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(54) Titre : VECTEURS ADENOVIRAUX SEROTYPE 34, ACIDES NUCLEIQUES ET VIRUS PRODUITS PAR CES MOYENS

(54) Title: ADENOVIRUS SEROTYPE 34 VECTORS, NUCLEIC ACIDS AND VIRUS PRODUCED THEREBY

(57) **Abrégé/Abstract:**

Adenoviral serotypes differ in their natural tropism. The various serotypes of adenovirus have been found to differ in at least their capsid proteins (e.g., penton-base and hexon proteins), proteins responsible for cell binding (e.g, fiber proteins), and proteins involved in adenovirus replication. This difference in tropism and capsid proteins among serotypes has led to the many research efforts aimed at redirecting the adenovirus tropism by modification of the capsid proteins. The present invention bypasses such requirement for capsid protein modification as it presents a recombinant, replication-defective adenovirus of serotype 34, a rare adenoviral serotype, and methods for generating the alternative, recombinant adenovirus. Additionally, means of employing the recombinant adenovirus for the delivery and expression of exogenous genes are provided.



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(54) Title: ADENOVIRUS SEROTYPE 34 VECTORS, NUCLEIC ACIDS AND VIRUS PRODUCED THEREBY

(57) Abstract: Adenoviral serotypes differ in their natural tropism. The various serotypes of adenovirus have been found to differ in at least their capsid proteins (*e.g.*, penton-base and hexon proteins), proteins responsible for cell binding (*e.g.*, fiber proteins), and proteins involved in adenovirus replication. This difference in tropism and capsid proteins among serotypes has led to the many research efforts aimed at redirecting the adenovirus tropism by modification of the capsid proteins. The present invention bypasses such requirement for capsid protein modification as it presents a recombinant, replication-defective adenovirus of serotype 34, a rare adenoviral serotype, and methods for generating the alternative, recombinant adenovirus. Additionally, means of employing the recombinant adenovirus for the delivery and expression of exogenous genes are provided.

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## TITLE OF THE INVENTION

ADENOVIRUS SEROTYPE 34 VECTORS, NUCLEIC ACIDS AND VIRUS PRODUCED THEREBY

## 5 CROSS-REFERENCE TO RELATED APPLICATIONS

This application claims the benefit of U.S. provisional application serial no. 60/458,825, filed on March 28, 2003.

## BACKGROUND OF THE INVENTION

10 Adenoviruses are nonenveloped, icosahedral viruses that have been identified in several avian and mammalian hosts; Horne *et al.*, 1959 *J. Mol. Biol.* 1:84-86; Horwitz, 1990 In *Virology*, eds. B.N. Fields and D.M. Knipe, pps. 1679-1721. The first human adenoviruses (Ads) were isolated over four decades ago. Since then, over 100 distinct adenoviral serotypes have been isolated which infect various mammalian species, 51 of which are of human origin; 15 Straus, 1984, In *The Adenoviruses*, ed. H. Ginsberg, pps. 451-498, New York:Plenus Press; Hierholzer *et al.*, 1988 *J. Infect. Dis.* 158:804-813; Schnurr and Dondero, 1993, *Intervirology*; 36:79-83; Jong *et al.*, 1999 *J Clin Microbiol.*, 37:3940-5. The human serotypes have been categorized into six subgenera (A-F) based on a number of biological, chemical, immunological and structural criteria which include hemagglutination properties of rat and rhesus monkey 20 erythrocytes, DNA homology, restriction enzyme cleavage patterns, percentage G+C content and oncogenicity; Straus, *supra*; Horwitz, *supra*.

The adenovirus genome is very well characterized. It consists of a linear double-stranded DNA molecule of approximately 36,000 base pairs, and despite the existence of several distinct serotypes, there is some general conservation in the overall organization of the 25 adenoviral genome with specific functions being similarly positioned.

Adenovirus has been a very attractive target for delivery of exogenous genes. The biology of adenoviruses is very well understood. Adenovirus has not been found to be associated with severe human pathology in immuno-competent individuals. The virus is extremely efficient in introducing its DNA into the host cell and is able to infect a wide variety 30 of cells. Furthermore, the virus can be produced at high virus titers in large quantities. In addition, the virus can be rendered replication defective by deletion of the essential early-region 1 (E1) of the viral genome; Brody et al, 1994 *Ann N Y Acad Sci.*, 716:90-101.

Replication-defective adenovirus vectors have been used extensively as gene transfer vectors for vaccine and gene therapy purposes. These vectors are propagated in cell



lines that provide E1 gene products *in trans*. Supplementation of the essential E1 gene products *in trans* is very effective when the vectors are from the same or a very similar serotype. E1-deleted group C serotypes (Ad1, Ad2, Ad5 and Ad6), for instance, grow well in 293 or PER.C6 cells which contain and express the Ad5 E1 region. However, the Ad5 E1 sequences in 293 or PER.C6 cells do not fully complement the replication of all serotypes other than group C. This is perhaps due to the inability of the Ad5 (group C) E1B 55K gene product to functionally interact with the E4 gene product(s) of the non-group C serotypes. Although the interaction is conserved within members of the same subgroup, it has not been found to be well conserved between subgroups. In order to successfully and efficiently rescue recombinant adenovirus of alternative, non-group C serotypes, a cell line expressing the E1 region of the serotype of interest would have to be generated. Alternatively, available Ad5E1-expressing cell lines could be modified to express Ad5E4 (or Orf6) in addition to Ad5E1. These additional, sometimes tedious and daunting tasks, impeded the production of recombinant, non-group C adenoviral vectors.

An efficