacact attotoagaa tgacttggtt gagtactcac cagtcacaga
aaagcatctt acggatggca tgacagtaag agaattatgc agtgctgcca taaccatgag
tgataacact gcggccaact tacttctgac aacgatcgga ggaccgaagg agctaaccgc
ttttttgcac aacatggggg atcatgtaac tcgccttgat cgttgggaac cggagctgaa
tgaagccata ccaaacgacg agcgtgacac cacgatgcct gtagcaatgg caacaacgtt
                                                                  3240
gegeaaacta ttaactggeg aactacttac tetagettee eggeaacaat taatagactg
                                                                  3300
gatggaggcg gataaagttg caggaccact tetgegeteg gecetteegg etggetggtt
                                                                  3360
tattgctgat aaatctggag ccggtgagcg tgggtctcgc ggtatcattg cagcactggg
                                                                  3420
gccagatggt aagccctccc gtatcgtagt tatctacacg acggggagtc aggcaactat
                                                                  3480
ggatgaacga aatagacaga tcgctgagat aggtgcctca ctgattaagc attggtaact
                                                                  3540
3600
aaggatctag gtgaagatcc tttttgataa tctcatgacc aaaatccctt aacgtgagtt
                                                                  3660
ttcgttccac tgagcgtcag accccgtaga aaagatcaaa ggatcttctt gagatccttt
                                                                  3720
ttttctgcgc gtaatctgct gcttgcaaac aaaaaaacca ccgctaccag cggtggtttg
                                                                  3780
tttgccgqat caagagctac caactctttt tccgaaggta actggcttca gcagagcgca
                                                                  3840
gataccaaat actgtccttc tagtgtagcc gtagttaggc caccacttca agaactctgt
                                                                  3900
ageacegect acataceteg etetgetaat cetgttacea gtggetgetg ceagtggega
                                                                  3960
taagtcgtgt cttaccgggt tggactcaag acgatagtta ccggataagg cgcagcggtc
                                                                  4020
gggctgaacg gggggttcgt gcacacagcc cagcttggag cgaacgacct acaccgaact
                                                                  4080
gagataccta cagcgtgagc tatgagaaag cgccacgctt cccgaaggga gaaaggcgga
                                                                  4140
caggtateeg gtaageggea gggteggaac aggagagege acgagggage ttecaggggg
                                                                  4200
aaacgcctgg tatctttata gtcctgtcgg gtttcgccac ctctgacttg agcgtcgatt
                                                                  4260
tttgtgatgc tcgtcagggg ggcggagcct atggaaaaac gccagcaacg cggccttttt
                                                                  4320
acggttcctg gccttttgct ggccttttgc tcacatgttc tttcctgcgt tatcccctga
                                                                  4380
ttctgtggat aaccgtatta ccgcctttga gtgagctgat accgctcgcc gcagccgaac
                                                                  4440
gaccgagcgc agcgagtcag tgagcgagga agcggaagag cgcccaatac gcaaaccgcc
                                                                  4500
teteceegeg egttggeega tteattaatg eageagetge gegetegete geteaetgag
                                                                  4560
geogeologig caaageologi gegtegggeg acetttggte geologicete agtgagegag
                                                                  4620
cgagcgcgca gagagggagt ggccaactcc atcactaggg gttccttgta gttaatgatt
                                                                  4680
aacccgccat gctacttatc tacgtagcca tgctctagga cattgattat tgactagtgg
                                                                  4740
agttccgcgt tacataactt acggtaaatg gcccgcctgg ctgaccgccc aacgaccccc
                                                                  4800
gcccattgac gtcaataatg acgtatgttc ccatagtaac gccaataggg actttccatt
                                                                  4860
gacgtcaatg ggtggagtat ttacggtaaa ctgcccactt ggcagtacat caagtgtatc
                                                                  4920
atatgccaag tacgcccct attgacgtca atgacggtaa atggcccgcc tggcattatg
                                                                  4980
cccagtacat gaccttatgg gactttccta cttggcagta catctacgta ttagtcatcg
                                                                  5040
ctattaccat ggtcgaggtg agcccacgt tctgcttcac tctccccatc tccccccct
                                                                  5100
cccacccc aattitqtat ttatttattt tttaattatt ttqtqcaqcq atqqqqqcqq
                                                                  5160
ggggggggg gggcgcgcc caggcggggc ggggcggggc gaggggcggg gcggggcgag
                                                                  5220
gcggagaggt gcggcggcag ccaatcagag cggcgcgctc cgaaagtttc cttttatggc
                                                                  5280
gaggcggcgg cggcggcggc cctataaaaa gcgaagcgcg cggcgggcgg gagtcgctgc
                                                                  5340
gegetgeett egeceegtge eeegeteege egeegeeteg egeegeeege eeeggetetg
actgaccgcg ttactaaaac aggtaagtcc ggcctccgcg ccgggttttg gcgcctcccg
                                                                  5460
cgggcgcccc cctcctcacg gcgagcgctg ccacgtcaga cgaagggcgc agcgagcgtc
                                                                  5520
ctgatcette egeceggaeg eteaggaeag eggeeegetg eteataagae teggeettag
aaccccagta tcagcagaag gacattttag gacgggactt gggtgactct agggcactgg
ttttctttcc agagagaga acaggagagg aaaagtagtc ccttctcggc gattctqcgg
agggatetee gtggggeggt gaacgeegat gatgeeteta etaaceatgt teatgtttte
ttttttttc tacaggtcct gggtgacgaa cagggtacc
SEO ID NO: 2
                      moltype = DNA length = 540
FEATURE
                      Location/Qualifiers
source
                      1..540
                      mol type = other DNA
                      organism = synthetic construct
SEOUENCE: 2
gccaccatgg ccaccggctc tcgcacaagc ctgctgctgg ctttcggact gctgtgcctg
cettggetee aggagggete egeegetage ategataceg tegetatgtg etggaggett
gctgaagget gtatgctgag atctteggtt gcaccatget egttttggee tetgactgae
gagcatggtg accgaagatc tcaggacaca aggcctgtta ctagcactca catggaacaa
atggcctcta gcctggaggc ttgctgaagg ctgtatgctg acagatcttc gatgtcacca
                                                                  300
tgcgttttgg cctctgactg acgcatggtg accgaagatc tgtcaggaca caaggcctgt
```

tactagcact cacatggaac aaatggcctc tagcctggag gcttgctgaa ggctgtatgc 420

```
tgacagatct tcgaagtcac catgcgtttt ggcctctgac tgacgcatgg tgaccgaaga
tetgteagga cacaaggeet gttactagea etcacatgga acaaatggee tetetagaat
SEQ ID NO: 3
                      moltype = DNA length = 540
FEATURE
                      Location/Qualifiers
                      1..540
source
                      mol_type = other DNA
                      organism = synthetic construct
SEQUENCE: 3
gccaccatgg ccaccggctc tcgcacaagc ctgctgctgg ctttcggact gctgtgcctg
cettggetee aggagggete egeegetage ategataceg tegetatgtg etggaggett
gctgaaggct gtatgctgag ttatggcact gagatgctgg cgttttggcc tctgactgac
gccagcatct gtgccataac tcaggacaca aggcctgtta ctagcactca catggaacaa
atggcctcta gcctggaggc ttgctgaagg ctgtatgctg tgatcatacg gatgcaatgc
accepttttgg cctctgactg acgeptgcatt gcccgtatga tcacaggaca caaggcctgt
tactagcact cacatggaac aaatggcctc tagcctggag gcttgctgaa ggctgtatgc
tgagttatag cttaatcatc ctggcgtttt ggcctctgac tgacgccagg atgaaagcta
taactcagga cacaaggeet gttactagea eteacatgga acaaatggee tetetagaat
SEO ID NO: 4
                      moltype = DNA length = 540
FEATURE
                      Location/Qualifiers
                      1..540
source
                      mol_type = other DNA
                      organism = synthetic construct
SEOUENCE: 4
gccaccatgg ccaccggctc tcgcacaagc ctgctgctgg ctttcggact gctgtgcctg
cettggetee aggagggete egeegetage ategataceg tegetatgtg etggaggett
                                                                  120
getgaagget gtatgetgat agtteggget gteagaattt egttttggee tetgaetgae
                                                                  180
gaaattotga goocgaacta toaggacaca aggootgtta otagcactoa catggaacaa
                                                                  240
atggcctcta gcctggaggc ttgctgaagg ctgtatgctg agatgaatgc ctgatagcct
teegttttgg cetetgactg aeggaagget atggeattea teteaggaca eaaggeetgt
                                                                  360
tactagcact cacatggaac aaatggcctc tagcctggag gcttgctgaa ggctgtatgc
                                                                  420
tgagatgaat gccaaatagc cttccgtttt ggcctctgac tgacggaagg ctatggcatt
                                                                  480
catctcaqqa cacaaqqcct qttactaqca ctcacatqqa acaaatqqcc tctctaqaat
SEO ID NO: 5
                      moltype = DNA length = 540
FEATURE
                      Location/Qualifiers
source
                      1..540
                      mol_type = other DNA
organism = synthetic construct
SEQUENCE: 5
gccaccatgg ccaccggctc tcgcacaagc ctgctgctgg ctttcggact gctgtgcctg
cettggetee aggagggete egeegetage ategataceg tegetatgtg etggaggett
                                                                  120
gctgaaggct gtatgctgtt cagatcatct gggtatccgg cgttttggcc tctgactgac
                                                                  180
gccggatacc gatgatctga acaggacaca aggcctgtta ctagcactca catggaacaa
                                                                  240
atggcctcta gcctggaggc ttgctgaagg ctgtatgctg aataatcaga tggtgttgcg
                                                                  300
atcgttttgg cctctgactg acgatcgcaa caccctgatt attcaggaca caaggcctgt
tactagcact cacatggaac aaatggcctc tagcctggag gcttgctgaa ggctgtatgc
                                                                  420
tgtactgatg cacaatttaa acgccgtttt ggcctctgac tgacggcgtt taaagtgcat
cagtacagga cacaaggeet gttactagea etcacatgga acaaatggee tetetagaat
SEQ ID NO: 6
                      moltype = DNA length = 534
FEATURE
                      Location/Qualifiers
                      1..534
source
                      mol_type = other DNA
                      organism = synthetic construct
SEQUENCE: 6
atggccaccg gctctcgcac aagcetgctg ctggctttcg gactgctgtg cctgccttgg
ctccaggagg gctccgccgc tagcatcgat accgtcgcta tgtgctggag gcttgctgaa
ggctgtatgc tgaggtaatc ggctgatgcg tttgcgtttt ggcctctgac tgacgcaaac
gcatgccgat tacctcagga cacaaggcct gttactagca ctcacatgga acaaatggcc
totageotgg aggettgetg aaggetgtat getgttatae aeggtatgee ttteagegtt
ttqqcctctq actqacqctq aaaqqcaccq tqtataacaq qacacaaqqc ctqttactaq
cactcacatg gaacaaatgg cctctagcct ggaggcttgc tgaaggctgt atgctgatca
                                                                  420
480
aggacacaag gcctgttact agcactcaca tggaacaaat ggcctctcta gaat
SEO ID NO: 7
                      moltype = DNA length = 540
FEATURE
                      Location/Qualifiers
source
                      1..540
                      mol_type = other DNA
                      organism = synthetic construct
SEOUENCE: 7
gccaccatgg ccaccggctc tcgcacaagc ctgctgctgg ctttcggact gctgtgcctg 60
cettggetee aggagggete egeegetage ategataceg tegetatgtg etggaggett
gctgaaggct gtatgctgac ataaagctat gtcctggcag cgttttggcc tctgactgac 180
```

```
gctgccagga tagctttatg tcaggacaca aggcctgtta ctagcactca catggaacaa
atggcctcta gcctggaggc ttgctgaagg ctgtatgctg atacagcaga tatcgcgaat
                                                                   300
ttcgttttgg cctctgactg acgaaattcg cgatctgctg tatcaggaca caaggcctgt
                                                                   360
                                                                   420
tactagcact cacatggaac aaatggcctc tagcctggag gcttgctgaa ggctgtatgc
tgtttcagtt taaaatcagc gccacgtttt ggcctctgac tgacgtggcg ctgattaaac
                                                                   480
tgaaacagga cacaaggcct gttactagca ctcacatgga acaaatggcc tctctagaat
SEO ID NO: 8
                       moltype = DNA length = 540
FEATURE
                       Location/Qualifiers
                       1..540
source
                       mol type = other DNA
                       organism = synthetic construct
SEQUENCE: 8
gccaccatgg ccaccggctc tcgcacaagc ctgctgctgg ctttcggact gctgtgcctg
cettggetee aggagggete egeegetage ategataceg tegetatgtg etggaggett
gctgaaggct gtatgctgat acaggcgttt gtgctcggtt cgttttggcc tctgactgac
gaaccgagca aacgcctgta tcaggacaca aggcctgtta ctagcactca catggaacaa
atggcctcta gcctggaggc ttgctgaagg ctgtatgctg tcatgatcaa tatggtatcc
ggcgttttgg cctctgactg acgccggata ccattgatca tgacaggaca caaggcctgt
tactaqcact cacatqqaac aaatqqcctc taqcctqqaq qcttqctqaa qqctqtatqc
tgtcatgatc aataaggtat ccggcgtttt ggcctctgac tgacgccgga taccattgat
                                                                   480
catgacagga cacaaggeet gttactagea etcacatgga acaaatggee tetetagaat
                       moltype = DNA length = 540
SEO ID NO: 9
FEATURE
                       Location/Qualifiers
                       1..540
source
                       mol type = other DNA
                       organism = synthetic construct
SEOUENCE: 9
gccaccatgg ccaccggctc tcgcacaagc ctgctgctgg ctttcggact gctgtgcctg
cettggetee aggagggete egeegetage ategataceg tegetatgtg etggaggett
                                                                   120
gctgaaggct gtatgctgat aaacaggctt gttcgcccag cgttttggcc tctgactgac
                                                                   180
gctgggcgaa agcctgttta tcaggacaca aggcctgtta ctagcactca catggaacaa
                                                                   240
atgqcctcta qcctqqaqqc ttqctqaaqq ctqtatqctq tcataqctqc aatqqtttct
                                                                   300
teegttttgg cetetgaetg aeggaagaaa eetgeageta tgaeaggaea eaaggeetgt
                                                                   360
tactagcact cacatggaac aaatggcctc tagcctggag gcttgctgaa ggctgtatgc
                                                                   420
tgaccacttc cacaattcat gcaccgtttt ggcctctgac tgacggtgca tgaagtggaa
                                                                   480
gtggtcagga cacaaggcct gttactagca ctcacatgga acaaatggcc tctctagaat
SEQ ID NO: 10
                       moltype = DNA length = 6339
FEATURE
                      Location/Qualifiers
source
                      1..6339
                       mol_type = other DNA
                       organism = synthetic construct
SECUENCE: 10
aatcaacctc tggattacaa aatttgtgaa agattgactg gtattcttaa ctatgttgct
ccttttacgc tatgtggata cgctgcttta atgcctttgt atcatgctat tgcttcccgt
                                                                   120
atggetttea tttteteete ettgtataaa teetggttge tgtetettta tgaggagttg
                                                                   180
tggcccgttg tcaggcaacg tggcgtggtg tgcactgtgt ttgctgacgc aacccccact
ggttggggca ttgccaccac ctgtcagctc ctttccggga ctttcgcttt ccccctccct
                                                                   300
attgccacgg cggaactcat cgccgcctgc cttgcccgct gctggacagg ggctcggctg
ttgggcactg acaattccgt ggtgttgtcg gggaaatcat cgtcctttcc ttggctgctc
gcctgtgttg ccacctggat tctgcgcggg acgtccttct gctacgtccc ttcggccctc
aatccagcgg accttectte eegeggeetg etgeeggete tgeggeetet teegegtett
                                                                   540
cgccttcgcc ctcagacgag tcggatctcc ctttgggccg cctccccgcc taagcttatc
gataccgtcg agatctaact tgtttattgc agcttataat ggttacaaat aaagcaatag
catcacaaat ttcacaaata aagcattttt ttcactgcat tctagttgtg gtttgtccaa
acticated gtatettate atgtetggat etegaceteg actagageat ggetaegtag
ataagtagca tggcgggtta atcattaact acaaggaacc cctagtgatg gagttggcca
cteectetet gegegetege tegeteactg aggeogggeg accaaaggte geoogaegee
cgggctttgc ccgggcggcc tcagtgagcg agcgagcgcg cagctggcgt aatagcgaag
aggeorgeac egategeet teccaacagt tgegeageet gaatggegaa tggegattee
qttqcaatqq ctqqcqqtaa tattqttctq qatattacca qcaaqqccqa taqtttqaqt
                                                                   1080
tettetaete aggeaagtga tgttattaet aateaaagaa gtattgegae aaeggttaat
                                                                   1140
ttgcgtgatg gacagactct tttactcggt ggcctcactg attataaaaa cacttctcag
                                                                   1200
gattetggcg taccgtteet gtetaaaate cetttaateg geeteetgtt tageteeege
totgattota acgaggaaag cacgttatac gtgctcgtca aagcaaccat agtacgcgcc
                                                                   1320
ctgtagcggc gcattaagcg cggcgggtgt ggtggttacg cgcagcgtga ccgctacact
                                                                   1380
tgccagcgcc ctagcgcccg ctcctttcgc tttcttccct tcctttctcg ccacgttcgc
                                                                   1440
cggctttccc cgtcaagctc taaatcgggg gctcccttta gggttccgat ttagtgcttt
                                                                   1500
acggcacctc gaccccaaaa aacttgatta gggtgatggt tcacgtagtg ggccatcgcc
ctgatagacg gtttttcgcc ctttgacgtt ggagtccacg ttctttaata gtggactctt
                                                                   1620
gttccaaact ggaacaacac tcaaccctat ctcggtctat tcttttgatt tataagggat
tttgccgatt tcggcctatt ggttaaaaaa tgagctgatt taacaaaaat ttaacgcgaa
                                                                   1740
ttttaacaaa atattaacqt ttacaattta aatatttqct tatacaatct tcctqttttt
ggggcttttc tgattatcaa ccggggtaca tatgattgac atgctagttt tacgattacc
```

| gttcatcgat | tctcttqttt | gctccagact | ctcaggcaat | gacctgatag | cctttqtaqa | 1920 |
|------------|------------|------------|------------|------------|------------|------|
| | | | | cagctagaac | | 1980 |
| | | | | cgtttgaatc | | 2040 |
| | | | | aaaatttta | | 2100 |
| gaaataaagg | cttctcccgc | aaaagtatta | cagggtcata | atgtttttgg | tacaaccgat | 2160 |
| | | | | attctttgcc | | 2220 |
| | | | | cttacgcatc | | 2280 |
| | | | | gatgccgcat | | 2340 |
| gccccgacac | ccgccaacac | ccgctgacgc | gccctgacgg | gcttgtctgc | teceggeate | 2400 |
| | | | | tgtcagaggt | | 2460 |
| | | | | ctatttttat | | 2520 |
| catgataata | atggtttctt | agacgtcagg | tggcactttt | cggggaaatg | tgcgcggaac | 2580 |
| | | | | ccgctcatga | | 2640 |
| ctgataaatg | cttcaataat | attgaaaaag | gaagagtatg | agtattcaac | atttccgtgt | 2700 |
| cgcccttatt | cccttttttg | cggcattttg | ccttcctgtt | tttgctcacc | cagaaacgct | 2760 |
| | | | | gtgggttaca | | 2820 |
| tctcaacagc | ggtaagatcc | ttgagagttt | tcgccccgaa | gaacgttttc | caatgatgag | 2880 |
| cacttttaaa | gttctgctat | gtggcgcggt | attatcccgt | attgacgccg | ggcaagagca | 2940 |
| actcggtcgc | cgcatacact | attctcagaa | tgacttggtt | gagtactcac | cagtcacaga | 3000 |
| | | | | agtgctgcca | | 3060 |
| tgataacact | gcggccaact | tacttctgac | aacgatcgga | ggaccgaagg | agctaaccgc | 3120 |
| ttttttgcac | aacatggggg | atcatgtaac | tcgccttgat | cgttgggaac | cggagctgaa | 3180 |
| | | | | gtagcaatgg | | 3240 |
| | | | | cggcaacaat | | 3300 |
| gatggaggcg | gataaagttg | caggaccact | tetgegeteg | gcccttccgg | ctggctggtt | 3360 |
| | | | | ggtatcattg | | 3420 |
| | | | | acggggagtc | | 3480 |
| ggatgaacga | aatagacaga | tcgctgagat | aggtgcctca | ctgattaagc | attggtaact | 3540 |
| gtcagaccaa | gtttactcat | atatacttta | gattgattta | aaacttcatt | tttaatttaa | 3600 |
| | | | | aaaatccctt | | 3660 |
| ttcgttccac | tgagcgtcag | accccgtaga | aaagatcaaa | ggatcttctt | gagatccttt | 3720 |
| ttttctgcgc | gtaatctgct | gcttgcaaac | aaaaaaacca | ccgctaccag | cggtggtttg | 3780 |
| tttgccggat | caagagctac | caactctttt | tccgaaggta | actggcttca | gcagagcgca | 3840 |
| gataccaaat | actgtccttc | tagtgtagcc | gtagttaggc | caccacttca | agaactctgt | 3900 |
| agcaccgcct | acatacctcg | ctctgctaat | cctgttacca | gtggctgctg | ccagtggcga | 3960 |
| taagtcgtgt | cttaccgggt | tggactcaag | acgatagtta | ccggataagg | cgcagcggtc | 4020 |
| gggctgaacg | gggggttcgt | gcacacagcc | cagcttggag | cgaacgacct | acaccgaact | 4080 |
| gagataccta | cagcgtgagc | tatgagaaag | cgccacgctt | cccgaaggga | gaaaggcgga | 4140 |
| caggtatccg | gtaagcggca | gggtcggaac | aggagagcgc | acgagggagc | ttccaggggg | 4200 |
| aaacgcctgg | tatctttata | gtcctgtcgg | gtttcgccac | ctctgacttg | agcgtcgatt | 4260 |
| tttgtgatgc | tcgtcagggg | ggcggagcct | atggaaaaac | gccagcaacg | cggccttttt | 4320 |
| acggttcctg | gccttttgct | ggccttttgc | tcacatgttc | tttcctgcgt | tatcccctga | 4380 |
| ttctgtggat | aaccgtatta | ccgcctttga | gtgagctgat | accgctcgcc | gcagccgaac | 4440 |
| gaccgagcgc | agcgagtcag | tgagcgagga | agcggaagag | cgcccaatac | gcaaaccgcc | 4500 |
| tctccccgcg | cgttggccga | ttcattaatg | cagcagctgc | gcgctcgctc | gctcactgag | 4560 |
| gccgcccggg | caaagcccgg | gcgtcgggcg | acctttggtc | gcccggcctc | agtgagcgag | 4620 |
| | | | | gttccttgta | | 4680 |
| | | | | cattgattat | | 4740 |
| agttccgcgt | tacataactt | acggtaaatg | gcccgcctgg | ctgaccgccc | aacgaccccc | 4800 |
| | | | | gccaataggg | | 4860 |
| | | | | ggcagtacat | | 4920 |
| | | | | atggcccgcc | | 4980 |
| | | | | catctacgta | | 5040 |
| | | | | tctccccatc | | 5100 |
| | _ | | | ttgtgcagcg | | 5160 |
| | | | | gaggggggg | | 5220 |
| | | | | cgaaagtttc | | 5280 |
| | | | | cggcgggcgg | | 5340 |
| | | | | cgccgcccgc | | 5400 |
| | | | | ccgggttttg | | 5460 |
| | _ | | | cgaagggcgc | | 5520 |
| | | | | ctcataagac | | 5580 |
| _ | | - | | gggtgactct | | 5640 |
| | | | | ccttctcggc | | 5700 |
| | | | | ctaaccatgt | | 5760 |
| tttttttc | tacaggtcct | gggtgacgaa | cagggtaccg | ccaccatggc | caccggctct | 5820 |
| cgcacaagcc | tgctgctggc | tttcggactg | ctgtgcctgc | cttggctcca | ggagggctcc | 5880 |
| gccgctagca | tcgataccgt | cgctatgtgc | tggaggcttg | ctgaaggctg | tatgctgaga | 5940 |
| | | | | agcatggtga | | 6000 |
| | | | | tggcctctag | | 6060 |
| | | | | gcgttttggc | | 6120 |
| | | | | actagcactc | | 6180 |
| | | | | gacagatett | | 6240 |
| | | | | ctgtcaggac | | 6300 |
| | | caaatggcct | | 5 45540 | | 6339 |
| | | Janusyyeet | Josepaac | | | 2229 |

moltype = DNA length = 6339 SEO ID NO: 11 FEATURE Location/Qualifiers source 1..6339 mol_type = other DNA organism = synthetic construct SEOUENCE: 11 aatcaacctc tggattacaa aatttgtgaa agattgactg gtattcttaa ctatgttgct cettttacgc tatgtggata cgctgcttta atgcctttgt atcatgctat tgcttcccgt atggetttea tttteteete ettgtataaa teetggttge tgtetettta tgaggagttg tggcccgttg tcaggcaacg tggcgtggtg tgcactgtgt ttgctgacgc aacccccact ggttggggca ttgccaccac ctgtcagctc ctttccggga ctttcgcttt ccccctccct 300 attgccacgg cggaactcat cgccgcctgc cttgcccgct gctggacagg ggctcggctg ttgggcactg acaattccgt ggtgttgtcg gggaaatcat cgtcctttcc ttggctgctc gcctgtgttg ccacctggat tctgcgcggg acgtccttct gctacgtccc ttcggccctc aatccagegg accttectte eegeggeetg etgeeggete tgeggeetet teegegtett cgccttcgcc ctcagacgag tcggatctcc ctttgggccg cctccccgcc taagcttatc gataccgtcg agatctaact tgtttattgc agcttataat ggttacaaat aaagcaatag catcacaaat ttcacaaata aagcattttt ttcactgcat tctagttgtg gtttgtccaa actcatcaat gtatcttatc atgtctggat ctcgacctcg actagagcat ggctacgtag ataagtagca tggcgggtta atcattaact acaaggaacc cctagtgatg gagttggcca ctccctctct gcgcgctcgc tcgctcactg aggccgggcg accaaaggtc gcccgacgcc 900 cgggctttgc ccgggcggcc tcagtgagcg agcgagcgcg cagctggcgt aatagcgaag 960 aggcccqcac cqatcqccct tcccaacaqt tqcqcaqcct qaatqqcqaa tqqcqattcc 1020 gttgcaatgg ctggcggtaa tattgttctg gatattacca gcaaggccga tagtttgagt 1080 tottotacto aggicalita tottattact altolaagaa gtattgogac aloggitaat 1140 ttgcgtgatg gacagactct tttactcggt ggcctcactg attataaaaa cacttctcag 1200 gattetggcg taccgtteet gtetaaaate cetttaateg geeteetgtt tageteeege 1260 tetgatteta acgaggaaag caegttatae gtgetegtea aagcaaceat agtaegegee 1320 ctgtagcggc gcattaagcg cggcgggtgt ggtggttacg cgcagcgtga ccgctacact 1380 tgccagcgcc ctagcgcccg ctcctttcgc tttcttccct tcctttctcg ccacgttcgc 1440 eggettteee egteaagete taaategggg geteeettta gggtteegat ttagtgettt 1500 1560 acggcacctc gaccccaaaa aacttgatta gggtgatggt tcacgtagtg ggccatcgcc ctgatagacg gtttttcgcc ctttgacgtt ggagtccacg ttctttaata gtggactctt 1620 qttccaaact qqaacaacac tcaaccctat ctcqqtctat tcttttqatt tataaqqqat 1680 tttgccgatt tcggcctatt ggttaaaaaa tgagctgatt taacaaaaat ttaacgcgaa 1740 ttttaacaaa atattaacqt ttacaattta aatatttqct tatacaatct tcctqttttt 1800 ggggcttttc tgattatcaa ccggggtaca tatgattgac atgctagttt tacgattacc 1860 gttcatcgat tctcttgttt gctccagact ctcaggcaat gacctgatag cctttgtaga 1920 gacctctcaa aaatagctac cctctccggc atgaatttat cagctagaac ggttgaatat 1980 catattgatg gtgatttgac tgtctccggc ctttctcacc cgtttgaatc tttacctaca 2040 cattactcag gcattgcatt taaaatatat gagggttcta aaaattttta tccttgcgtt 2100 gaaataaagg cttctcccgc aaaagtatta cagggtcata atgtttttgg tacaaccgat 2160 ttagctttat gctctgaggc tttattgctt aattttgcta attctttgcc ttgcctgtat 2220 gatttattgg atgttggaat tcctgatgcg gtattttctc cttacgcatc tgtgcggtat 2280 ttcacacege atatggtgca eteteagtae aatetgetet gatgeegeat agttaageea 2340 geologica cogecaacae cegetgaege geologiaegg gettgtetge teeleggeate 2400 cgcttacaga caagctgtga ccgtctccgg gagctgcatg tgtcagaggt tttcaccgtc 2460 atcaccgaaa cgcgcgagac gaaagggcct cgtgatacgc ctatttttat aggttaatgt 2520 catgataata atggtttctt agacgtcagg tggcactttt cggggaaatg tgcgcggaac 2580 ccctatttgt ttatttttct aaatacattc aaatatgtat ccgctcatga gacaataacc 2640 ctgataaatg cttcaataat attgaaaaag gaagagtatg agtattcaac atttccgtgt cgcccttatt cccttttttg cggcattttg ccttcctgtt tttgctcacc cagaaacgct 2760 ggtgaaagta aaagatgctg aagatcagtt gggtgcacga gtgggttaca tcgaactgga 2820 totcaacago ggtaagatoo ttgagagttt togocoogaa gaacgtttto caatgatgag 2880 cacttttaaa gttctgctat gtggcgcggt attatcccgt attgacgccg ggcaagagca actoggtogo ogcatacact attotoagaa tgacttggtt gagtactcac cagtcacaga aaagcatctt acggatggca tgacagtaag agaattatgc agtgctgcca taaccatgag tgataacact gcggccaact tacttctgac aacgatcgga ggaccgaagg agctaaccgc ttttttgcac aacatggggg atcatgtaac tcgccttgat cgttgggaac cggagctgaa tgaagccata ccaaacgacg agcgtgacac cacgatgcct gtagcaatgg caacaacgtt 3240 gcgcaaacta ttaactggcg aactacttac tctagcttcc cggcaacaat taatagactg gatqqaqqcq qataaaqttq caqqaccact tctqcqctcq qcccttccqq ctqqctqqtt 3360 tattgctgat aaatctggag ccggtgagcg tgggtctcgc ggtatcattg cagcactggg 3420 gccagatggt aagccttcc gtatcgtagt tatctacacg acggggagtc aggcaactat 3480 ggatgaacga aatagacaga tcgctgagat aggtgcctca ctgattaagc attggtaact 3540 aaggatctag gtgaagatcc tttttgataa tctcatgacc aaaatccctt aacgtgagtt 3660 ttcqttccac tqaqcqtcaq accccqtaqa aaaqatcaaa qqatcttctt qaqatccttt 3720 ttttctgcgc gtaatctgct gcttgcaaac aaaaaaacca ccgctaccag cggtggtttg 3780 tttgccggat caagagctac caactctttt tccgaaggta actggcttca gcagagcgca 3840 gataccaaat actgtccttc tagtgtagcc gtagttaggc caccacttca agaactctgt agcaccgcct acatacctcg ctctgctaat cctgttacca gtggctgctg ccagtggcga 3960 taagtcgtgt cttaccgggt tggactcaag acgatagtta ccggataagg cgcagcggtc 4020 gggctgaacg gggggttcgt gcacacagcc cagcttggag cgaacgacct acaccgaact 4080 gagataccta cagcgtgagc tatgagaaag cgccacgctt cccgaaggga gaaaggcgga 4140 caggtateeg gtaageggea gggteggaae aggagagege acgagggage ttecaggggg

```
aaacgcctgg tatctttata gtcctgtcgg gtttcgccac ctctgacttg agcgtcgatt
tttgtgatgc tcgtcagggg ggcggagcct atggaaaaac gccagcaacg cggccttttt
                                                                  4320
acggttcctg gccttttgct ggccttttgc tcacatgttc tttcctgcgt tatcccctga
                                                                  4380
ttctgtggat aaccgtatta ccgcctttga gtgagctgat accgctcgcc gcagccgaac
                                                                  4440
gaccgagcgc agcgagtcag tgagcgagga agcggaagag cgcccaatac gcaaaccgcc
                                                                  4500
teteceegeg egttggeega tteattaatg eageagetge gegetegete geteactgag
                                                                  4560
geogeologic caaageologic geotegges acctttggto geologicto agtgagegag
                                                                  4620
cgagcgcgca gagagggagt ggccaactcc atcactaggg gttccttgta gttaatgatt
                                                                  4680
aacccgccat gctacttatc tacgtagcca tgctctagga cattgattat tgactagtgg
                                                                  4740
agttccgcgt tacataactt acggtaaatg gcccgcctgg ctgaccgccc aacgaccccc
gcccattgac gtcaataatg acgtatgttc ccatagtaac gccaataggg actttccatt
                                                                  4860
gacgtcaatg ggtggagtat ttacggtaaa ctgcccactt ggcagtacat caagtgtatc
                                                                  4920
atatgccaag tacgcccct attgacgtca atgacggtaa atggcccgcc tggcattatg
                                                                  4980
cccagtacat gaccttatgg gactttccta cttggcagta catctacgta ttagtcatcg
ctattaccat ggtcgaggtg agccccacgt tctgcttcac tctccccatc tccccccct
ccccacccc aattttgtat ttatttattt tttaattatt ttgtgcagcg atggggggg
ggggggggg gggcgcgcgc caggcggggc ggggcggggc gaggggggg gcggggcgag
gcggagaggt gcggcggcag ccaatcagag cggcgcgctc cgaaagtttc cttttatggc
gaggeggegg eggeggegge cetataaaaa gegaagegeg eggegggegg gagtegetge
                                                                  5340
gegetgeett egeceegtge eeegeteege egecgeeteg egecgeeege eeeggetetg
                                                                  5400
actgaccgcg ttactaaaac aggtaagtcc ggcctccgcg ccgggttttg gcgcctcccg
                                                                  5460
cgggcgcccc cctcctcacg gcgagcgctg ccacgtcaga cgaagggcgc agcgagcgtc
                                                                  5520
ctgatccttc cgcccggacg ctcaggacag cggcccgctg ctcataagac tcggccttag
                                                                  5580
aaccccagta tcagcagaag gacattttag gacgggactt gggtgactct agggcactgg
                                                                  5640
ttttctttcc agagagcgga acaggcgagg aaaagtagtc ccttctcggc gattctgcgg
                                                                  5700
agggatctcc gtggggcggt gaacgccgat gatgcctcta ctaaccatgt tcatgttttc
                                                                  5760
ttttttttc tacaggtcct gggtgacgaa cagggtaccg ccaccatggc caccggctct
                                                                  5820
cgcacaagcc tgctgctggc tttcggactg ctgtgcctgc cttggctcca ggagggctcc
                                                                  5880
geogetagea tegatacegt egetatgtge tggaggettg etgaaggetg tatgetgagt
                                                                  5940
tatqqcactq aqatqctqqc qttttqqcct ctqactqacq ccaqcatctq tqccataact
                                                                  6000
caggacacaa ggcctgttac tagcactcac atggaacaaa tggcctctag cctggaggct
                                                                  6060
                                                                  6120
eggtgeattg ceegtatgat cacaggacae aaggeetgtt actageacte acatggaaca
                                                                  6180
aatqqcctct aqcctqqaqq cttqctqaaq qctqtatqct qaqttataqc ttaatcatcc
                                                                  6240
tggcgttttg gcctctgact gacgccagga tgaaagctat aactcaggac acaaggcctg
                                                                  6300
ttactagcac tcacatggaa caaatggcct ctctagaat
                                                                  6339
SEQ ID NO: 12
                      moltype = DNA length = 6339
FEATURE
                      Location/Qualifiers
source
                      1..6339
                      mol_type = other DNA
                      organism = synthetic construct
SEQUENCE: 12
aatcaacctc tggattacaa aatttgtgaa agattgactg gtattcttaa ctatgttgct
ccttttacgc tatgtggata cgctgcttta atgcctttgt atcatgctat tgcttcccgt
                                                                  120
atggetttea tttteteete ettgtataaa teetggttge tgtetettta tgaggagttg
                                                                  180
tggcccgttg tcaggcaacg tggcgtggtg tgcactgtgt ttgctgacgc aacccccact
                                                                  240
ggttggggca ttgccaccac ctgtcagctc ctttccggga ctttcgcttt ccccctccct
                                                                  300
attgccacgg cggaactcat cgccgcctgc cttgcccgct gctggacagg ggctcggctg
ttgggcactg acaattccgt ggtgttgtcg gggaaatcat cgtcctttcc ttggctgctc
                                                                  420
geetgtgttg ceacetggat tetgegeggg acgteettet getacgteee tteggeeete
aatccagcgg accttectte eegeggeetg etgeeggete tgeggeetet teegegtett
egeettegee eteagaegag teggatetee etttgggeeg eeteeegee taagettate
gataccgtcg agatctaact tgtttattgc agcttataat ggttacaaat aaagcaatag
catcacaaat ttcacaaata aagcattttt ttcactgcat tctagttgtg gtttgtccaa
actcatcaat gtatcttatc atgtctggat ctcgacctcg actagagcat ggctacgtag
ataagtagca tggcgggtta atcattaact acaaggaacc cctagtgatg gagttggcca
ctcctctct gcgcgctcgc tcgctcactg aggccgggcg accaaaggtc gcccgacgcc
cgggctttgc ccgggcggcc tcagtgagcg agcgagcgcg cagctggcgt aatagcgaag
aggeorgeac egategeet teccaacagt tgegeageet gaatggegaa tggegattee
gttgcaatgg ctggcggtaa tattgttctg gatattacca gcaaggccga tagtttgagt
tottotacto aggoaagtga tgttattact aatcaaagaa gtattgogac aacggttaat
                                                                  1140
ttgcgtgatg gacagactct tttactcggt ggcctcactg attataaaaa cacttctcag
                                                                  1200
gattetggeg tacegtteet gtetaaaate eetttaateg geeteetgtt tageteeege
                                                                  1260
totgattota acgaggaaag cacgttatac gtgctcgtca aagcaaccat agtacgcgcc
                                                                  1320
ctgtagcggc gcattaagcg cggcgggtgt ggtggttacg cgcagcgtga ccgctacact
tgccagegcc ctagegcceg etcetttege tttetteeet teettteteg ccaegttege
                                                                  1440
cggctttccc cgtcaagctc taaatcgggg gctcccttta gggttccgat ttagtgcttt
                                                                  1500
acggcacctc gaccccaaaa aacttgatta gggtgatggt tcacgtagtg ggccatcgcc
                                                                  1560
ctgatagacg gtttttcgcc ctttgacgtt ggagtccacg ttctttaata gtggactctt
                                                                  1620
gttccaaact ggaacaacac tcaaccctat ctcggtctat tcttttgatt tataagggat
tttgccgatt tcggcctatt ggttaaaaaa tgagctgatt taacaaaaat ttaacgcgaa
                                                                  1740
ttttaacaaa atattaacgt ttacaattta aatatttgct tatacaatct tcctgttttt
                                                                  1800
```

1860

ggggcttttc tgattatcaa ccggggtaca tatgattgac atgctagttt tacgattacc

gttcatcgat totottgttt gotocagact otcaggoaat gacctgatag cotttgtaga gacctotcaa aaatagotac cototcoggo atgaatttat cagotagaac ggttgaatat

| catattgatg | gtgatttgac | tgtctccggc | ctttctcacc | cgtttgaatc | tttacctaca | 2040 |
|------------|------------|------------|------------|------------|------------|------|
| | gcattgcatt | | | | | 2100 |
| | | | | | | |
| gaaataaagg | cttctcccgc | aaaagtatta | cagggtcata | atgtttttgg | tacaaccgat | 2160 |
| ttaqctttat | gctctgaggc | tttattqctt | aattttqcta | attctttqcc | ttqcctqtat | 2220 |
| | atgttggaat | | | | | 2280 |
| | | | | | | |
| ttcacaccgc | atatggtgca | ctctcagtac | aatctgctct | gatgccgcat | agttaagcca | 2340 |
| qccccqacac | ccgccaacac | ccqctqacqc | gccctgacgg | acttatctac | teceggeate | 2400 |
| | | | | | | 2460 |
| | caagctgtga | | | | | |
| atcaccgaaa | cgcgcgagac | gaaagggcct | cgtgatacgc | ctatttttat | aggttaatgt | 2520 |
| catgataata | atggtttctt | agacgtcagg | tggcactttt | caaaaaata | tacacaaaac | 2580 |
| | | | | | | 2640 |
| | ttatttttct | | | | | |
| ctgataaatg | cttcaataat | attgaaaaag | gaagagtatg | agtattcaac | atttccgtgt | 2700 |
| cacccttatt | cccttttttg | caacatttta | ccttcctatt | tttgctcacc | cagaaacgct | 2760 |
| | | | | | | 2820 |
| | aaagatgctg | | | | | |
| tctcaacagc | ggtaagatcc | ttgagagttt | tegeceegaa | gaacgttttc | caatgatgag | 2880 |
| cacttttaaa | gttctgctat | ataacacaat | attatcccqt | attgacgccg | aacaaaaca | 2940 |
| | | | | | | 3000 |
| | cgcatacact | | | | | |
| aaagcatctt | acggatggca | tgacagtaag | agaattatgc | agtgctgcca | taaccatgag | 3060 |
| tgataacact | gcggccaact | tacttctgac | aacqatcqqa | agaccgaagg | agctaaccgc | 3120 |
| | | | | | | 3180 |
| | aacatggggg | | | | | |
| tgaagccata | ccaaacgacg | agcgtgacac | cacgatgcct | gtagcaatgg | caacaacgtt | 3240 |
| gcgcaaacta | ttaactggcg | aactacttac | tetagettee | cggcaacaat | taatagactg | 3300 |
| | | | | | | |
| garggaggeg | gataaagttg | caggaccact | rergegereg | geeetteegg | erggergger | 3360 |
| tattgctgat | aaatctggag | ccggtgagcg | tgggtctcgc | ggtatcattg | cagcactggg | 3420 |
| | aagccctccc | | | | | 3480 |
| | | | | | | |
| ggatgaacga | aatagacaga | tcgctgagat | aggtgcctca | ctgattaagc | attggtaact | 3540 |
| qtcaqaccaa | gtttactcat | atatacttta | qattqattta | aaacttcatt | tttaatttaa | 3600 |
| | | | | | | 3660 |
| | gtgaagatcc | _ | _ | | | |
| ttcgttccac | tgagcgtcag | accccgtaga | aaagatcaaa | ggatcttctt | gagatccttt | 3720 |
| ttttctacac | gtaatctgct | acttacaaac | aaaaaaacca | ccactaccaa | caataattta | 3780 |
| | | | | | | |
| | caagagctac | | | | | 3840 |
| gataccaaat | actgtccttc | tagtgtagcc | gtagttaggc | caccacttca | agaactctgt | 3900 |
| aggaggggt | acatacctcg | ctctcctaat | cctqttacca | ataactacta | ccagtagcga | 3960 |
| | | | | | | 4020 |
| | cttaccgggt | | | | | |
| gggctgaacg | gggggttcgt | gcacacagcc | cagcttggag | cgaacgacct | acaccgaact | 4080 |
| | cagcgtgagc | | | | | 4140 |
| | | | | | | |
| | gtaagcggca | | | | | 4200 |
| aaacgcctgg | tatctttata | gtcctgtcgg | gtttcgccac | ctctgacttg | agcgtcgatt | 4260 |
| | tcgtcagggg | | | | | 4320 |
| | | | | | | |
| | gccttttgct | | | | | 4380 |
| ttctgtggat | aaccgtatta | ccgcctttga | gtgagctgat | accgctcgcc | gcagccgaac | 4440 |
| | agcgagtcag | | | | | 4500 |
| | | | | | | |
| | cgttggccga | | | | | 4560 |
| qccqcccqqq | caaagcccgg | qcqtcqqqcq | acctttqqtc | qcccqqcctc | aqtqaqcqaq | 4620 |
| | gagagggagt | | | | | 4680 |
| | | | | | | |
| aacccgccat | gctacttatc | tacgtagcca | tgctctagga | cattgattat | tgactagtgg | 4740 |
| aqttccqcqt | tacataactt | acqqtaaatq | qcccqcctqq | ctgaccgccc | aacqaccccc | 4800 |
| | gtcaataatg | | | | | 4860 |
| | | | | | | |
| gacgtcaatg | ggtggagtat | ttacggtaaa | ctgcccactt | ggcagtacat | caagtgtatc | 4920 |
| atatqccaaq | tacgccccct | attqacqtca | atqacqqtaa | atqqcccqcc | tqqcattatq | 4980 |
| | gaccttatgg | | | | | 5040 |
| | | | | | | |
| | ggtcgaggtg | | | | | 5100 |
| ccccaccccc | aattttgtat | ttatttattt | tttaattatt | ttgtgcagcg | atgggggcgg | 5160 |
| | gggcgcgcgc | | | | | 5220 |
| | | | | | | |
| | gcggcggcag | | | | | 5280 |
| gaggcggcgg | cggcggcggc | cctataaaaa | gcgaagcgcg | cggcgggcgg | gagtcgctgc | 5340 |
| | cgccccgtgc | | | | | 5400 |
| | | | | | | |
| | ttactaaaac | | | | | 5460 |
| cgggcgcccc | cctcctcacg | gcgagcgctg | ccacgtcaga | cgaagggcgc | agcgagcgtc | 5520 |
| | cgcccggacg | | | | | 5580 |
| | | | | | | |
| aaccccagta | tcagcagaag | gacattttag | gacgggactt | gggtgactct | agggcactgg | 5640 |
| ttttctttcc | agagagcgga | acaggcgagg | aaaagtagtc | ccttctcggc | gattctgcgg | 5700 |
| | | | | | | 5760 |
| | gtggggcggt | | | | | |
| tttttttc | tacaggtcct | gggtgacgaa | cagggtaccg | ccaccatggc | caccggctct | 5820 |
| | tgctgctggc | | | | | 5880 |
| | | | | | | |
| gccgctagca | tcgataccgt | cgctatgtgc | tggaggcttg | ctgaaggctg | tatgctgata | 5940 |
| atteaaaeta | tcagaatttc | attttaacct | ctgactgacg | aaattotoad | cccgaactat | 6000 |
| | | | | | | |
| | ggcctgttac | - | | | | 6060 |
| tgctgaaqqc | tgtatgctga | gatgaatqcc | tgataqcctt | ccgttttqqc | ctctgactga | 6120 |
| | | | | | | 6180 |
| | tggcattcat | | | | | |
| aatggcctct | agcctggagg | cttgctgaag | gctgtatgct | gagatgaatg | ccaaatagcc | 6240 |
| ttccatttta | gcctctgact | dacddaaddd | tatogcatto | atctcaddad | acaaddcctd | 6300 |
| | | | | | | |
| | ccacatqqaa | caaatggcct | ciclagaat | | | 6339 |

SEQ ID NO: 13 moltype = DNA length = 6339 FEATURE Location/Qualifiers

| source | | 16339 | | | | |
|------------|------------|--------------------------|-------------|------------|------------|--------------|
| | | mol_type : | other DNA | | | |
| | | organism : | = synthetic | construct | | |
| SEQUENCE: | | | | | | |
| | | aatttgtgaa | | | | 60 |
| | | cgctgcttta | | | | 120 180 |
| | | cttgtataaa tggcgtggtg | | | | 240 |
| | | ctgtcagctc | | | | 300 |
| | | cgccgcctgc | | | | 360 |
| | | ggtgttgtcg | | | | 420 |
| | | tctgcgcggg | | | | 480 |
| aatccagcgg | accttccttc | ccgcggcctg | ctgccggctc | tgcggcctct | teegegtett | 540 |
| | | tcggatctcc | | | | 600 |
| | | tgtttattgc | | | | 660 |
| | | aagcattttt | _ | | | 720 |
| | | atgtctggat | | | | 780 840 |
| | | atcattaact tcgctcactg | | | | 900 |
| | | tcagtgagcg | | | | 960 |
| | | tcccaacagt | | | | 1020 |
| | | tattgttctg | | | | 1080 |
| | | tgttattact | | | | 1140 |
| ttgcgtgatg | gacagactct | tttactcggt | ggcctcactg | attataaaaa | cacttctcag | 1200 |
| gattctggcg | taccgttcct | gtctaaaatc | cctttaatcg | gcctcctgtt | tagctcccgc | 1260 |
| | | cacgttatac | | | | 1320 |
| | | cggcgggtgt | | | | 1380 |
| | | ctcctttcgc | | | | 1440 |
| | | taaatcgggg | - | | | 1500 1560 |
| | | aacttgatta ctttgacgtt | | | | 1620 |
| | | tcaaccctat | | | | 1680 |
| | | ggttaaaaaa | | | | 1740 |
| | | ttacaattta | | | | 1800 |
| ggggcttttc | tgattatcaa | ccggggtaca | tatgattgac | atgctagttt | tacgattacc | 1860 |
| gttcatcgat | tctcttgttt | gctccagact | ctcaggcaat | gacctgatag | cctttgtaga | 1920 |
| | | cctctccggc | | | | 1980 |
| | | tgtctccggc | | | | 2040 |
| | | taaaatatat | | | | 2100 |
| | | aaaagtatta tttattgctt | | | | 2160 2220 |
| | | tcctgatgcg | | | | 2280 |
| | | ctctcagtac | | | | 2340 |
| | | ccgctgacgc | | | | 2400 |
| | | ccgtctccgg | | | | 2460 |
| atcaccgaaa | cgcgcgagac | gaaagggcct | cgtgatacgc | ctatttttat | aggttaatgt | 2520 |
| | | agacgtcagg | | | | 2580 |
| | | aaatacattc | | | | 2640 |
| | | attgaaaaag | | | | 2700 |
| | | cggcattttg | | | | 2760 2820 |
| | | aagatcagtt ttgagagttt | | | | 2880 |
| | | gtggcgcggt | | | | 2940 |
| | | attctcagaa | | | | 3000 |
| | | tgacagtaag | | | | 3060 |
| | | tacttctgac | | | | 3120 |
| | | atcatgtaac | | | | 3180 |
| | | agcgtgacac | | | _ | 3240 |
| | | aactacttac | | | | 3300 |
| | | caggaccact | | | | 3360 3420 |
| | | ccggtgagcg gtatcgtagt | | | | 3480 |
| | | tcgctgagat | | | | 3540 |
| | | atatacttta | | | | 3600 |
| | | tttttgataa | | | | 3660 |
| | | accccgtaga | _ | | | 3720 |
| | | gcttgcaaac | | | | 3780 |
| | | caactctttt | | | | 3840 |
| | | tagtgtagcc | | | | 3900 |
| | | ctctgctaat | | | | 3960 |
| | _ | tggactcaag | - | | | 4020 |
| | | gcacacagcc | | | | 4080 |
| | | tatgagaaag | | | | 4140 |
| caggtatccg | gtaagcggca | gggtcggaac | aggagagcgc | acgagggagc | ttccaggggg | 4200 |
| aaacgcctgg | tatctttata | gtcctgtcgg | gtttcgccac | ctctgacttg | agcgtcgatt | 4260 |
| tttgtgatgc | tcgtcagggg | ggcggagcct | atggaaaaac | gccagcaacg | cggccttttt | 4320 |
| | | | | | | |

```
acggttcctg gccttttgct ggccttttgc tcacatgttc tttcctgcgt tatcccctga
ttctgtggat aaccgtatta ccgcctttga gtgagctgat accgctcgcc gcagccgaac
                                                                   4440
gaccgagcgc agcgagtcag tgagcgagga agcggaagag cgcccaatac gcaaaccgcc
                                                                   4500
teteceegeg egttggeega tteattaatg eageagetge gegetegete geteaetgag
                                                                   4560
gccgcccggg caaagcccgg gcgtcgggcg acctttggtc gcccggcctc agtgagcgag
                                                                   4620
cgagcgcgca gagagggagt ggccaactcc atcactaggg gttccttgta gttaatgatt
                                                                   4680
aacccgccat gctacttatc tacgtagcca tgctctagga cattgattat tgactagtgg
                                                                   4740
agttccgcgt tacataactt acggtaaatg gcccgcctgg ctgaccgccc aacgaccccc
                                                                   4800
gcccattgac gtcaataatg acgtatgttc ccatagtaac gccaataggg actttccatt
                                                                   4860
gacgtcaatg ggtggagtat ttacggtaaa ctgcccactt ggcagtacat caagtgtatc
atatgccaag tacgcccct attgacgtca atgacggtaa atggcccgcc tggcattatg
                                                                   4980
cccagtacat gaccttatgg gactttccta cttggcagta catctacgta ttagtcatcg
                                                                   5040
ctattaccat ggtcgaggtg agccccacgt tctgcttcac tctccccatc tccccccct
ccccacccc aattttgtat ttatttattt tttaattatt ttgtgcagcg atggggggg
ggggggggg gggcgcgcgc caggcggggc ggggcggggc gaggggggg gcggggcgag
gcggagaggt gcggcggcag ccaatcagag cggcgcgctc cgaaagtttc cttttatggc
gaggeggegg eggeggegge ectataaaaa gegaagegeg eggegggegg gagtegetge
gegetgeett egeceegtge eeegeteege egecgeeteg egecgeeege eeeggetetg
actgaccgcg ttactaaaac aggtaagtcc ggcctccgcg ccgggttttg gcgcctcccg
                                                                   5460
eggegeeee ectecteacg gegagegetg ceacqteaga egaagggege agegagegte
                                                                   5520
ctgatectte egeceggaeg eteaggaeag eggeeegetg eteataagae teggeettag
                                                                   5580
aaccccagta tcagcagaag gacattttag gacgggactt gggtgactct agggcactgg
                                                                   5640
ttttctttcc agagagcgga acaggcgagg aaaagtagtc ccttctcggc gattctgcgg
                                                                   5700
agggatetee gtggggeggt gaacgeegat gatgeeteta etaaceatgt teatgtttte
                                                                   5760
ttttttttt tacaggtcct gggtgacgaa cagggtaccg ccaccatggc caccggctct
                                                                   5820
cgcacaagcc tgctgctggc tttcggactg ctgtgcctgc cttggctcca ggagggctcc
                                                                   5880
geogetagea tegatacegt egetatgtge tggaggettg etgaaggetg tatgetgtte
                                                                   5940
agatcatctg ggtatccggc gttttggcct ctgactgacg ccggataccg atgatctgaa
                                                                   6000
caggacacaa qqcctqttac tagcactcac atqqaacaaa tqqcctctaq cctqqaqqct
                                                                   6060
tgctgaaggc tgtatgctga ataatcagat ggtgttgcga tcgttttggc ctctgactga
                                                                   6120
cgategeaac accetgatta tteaggacae aaggeetgtt actageacte acatggaaca
                                                                   6180
aatggcctct agcctggagg cttgctgaag gctgtatgct gtactgatgc acaatttaaa
                                                                   6240
cgccgttttg gcctctgact gacggcgttt aaagtgcatc agtacaggac acaaggcctg
                                                                   6300
ttactagcac tcacatggaa caaatggcct ctctagaat
                                                                   6339
SEO ID NO: 14
                       moltype = DNA length = 6333
FEATURE
                       Location/Qualifiers
source
                       1..6333
                       mol_type = other DNA
organism = synthetic construct
SEQUENCE: 14
aatcaacctc tggattacaa aatttgtgaa agattgactg gtattcttaa ctatgttgct
                                                                   120
ccttttacgc tatgtggata cgctgcttta atgcctttgt atcatgctat tgcttcccgt
atggetttea tttteteete ettgtataaa teetggttge tgtetettta tgaggagttg
                                                                   180
tggcccgttg tcaggcaacg tggcgtggtg tgcactgtgt ttgctgacgc aacccccact
                                                                   240
ggttggggca ttgccaccac ctgtcagctc ctttccggga ctttcgcttt ccccctccct
                                                                   300
attgccacgg cggaactcat cgccgcctgc cttgcccgct gctggacagg ggctcggctg
ttgggcactg acaattccgt ggtgttgtcg gggaaatcat cgtcctttcc ttggctgctc
                                                                   420
geetgtgttg ceacetggat tetgegeggg acgteettet getacgteee tteggeeete
aatccagegg accttectte eegeggeetg etgeeggete tgeggeetet teegegtett
                                                                   540
cgccttcgcc ctcagacgag tcggatctcc ctttgggccg cctccccgcc taagcttatc
gataccgtcg agatctaact tgtttattgc agcttataat ggttacaaat aaagcaatag
catcacaaat ttcacaaata aagcattttt ttcactgcat tctagttgtg gtttgtccaa
actcatcaat gtatcttatc atgtctggat ctcgacctcg actagagcat ggctacgtag
                                                                   780
ataagtagca tggcgggtta atcattaact acaaggaacc cctagtgatg gagttggcca
ctcctctct gcgcgctcgc tcgctcactg aggccgggcg accaaaggtc gcccgacgcc
cgggctttgc ccgggcggcc tcagtgagcg agcgagcgcg cagctggcgt aatagcgaag
aggeegeac egategeet teccaacagt tgegeageet gaatggegaa tggegattee
gttgcaatgg ctggcggtaa tattgttctg gatattacca gcaaggccga tagtttgagt
tettetaete aggeaagtga tgttattaet aateaaagaa gtattgegae aacggttaat
                                                                   1140
ttgcgtgatg gacagactct tttactcggt ggcctcactg attataaaaa cacttctcag
gattetggeg taccettect gtetaaaate cetttaateg geeteetgtt tageteeege
                                                                   1260
tctgattcta acgaggaaag cacgttatac gtgctcgtca aagcaaccat agtacgcgcc
                                                                   1320
ctgtagcggc gcattaagcg cggcgggtgt ggtggttacg cgcagcgtga ccgctacact
                                                                   1380
tgccagcgcc ctagcgcccg ctcctttcgc tttcttccct tcctttctcg ccacgttcgc
                                                                   1440
eggettteee egteaagete taaategggg geteeettta gggtteegat ttagtgettt
                                                                   1500
acggcacctc gaccccaaaa aacttgatta gggtgatggt tcacgtagtg ggccatcgcc
                                                                   1560
ctgatagacg gtttttcgcc ctttgacgtt ggagtccacg ttctttaata gtggactctt
                                                                   1620
gttccaaact ggaacaacac tcaaccctat ctcggtctat tcttttgatt tataagggat
                                                                   1680
tttgccgatt tcggcctatt ggttaaaaaa tgagctgatt taacaaaaat ttaacgcgaa
                                                                   1740
ttttaacaaa atattaacgt ttacaattta aatatttgct tatacaatct tcctgttttt
ggggcttttc tgattatcaa ccggggtaca tatgattgac atgctagttt tacgattacc
                                                                   1860
gttcatcgat tctcttgttt gctccagact ctcaggcaat gacctgatag cctttgtaga
                                                                   1920
gacctctcaa aaatagctac cctctccggc atgaatttat cagctagaac ggttgaatat
                                                                   1980
catattgatg gtgatttgac tgtctccggc ctttctcacc cgtttgaatc tttacctaca
                                                                   2040
```

cattactcag gcattgcatt taaaatatat gagggttcta aaaattttta tccttgcgtt

| gaaataaagg | cttctcccgc | aaaagtatta | cagggtcata | atgtttttgg | tacaaccgat | 2160 |
|------------|-------------|------------|------------|------------|------------|--------------|
| ttagctttat | actictaaaac | tttattgctt | aattttgcta | attetttgcc | ttacctatat | 2220 |
| | | | gtattttctc | | | 2280 |
| | | | | | | |
| ttcacaccgc | atatggtgca | ctctcagtac | aatctgctct | gatgccgcat | agttaagcca | 2340 |
| gccccgacac | ccgccaacac | ccgctgacgc | gccctgacgg | gcttgtctgc | tcccggcatc | 2400 |
| cacttacaga | caagctgtga | ccatctccaa | gagctgcatg | tatcagaggt | tttcaccatc | 2460 |
| | | | | | | 2520 |
| | | | cgtgatacgc | | | |
| catgataata | atggtttctt | agacgtcagg | tggcactttt | cggggaaatg | tgcgcggaac | 2580 |
| ccctatttgt | ttatttttct | aaatacattc | aaatatgtat | ccgctcatga | gacaataacc | 2640 |
| | | | gaagagtatg | | | 2700 |
| | | | | | | 2760 |
| | | | ccttcctgtt | | | |
| ggtgaaagta | aaagatgctg | aagatcagtt | gggtgcacga | gtgggttaca | tcgaactgga | 2820 |
| tctcaacagc | ggtaagatcc | ttgagagttt | tcgccccgaa | gaacgttttc | caatgatgag | 2880 |
| | | | attatcccgt | | | 2940 |
| | | | - | | | 3000 |
| | | | tgacttggtt | | | |
| aaagcatctt | acggatggca | tgacagtaag | agaattatgc | agtgctgcca | taaccatgag | 3060 |
| tgataacact | gcggccaact | tacttctgac | aacgatcgga | ggaccgaagg | agctaaccgc | 3120 |
| | | | tcgccttgat | | | 3180 |
| | | | | | | |
| | | | cacgatgcct | | | 3240 |
| gcgcaaacta | ttaactggcg | aactacttac | tctagcttcc | cggcaacaat | taatagactg | 3300 |
| qatqqaqqcq | qataaaqttq | caqqaccact | tctgcgctcg | qcccttccqq | ctqqctqqtt | 3360 |
| | | | tgggtctcgc | | | 3420 |
| | | | | | | 3480 |
| | | | tatctacacg | | | |
| ggatgaacga | aatagacaga | tcgctgagat | aggtgcctca | ctgattaagc | attggtaact | 3540 |
| gtcagaccaa | gtttactcat | atatacttta | gattgattta | aaacttcatt | tttaatttaa | 3600 |
| | _ | | tctcatgacc | | | 3660 |
| | | | | | | 3720 |
| | | | aaagatcaaa | | | |
| | | | aaaaaaacca | | | 3780 |
| tttgccggat | caagagctac | caactctttt | tccgaaggta | actggcttca | gcagagcgca | 3840 |
| gataccaaat | actotectte | tagtgtagcc | gtagttaggc | caccacttca | agaactctgt | 3900 |
| | | | cctgttacca | | | 3960 |
| | | | | | | 4020 |
| | | | acgatagtta | | | |
| | | | cagcttggag | | | 4080 |
| gagataccta | cagcgtgagc | tatgagaaag | cgccacgctt | cccgaaggga | gaaaggcgga | 4140 |
| caggtatccg | qtaaqcqqca | gggtcggaac | aggagagcgc | acqaqqqaqc | ttccaqqqqq | 4200 |
| | | | gtttcgccac | | | 4260 |
| | | | | | | 4320 |
| | | | atggaaaaac | | | |
| acggttcctg | gccttttgct | ggccttttgc | tcacatgttc | tttcctgcgt | tatcccctga | 4380 |
| ttctgtggat | aaccgtatta | ccgcctttga | gtgagctgat | accgctcgcc | gcagccgaac | 4440 |
| qaccqaqcqc | agcgagtcag | tqaqcqaqqa | agcggaagag | cqcccaatac | gcaaaccgcc | 4500 |
| | | | cagcagctgc | | | 4560 |
| | | | acctttggtc | | | 4620 |
| | | | | | | |
| | | | atcactaggg | | | 4680 |
| aacccgccat | gctacttatc | tacgtagcca | tgctctagga | cattgattat | tgactagtgg | 4740 |
| aqttccqcqt | tacataactt | acqqtaaatq | gcccgcctgg | ctqaccqccc | aacqaccccc | 4800 |
| | | | ccatagtaac | | | 4860 |
| | | | | | | 4920 |
| | | | ctgcccactt | | | |
| | | | atgacggtaa | | | 4980 |
| cccagtacat | gaccttatgg | gactttccta | cttggcagta | catctacgta | ttagtcatcg | 5040 |
| ctattaccat | qqtcqaqqtq | agccccacqt | tctgcttcac | tctccccatc | tccccccct | 5100 |
| | | | tttaattatt | | | 5160 |
| | | | ggggcggggc | | | 5220 |
| | | | | | | |
| | | | cggcgcgctc | | | 5280 |
| gaggcggcgg | cggcggcggc | cctataaaaa | gcgaagcgcg | cggcgggcgg | gagtcgctgc | 5340 |
| qcqctqcctt | cqccccqtqc | cccqctccqc | cgccgcctcg | cqccqcccqc | cccqqctctq | 5400 |
| | | | ggcctccgcg | | | 5460 |
| | | | | | | 5520 |
| | | | ccacgtcaga | | | |
| | | | cggcccgctg | | | 5580 |
| aaccccagta | tcagcagaag | gacattttag | gacgggactt | gggtgactct | agggcactgg | 5640 |
| ttttctttcc | agagagcgga | acaggcgagg | aaaagtagtc | ccttctcggc | gattctgcgg | 5700 |
| | | | gatgcctcta | | | 5760 |
| | | | cagggtacca | _ | _ | 5820 |
| | | | | | | |
| | | | ctgccttggc | | | 5880 |
| agcatcgata | ccgtcgctat | gtgctggagg | cttgctgaag | gctgtatgct | gaggtaatcg | 5940 |
| gctgatgcgt | ttqcqtttta | gcctctgact | gacgcaaacg | catqccqatt | acctcaqqac | 6000 |
| | | | | | | 6060 |
| | _ | | caaatggcct | | | |
| | | | ttcagcgttt | | | 6120 |
| aaggcaccgt | gtataacagg | acacaaggcc | tgttactagc | actcacatgg | aacaaatggc | 6180 |
| ctctagcctg | | gaaggetgta | tactgatcag | atcatcaaqt | tcaqcaqcqt | 6240 |
| | qaqqcttqct | | | | | |
| tttaacctct | | | | | | 6300 |
| | gactgacgct | gctgaacgat | gatctgatca | | | 6300 |
| | | gctgaacgat | gatctgatca | | | 6300 6333 |
| | gactgacgct | gctgaacgat | gatctgatca | | | |

SEQ ID NO: 15 moltype = DNA length = 6339
FEATURE Location/Qualifiers
source 1..6339
mol_type = other DNA

| | | organism : | synthetic | construct | | |
|-------------|------------|------------|------------|--------------------------|------------|--------------|
| SEQUENCE: 3 | 15 | | | | | |
| aatcaacctc | tggattacaa | aatttgtgaa | agattgactg | gtattcttaa | ctatgttgct | 60 |
| | | | | atcatgctat | | 120 |
| | | | | tgtctcttta | | 180 |
| | | | | ttgctgacgc | | 240 300 |
| | | | | ctttcgcttt gctggacagg | | 360 |
| | | | | cgtcctttcc | | 420 |
| | | | | gctacgtccc | | 480 |
| | | | | tgcggcctct | | 540 |
| cgccttcgcc | ctcagacgag | tcggatctcc | ctttgggccg | cctccccgcc | taagcttatc | 600 |
| | | | | ggttacaaat | | 660 |
| | | | | tctagttgtg | | 720 |
| | | | | actagagcat | | 780 |
| | | | | cctagtgatg | | 840 900 |
| | | | | accaaaggtc cagctggcgt | | 960 |
| | | | | gaatggcgaa | | 1020 |
| | | | | gcaaggccga | | 1080 |
| | | | | gtattgcgac | | 1140 |
| ttgcgtgatg | gacagactct | tttactcggt | ggcctcactg | attataaaaa | cacttctcag | 1200 |
| | | | | gcctcctgtt | | 1260 |
| | | | | aagcaaccat | | 1320 |
| | | | | cgcagcgtga | | 1380 |
| | | - | | tcctttctcg | | 1440 1500 |
| | | | | gggttccgat tcacgtagtg | | 1560 |
| | | | | ttctttaata | | 1620 |
| | | | | tcttttgatt | | 1680 |
| - | | | | taacaaaaat | | 1740 |
| | | | | tatacaatct | | 1800 |
| ggggcttttc | tgattatcaa | ccggggtaca | tatgattgac | atgctagttt | tacgattacc | 1860 |
| | | | | gacctgatag | | 1920 |
| - | _ | | - | cagctagaac | | 1980 |
| | | | | cgtttgaatc | | 2040 2100 |
| | | | | aaaattttta atgtttttgg | | 2160 |
| | | | | attetttgee | | 2220 |
| | | | | cttacgcatc | | 2280 |
| | | | | gatgccgcat | | 2340 |
| gccccgacac | ccgccaacac | ccgctgacgc | gccctgacgg | gcttgtctgc | teceggeate | 2400 |
| | | | | tgtcagaggt | | 2460 |
| | | | | ctatttttat | | 2520 |
| | | | | cggggaaatg | | 2580 |
| | | | | ccgctcatga | | 2640 2700 |
| | | | | agtattcaac tttgctcacc | | 2760 |
| | | | | gtgggttaca | | 2820 |
| | | | | gaacgttttc | | 2880 |
| | | | | attgacgccg | | 2940 |
| actcggtcgc | cgcatacact | attctcagaa | tgacttggtt | gagtactcac | cagtcacaga | 3000 |
| | | | | agtgctgcca | | 3060 |
| | | | | ggaccgaagg | | 3120 |
| | | | | cgttgggaac | | 3180 3240 |
| | | | | gtagcaatgg cggcaacaat | | 3300 |
| | | | | gcccttccgg | | 3360 |
| | | | | ggtatcattg | | 3420 |
| | | | | acggggagtc | | 3480 |
| | | | | ctgattaagc | | 3540 |
| gtcagaccaa | gtttactcat | atatacttta | gattgattta | aaacttcatt | tttaatttaa | 3600 |
| aaggatctag | gtgaagatcc | tttttgataa | tctcatgacc | aaaatccctt | aacgtgagtt | 3660 |
| | | | | ggatcttctt | | 3720 |
| | | | | ccgctaccag | | 3780 |
| | | | | actggcttca | | 3840 |
| | | | | caccacttca | | 3900 |
| | | | | gtggctgctg | | 3960 |
| | | | | ccggataagg | | 4020 4080 |
| | | | | cgaacgacct | | 4140 |
| | | | | cccgaaggga | | 4200 |
| | | | | acgagggagc ctctgacttg | | 4260 |
| | | | | gccagcaacg | | 4320 |
| | | | | tttcctgcgt | | 4380 |
| | | | | accgctcgcc | | 4440 |
| | - | - | | | - | |

```
gaccgagcgc agcgagtcag tgagcgagga agcggaagag cgcccaatac gcaaaccgcc
teteccegeg egttggeega tteattaatg cageagetge gegetegete geteactgag
                                                                   4560
geogeologic caaageologic gegteggeg acctttggte geologic agtgagegag
                                                                   4620
cgagcgcgca gagagggagt ggccaactcc atcactaggg gttccttgta gttaatgatt
                                                                   4680
aacccgccat gctacttatc tacgtagcca tgctctagga cattgattat tgactagtgg
                                                                   4740
agttccgcgt tacataactt acggtaaatg gcccgcctgg ctgaccgccc aacgaccccc
                                                                   4800
gcccattgac gtcaataatg acgtatgttc ccatagtaac gccaataggg actttccatt
                                                                   4860
gacgtcaatg ggtggagtat ttacggtaaa ctgcccactt ggcagtacat caagtgtatc
                                                                   4920
atatgccaag tacgcccct attgacgtca atgacggtaa atggcccgcc tggcattatg
                                                                   4980
cccagtacat gaccttatgg gactttccta cttggcagta catctacgta ttagtcatcg
ctattaccat ggtcgaggtg agcccacgt tctgcttcac tctccccatc tccccccct
                                                                   5100
ccccacccc aattitgtat ttatttattt tttaattatt ttgtgcagcg atggggggg
                                                                   5160
ggggggggg gggcgcgcc caggcggggc ggggcggggc gaggggcggg gcggggcgag
gcggagaggt gcggcggcag ccaatcagag cggcgcgctc cgaaagtttc cttttatggc
gaggeggegg eggeggege ectataaaaa gegaagegeg eggegggegg gagtegetge
                                                                   5340
gegetgeett egeceegtge eeegeteege egeegeeteg egeegeeege eeeggetetg
actgaccgcg ttactaaaac aggtaagtcc ggcctccgcg ccgggttttg gcgcctccccg
egggegeee ectecteacg gegagegetg ceacqteaga egaagggege agegagegte
ctgatccttc cgcccggacg ctcaggacag cggcccgctg ctcataagac tcggccttag
aaccccaqta tcaqcaqaaq qacattttaq qacqqqactt qqqtqactct aqqqcactqq
ttttctttcc agagagcgga acaggcgagg aaaagtagtc ccttctcggc gattctgcgg
                                                                   5700
agggatetee gtggggeggt gaacgeegat gatgeeteta etaaceatgt teatgtttte
                                                                   5760
ttttttttt tacaggtcct gggtgacgaa cagggtaccg ccaccatggc caccggctct
                                                                   5820
cgcacaagcc tgctgctggc tttcggactg ctgtgcctgc cttggctcca ggagggctcc
                                                                   5880
geogetagea tegatacegt egetatgtge tggaggettg etgaaggetg tatgetgaca
                                                                   5940
taaagctatg tcctggcagc gttttggcct ctgactgacg ctgccaggat agctttatgt
                                                                   6000
caggacacaa ggcctgttac tagcactcac atggaacaaa tggcctctag cctggaggct
                                                                   6060
tgctgaaggc tgtatgctga tacagcagat atcgcgaatt tcgttttggc ctctgactga
                                                                   6120
cgaaattcgc gatctgctgt atcaggacac aaggcctgtt actagcactc acatqqaaca
                                                                   6180
aatggeetet ageetggagg ettgetgaag getgtatget gttteagttt aaaateageg
                                                                   6240
ccaegttttg geetetgaet gaegtggege tgattaaaet gaaacaggae acaaggeetg
                                                                   6300
ttactagcac tcacatggaa caaatggcct ctctagaat
                                                                   6339
SEO ID NO. 16
                      moltype = DNA length = 6339
                      Location/Qualifiers
FEATURE
source
                      1..6339
                      mol_type = other DNA
                      organism = synthetic construct
SEQUENCE: 16
aatcaacctc tggattacaa aatttgtgaa agattgactg gtattcttaa ctatgttgct
                                                                   120
cettttacge tatgtggata egetgettta atgeetttgt ateatgetat tgetteeegt
atggetttea titteteete etigtataaa teetggtige tgtetettia tgaggagtig
                                                                   180
tggcccgttg tcaggcaacg tggcgtggtg tgcactgtgt ttgctgacgc aacccccact
                                                                   240
ggttggggca ttgccaccac ctgtcagctc ctttccggga ctttcgcttt ccccctccct
                                                                   300
attgccacgg cggaactcat cgccgcctgc cttgcccgct gctggacagg ggctcggctg
                                                                   360
ttgggcactg acaattccgt ggtgttgtcg gggaaatcat cgtcctttcc ttggctgctc
                                                                   420
geetgtgttg ccacetggat tetgegeggg aegteettet getaegteee tteggeeete
                                                                   480
aatccagegg accttectte eegeggeetg etgeeggete tgeggeetet teegegtett
                                                                   540
cgccttcgcc ctcagacgag tcggatctcc ctttgggccg cctccccgcc taagcttatc
                                                                   600
gataccgtcg agatctaact tgtttattgc agcttataat ggttacaaat aaagcaatag
                                                                   660
catcacaaat ttcacaaata aagcattttt ttcactgcat tctagttgtg gtttgtccaa
actcatcaat gtatcttatc atgtctggat ctcgacctcg actagagcat ggctacgtag
                                                                   780
ataagtagca tggcgggtta atcattaact acaaggaacc cctagtgatg gagttggcca
ctccctctct gcgcgctcgc tcgctcactg aggccgggcg accaaaggtc gcccgacgcc
                                                                   900
cgggctttgc ccgggcggcc tcagtgagcg agcgagcgcg cagctggcgt aatagcgaag
aggecegeae egategeeet teccaacagt tgegeageet gaatggegaa tggegattee
gttgcaatgg ctggcggtaa tattgttctg gatattacca gcaaggccga tagtttgagt
tottotacto aggoaagtga tgttattact aatcaaagaa gtattgogac aacggttaat
ttgcgtgatg gacagactct tttactcggt ggcctcactg attataaaaa cacttctcag
gattetggeg taccettect gtetaaaate cetttaateg geeteetgtt tageteeege
                                                                   1260
totgattota acgaggaaag cacgttatac gtgctcgtca aagcaaccat agtacgcgcc
ctqtaqcqqc qcattaaqcq cqqcqqtqt qqtqqttacq cqcaqcqtqa ccqctacact
tgccagcgcc ctagcgcccg ctcctttcgc tttcttccct tcctttctcg ccacgttcgc
                                                                   1440
eggettteee egteaagete taaategggg geteeettta gggtteegat ttagtgettt
                                                                   1500
acggcacctc gaccccaaaa aacttgatta gggtgatggt tcacgtagtg ggccatcgcc
                                                                   1560
ctgatagacg gtttttcgcc ctttgacgtt ggagtccacg ttctttaata gtggactctt
gttccaaact ggaacaacac tcaaccctat ctcggtctat tcttttgatt tataagggat
                                                                   1680
tttqccgatt tcggcctatt ggttaaaaaa tgagctgatt taacaaaaaat ttaacgcgaa
                                                                   1740
ttttaacaaa atattaacgt ttacaattta aatatttgct tatacaatct tcctgttttt
                                                                   1800
ggggcttttc tgattatcaa ccggggtaca tatgattgac atgctagttt tacgattacc
                                                                   1860
gttcatcgat tctcttgttt gctccagact ctcaggcaat gacctgatag cctttgtaga
qacctctcaa aaataqctac cctctccqqc atqaatttat caqctaqaac qqttqaatat
catattgatg gtgatttgac tgtctccggc ctttctcacc cgtttgaatc tttacctaca
cattactcag gcattgcatt taaaatatat gagggttcta aaaattttta tccttgcgtt
                                                                   2100
gaaataaagg cttctcccgc aaaagtatta cagggtcata atgtttttgg tacaaccgat
```

ttagetttat getetgagge tttattgett aattttgeta attetttgee ttgeetgtat

```
gatttattgg atgttggaat tcctgatgcg gtattttctc cttacgcatc tgtgcggtat
ttcacaccgc atatggtgca ctctcagtac aatctgctct gatgccgcat agttaagcca
                                                                   2340
geocegacae cegecaacae eegetgaege geoctgaegg gettgtetge teeeggeate
                                                                   2400
cgcttacaga caagctgtga ccgtctccgg gagctgcatg tgtcagaggt tttcaccgtc
                                                                   2460
atcaccgaaa cgcgcgagac gaaagggcct cgtgatacgc ctatttttat aggttaatgt
                                                                   2520
catgataata atggtttctt agacgtcagg tggcactttt cggggaaatg tgcgcggaac
                                                                   2580
ccctatttgt ttattttct aaatacattc aaatatgtat ccgctcatga gacaataacc
                                                                   2640
ctgataaatg cttcaataat attgaaaaag gaagagtatg agtattcaac atttccgtgt
                                                                   2700
cgcccttatt cccttttttg cggcattttg ccttcctgtt tttgctcacc cagaaacgct
                                                                   2760
ggtgaaagta aaagatgctg aagatcagtt gggtgcacga gtgggttaca tcgaactgga
teteaacage ggtaagatee ttgagagttt tegeceegaa gaacgtttte caatgatgag
                                                                   2880
cacttttaaa gttctgctat gtggcgcggt attatcccgt attgacgccg ggcaagagca
                                                                   2940
actoggtogo ogcatacact attotoagaa tgacttggtt gagtactoac cagtcacaga
                                                                   3000
aaagcatctt acggatggca tgacagtaag agaattatgc agtgctgcca taaccatgag
tgataacact geggecaact tacttetgac aacgategga ggacegaagg agetaacege
                                                                   3120
ttttttgcac aacatggggg atcatgtaac tcgccttgat cgttgggaac cggagctgaa
tgaagccata ccaaacgacg agcgtgacac cacgatgcct gtagcaatgg caacaacgtt
gegeaaacta ttaactggeg aactacttac tetagettee eggeaacaat taatagactg
gatggaggcg gataaagttg caggaccact tetgegeteg gecetteegg etggetggtt
                                                                   3360
tattgctgat aaatctggag ccggtgagcg tgggtctcgc ggtatcattg cagcactggg
                                                                   3420
gccagatggt aagccctccc gtatcgtagt tatctacacg acggggagtc aggcaactat
                                                                   3480
ggatgaacga aatagacaga tcgctgagat aggtgcctca ctgattaagc attggtaact
                                                                   3540
3600
aaggatctag gtgaagatcc tttttgataa tctcatgacc aaaatccctt aacgtgagtt
                                                                   3660
ttcgttccac tgagcgtcag accccgtaga aaagatcaaa ggatcttctt gagatccttt
                                                                   3720
ttttctgcgc gtaatctgct gcttgcaaac aaaaaaacca ccgctaccag cggtggtttg
                                                                   3780
tttgccggat caagagctac caactctttt tccgaaggta actggcttca gcagagcgca
                                                                   3840
gataccaaat actgtccttc tagtgtagcc gtagttaggc caccacttca agaactctgt
                                                                   3900
agcaccgcct acatacctcg ctctgctaat cctgttacca gtggctgctg ccagtggcga taagtcgtgt cttaccgggt tggactcaag acgatagtta ccggataagg cgcagcggtc
                                                                   3960
                                                                   4020
gggctgaacg gggggttcgt gcacacagcc cagcttggag cgaacgacct acaccgaact
                                                                   4080
gagataccta cagcgtgagc tatgagaaag cgccacgctt cccgaaggga gaaaggcgga
                                                                   4140
caggtatecg gtaageggea gggteggaac aggagagege acgagggage ttecaggggg
                                                                   4200
aaacgcctgg tatctttata gtcctgtcgg gtttcgccac ctctgacttg agcgtcgatt
                                                                   4260
tttgtgatgc tcgtcagggg ggcggagcct atggaaaaac gccagcaacg cggccttttt
                                                                   4320
acggttcctg gccttttgct ggccttttgc tcacatgttc tttcctgcgt tatcccctga
                                                                   4380
ttctgtggat aaccgtatta ccgcctttga gtgagctgat accgctcgcc gcagccgaac
                                                                   4440
gaccgagcgc agcgagtcag tgagcgagga agcggaagag cgcccaatac gcaaaccgcc
                                                                   4500
teteceegeg egitggeega ticattaatg eageagetge gegetegete geteactgag
                                                                   4560
geogeologic caaageologic gegtegggeg acetttggte geologicete agtgagegag
                                                                   4620
cgagcgcgca gagagggagt ggccaactcc atcactaggg gttccttgta gttaatgatt
                                                                   4680
aaccegecat getaettate taegtageca tgetetagga cattgattat tgaetagtgg
                                                                   4740
agttccgcgt tacataactt acggtaaatg gcccgcctgg ctgaccgccc aacgaccccc
                                                                   4800
gcccattgac gtcaataatg acgtatgttc ccatagtaac gccaataggg actttccatt
                                                                   4860
gacgtcaatg ggtggagtat ttacggtaaa ctgcccactt ggcagtacat caagtgtatc
                                                                   4920
atatgccaag tacgcccct attgacgtca atgacggtaa atggcccgcc tggcattatg
                                                                   4980
cccagtacat gaccttatgg gactttccta cttggcagta catctacgta ttagtcatcg
                                                                   5040
ctattaccat ggtcgaggtg agccccacgt tctgcttcac tctccccatc tccccccct
                                                                   5100
ccccacccc aattttgtat ttatttattt tttaattatt ttgtgcagcg atggggggg
                                                                   5160
ggggggggg gggcgcgcc caggcggggc ggggcggggc gaggggcggg gcggggcgag
                                                                   5220
gcggagaggt gcggcggcag ccaatcagag cggcgcgctc cgaaagtttc cttttatggc
                                                                   5280
gaggeggegg eggeggegge cetataaaaa gegaagegeg eggegggegg gagtegetge
                                                                   5340
gegetgeett egeceegtge eeegeteege egeegeeteg egeegeeege eeeggetetg
                                                                   5400
actgaccgcg ttactaaaac aggtaagtcc ggcctccgcg ccgggttttg gcgcctcccg
                                                                   5460
cgggcgcccc cctcctcacg gcgagcgctg ccacgtcaga cgaagggcgc agcgagcgtc
ctgatccttc cgcccggacg ctcaggacag cggcccgctg ctcataagac tcggccttag
aaccccagta tcagcagaag gacattttag gacgggactt gggtgactct agggcactgg
ttttctttcc agagagcgga acaggcgagg aaaagtagtc ccttctcggc gattctgcgg
agggatetee gtggggeggt gaacgeegat gatgeeteta etaaceatgt teatgtttte
tttttttttttttt tacaggtett gggtgaegaa cagggtaeeg ceaceatgge caeeggetet
cgcacaagcc tgctgctggc tttcggactg ctgtgcctgc cttggctcca ggagggctcc
geogetagea tegatacegt egetatgtee tggaggettg etgaaggetg tatgetgata
caggogtttg tgctcggttc gttttggcct ctgactgacg aaccgagcaa acgcctgtat
                                                                   6000
caggacacaa ggcctgttac tagcactcac atggaacaaa tggcctctag cctggaggct
                                                                   6060
tgctgaaggc tgtatgctgt catgatcaat atggtatccg gcgttttggc ctctgactga
                                                                   6120
cgccggatac cattgatcat gacaggacac aaggcctgtt actagcactc acatggaaca
aatggeetet ageetggagg ettgetgaag getgtatget gteatgatea ataaggtate
                                                                   6240
cggcgttttg gcctctgact gacgccggat accattgatc atgacaggac acaaggcctg
                                                                   6300
ttactagcac tcacatggaa caaatggcct ctctagaat
                                                                   6339
                       moltype = DNA length = 6339
SEO ID NO: 17
FEATURE
                       Location/Qualifiers
                       1..6339
source
                       mol_type = other DNA
```

organism = synthetic construct

SEQUENCE: 17

aatcaacctc tggattacaa aatttgtgaa agattgactg gtattcttaa ctatgttgct 120 ccttttacgc tatgtggata cgctgcttta atgcctttgt atcatgctat tgcttcccgt atggetttea tttteteete ettgtataaa teetggttge tgtetettta tgaggagttg 180 tggcccgttg tcaggcaacg tggcgtggtg tgcactgtgt ttgctgacgc aacccccact 240 ggttggggca ttgccaccac ctgtcagctc ctttccggga ctttcgcttt ccccctccct 300 attgccacgg cggaactcat cgccgcctgc cttgcccgct gctggacagg ggctcggctg 360 ttgggcactg acaattccgt ggtgttgtcg gggaaatcat cgtcctttcc ttggctgctc 420 geetgtgttg ceacetggat tetgegeggg aegteettet getaegteee tteggeeete 480 aatccagegg accttectte eegeggeetg etgeeggete tgeggeetet teegegtett 540 cgccttcgcc ctcagacgag tcggatctcc ctttgggccg cctccccgcc taagcttatc gataccgtcg agatctaact tgtttattgc agcttataat ggttacaaat aaagcaatag catcacaaat ttcacaaata aagcattttt ttcactgcat tctagttgtg gtttgtccaa actcatcaat gtatcttatc atgtctggat ctcgacctcg actagagcat ggctacgtag ataagtagca tggcgggtta atcattaact acaaggaacc cctagtgatg gagttggcca ctccctctct gcgcgctcgc tcgctcactg aggccgggcg accaaaggtc gcccgacgcc cgggctttgc ccgggcggcc tcagtgagcg agcgagcgcg cagctggcgt aatagcgaag aggecegeae egategeeet teccaacagt tgegeageet gaatggegaa tggegattee gttgcaatgg ctggcggtaa tattgttctg gatattacca gcaaggccga tagtttgagt tettetaete aggeaagtga tgttattaet aateaaagaa gtattgegae aaeggttaat 1140 ttgcgtgatg gacagactct tttactcggt ggcctcactg attataaaaa cacttctcag 1200 gattctggcg taccgttcct gtctaaaatc cctttaatcg gcctcctgtt tagctcccgc 1260 tetgatteta acgaggaaag caegttatae gtgetegtea aageaaceat agtaegegee 1320 ctgtagcggc gcattaagcg cggcgggtgt ggtggttacg cgcagcgtga ccgctacact 1380 tgccagcgcc ctagcgcccg ctcctttcgc tttcttccct tcctttctcg ccacgttcgc 1440 eggettteee egteaagete taaategggg geteeettta gggtteegat ttagtgettt 1500 acggcacctc gaccccaaaa aacttgatta gggtgatggt tcacgtagtg ggccatcgcc 1560 ctgatagacg gtttttcgcc ctttgacgtt ggagtccacg ttctttaata gtggactctt 1620 gttccaaact ggaacaacac tcaaccctat ctcggtctat tcttttgatt tataagggat 1680 tttgccgatt tcggcctatt ggttaaaaaa tgagctgatt taacaaaaat ttaacgcgaa 1740 ttttaacaaa atattaacgt ttacaattta aatatttgct tatacaatct tcctgttttt 1800 ggggcttttc tgattatcaa ccggggtaca tatgattgac atgctagttt tacgattacc 1860 gttcatcgat tctcttgttt gctccagact ctcaggcaat gacctgatag cctttgtaga 1920 gaccteteaa aaatagetae eeteteegge atgaatttat eagetagaae ggttgaatat 1980 catattgatg gtgatttgac tgtctccggc ctttctcacc cgtttgaatc tttacctaca 2040 cattactcag gcattgcatt taaaatatat gagggttcta aaaattttta tccttgcgtt 2100 gaaataaagg cttctcccgc aaaagtatta cagggtcata atgtttttgg tacaaccgat 2160 ttagctttat gctctgaggc tttattgctt aattttgcta attctttgcc ttgcctgtat 2220 gatttattgg atgttggaat tcctgatgcg gtattttctc cttacgcatc tgtgcggtat 2280 ttcacaccgc atatggtgca ctctcagtac aatctgctct gatgccgcat agttaagcca 2340 geocegacae eegecaacae eegetgaege geeetgaegg gettgtetge teeeggeate 2400 cgcttacaga caagctgtga ccgtctccgg gagctgcatg tgtcagaggt tttcaccgtc 2460 atcaccgaaa cgcgcgagac gaaagggcct cgtgatacgc ctatttttat aggttaatgt 2520 catgataata atggtttctt agacgtcagg tggcactttt cggggaaatg tgcgcggaac 2580 ccctatttgt ttatttttct aaatacattc aaatatgtat ccgctcatga gacaataacc 2640 ctgataaatg cttcaataat attgaaaaag gaagagtatg agtattcaac atttccgtgt 2700 cgcccttatt cccttttttg cggcattttg ccttcctgtt tttgctcacc cagaaacgct 2760 ggtgaaagta aaagatgctg aagatcagtt gggtgcacga gtgggttaca tcgaactgga 2820 totcaacago ggtaagatoo ttgagagttt togoocogaa gaacgtttto caatgatgag 2880 cacttttaaa gttctgctat gtggcgcggt attatcccgt attgacgccg ggcaagagca 2940 actoggtogo ogcatacact attotoagaa tgacttggtt gagtactcac cagtcacaga 3000 aaagcatctt acggatggca tgacagtaag agaattatgc agtgctgcca taaccatgag 3060 tgataacact gcggccaact tacttctgac aacgatcgga ggaccgaagg agctaaccgc 3120 ttttttgcac aacatggggg atcatgtaac tcgccttgat cgttgggaac cggagctgaa 3180 tgaagccata ccaaacgacg agcgtgacac cacgatgcct gtagcaatgg caacaacgtt 3240 gcgcaaacta ttaactggcg aactacttac tctagcttcc cggcaacaat taatagactg gatggaggcg gataaagttg caggaccact tetgegeteg gecetteegg etggetggtt tattgctgat aaatctggag ccggtgagcg tgggtctcgc ggtatcattg cagcactggg gccagatggt aagccctccc gtatcgtagt tatctacacg acggggagtc aggcaactat ggatgaacga aatagacaga tcgctgagat aggtgcctca ctgattaagc attggtaact aaggatctag gtgaagatcc tttttgataa tctcatgacc aaaatccctt aacgtgagtt tteqttecae tqaqeqteaq acceeqtaqa aaaqatcaaa qqatettett qaqateettt ttttctgcgc gtaatctgct gcttgcaaac aaaaaaacca ccgctaccag cggtggtttg tttgccggat caagagctac caactctttt tccgaaggta actggcttca gcagagcgca 3840 gataccaaat actgtccttc tagtgtagcc gtagttaggc caccacttca agaactctgt 3900 agcaccgcct acatacctcg ctctgctaat cctgttacca gtggctgctg ccagtggcga 3960 taagtegtgt ettacegggt tggaeteaag acgatagtta eeggataagg egeageggte 4020 gggctgaacg gggggttcgt gcacacagcc cagcttggag cgaacgacct acaccgaact 4080 gagataccta cagcgtgagc tatgagaaag cgccacgctt cccgaaggga gaaaggcgga 4140 caggtatccg gtaagcggca gggtcggaac aggagagcgc acgagggagc ttccaggggg 4200 aaacgcctgg tatctttata gtcctgtcgg gtttcgccac ctctgacttg agcgtcgatt tttgtgatgc tcgtcagggg ggcggagcct atggaaaaac gccagcaacg cggccttttt 4320 acggttectg geettttget ggeettttge teacatgtte ttteetgegt tateecetga 4380 ttctgtggat aaccgtatta ccgcctttga gtgagctgat accgctcgcc gcagccgaac 4440 gaccgagcgc agcgagtcag tgagcgagga agcggaagag cgcccaatac gcaaaccgcc 4500 teteceegeg egttggeega tteattaatg eageagetge gegetegete geteaetgag 4560

| geegeeeggg | caaagcccgg | gcgtcgggcg | acctttggtc | gcccggcctc | agtgagcgag | 4620 |
|------------|------------|------------|------------|------------|------------|------|
| cgagcgcgca | gagagggagt | ggccaactcc | atcactaggg | gttccttgta | gttaatgatt | 4680 |
| aacccgccat | gctacttatc | tacgtagcca | tgctctagga | cattgattat | tgactagtgg | 4740 |
| agttccgcgt | tacataactt | acggtaaatg | gcccgcctgg | ctgaccgccc | aacgaccccc | 4800 |
| gcccattgac | gtcaataatg | acgtatgttc | ccatagtaac | gccaataggg | actttccatt | 4860 |
| gacgtcaatg | ggtggagtat | ttacggtaaa | ctgcccactt | ggcagtacat | caagtgtatc | 4920 |
| atatgccaag | tacgccccct | attgacgtca | atgacggtaa | atggcccgcc | tggcattatg | 4980 |
| cccagtacat | gaccttatgg | gactttccta | cttggcagta | catctacgta | ttagtcatcg | 5040 |
| ctattaccat | ggtcgaggtg | agccccacgt | tctgcttcac | tctccccatc | tccccccct | 5100 |
| ccccaccccc | aattttgtat | ttatttattt | tttaattatt | ttgtgcagcg | atgggggcgg | 5160 |
| 9999999999 | gggcgcgcgc | caggcggggc | ggggcggggc | gaggggcggg | gcggggcgag | 5220 |
| gcggagaggt | gcggcggcag | ccaatcagag | cggcgcgctc | cgaaagtttc | cttttatggc | 5280 |
| gaggcggcgg | cggcggcggc | cctataaaaa | gcgaagcgcg | cggcgggcgg | gagtcgctgc | 5340 |
| gcgctgcctt | cgccccgtgc | cccgctccgc | cgccgcctcg | cgccgcccgc | cccggctctg | 5400 |
| actgaccgcg | ttactaaaac | aggtaagtcc | ggcctccgcg | ccgggttttg | gcgcctcccg | 5460 |
| cgggcgcccc | cctcctcacg | gcgagcgctg | ccacgtcaga | cgaagggcgc | agcgagcgtc | 5520 |
| ctgatccttc | cgcccggacg | ctcaggacag | cggcccgctg | ctcataagac | tcggccttag | 5580 |
| aaccccagta | tcagcagaag | gacattttag | gacgggactt | gggtgactct | agggcactgg | 5640 |
| ttttctttcc | agagagcgga | acaggcgagg | aaaagtagtc | ccttctcggc | gattctgcgg | 5700 |
| agggatctcc | gtggggcggt | gaacgccgat | gatgcctcta | ctaaccatgt | tcatgttttc | 5760 |
| tttttttc | tacaggtcct | gggtgacgaa | cagggtaccg | ccaccatggc | caccggctct | 5820 |
| cgcacaagcc | tgctgctggc | tttcggactg | ctgtgcctgc | cttggctcca | ggagggctcc | 5880 |
| gccgctagca | tcgataccgt | cgctatgtgc | tggaggcttg | ctgaaggctg | tatgctgata | 5940 |
| aacaggcttg | ttcgcccagc | gttttggcct | ctgactgacg | ctgggcgaaa | gcctgtttat | 6000 |
| caggacacaa | ggcctgttac | tagcactcac | atggaacaaa | tggcctctag | cctggaggct | 6060 |
| tgctgaaggc | tgtatgctgt | catagctgca | atggtttctt | ccgttttggc | ctctgactga | 6120 |
| cggaagaaac | ctgcagctat | gacaggacac | aaggcctgtt | actagcactc | acatggaaca | 6180 |
| aatggcctct | agcctggagg | cttgctgaag | gctgtatgct | gaccacttcc | acaattcatg | 6240 |
| caccgttttg | gcctctgact | gacggtgcat | gaagtggaag | tggtcaggac | acaaggcctg | 6300 |
| ttactagcac | tcacatggaa | caaatggcct | ctctagaat | | | 6339 |
| | | | | | | |

The invention claimed is:

- 1. A composition that comprises a recombinant plasmid (RP) with a sequence of nucleotides that encodes a sequence of micro-interfering ribonucleic acid (miRNA) that binds to and degrades and/or inactivates messenger ribonucleic acid (mRNA) that encodes complement C1r and the sequence of nucleotides comprises SEQ ID NO. 3.
- 2. The composition of claim 1, wherein the sequence of nucleotides is encased in a protein coat, a lipid vesicle or a viral vector.
- 3. The composition of claim 2, wherein the viral vector is one of a double stranded DNA virus, a single stranded DNA

virus, a single stranded RNA virus, or a double stranded RNA virus.

- **4**. The composition of claim **3**, wherein the viral vector is an adeno-associated virus.
- **5**. A composition that comprises a recombinant plasmid (RP) with a sequence of nucleotides that encodes a sequence of micro-interfering ribonucleic acid (miRNA) that binds to and degrades and/or inactivates messenger ribonucleic acid (mRNA) that encodes complement C1r and the sequence of nucleotides comprises SEQ ID NO. 11.

* * * * *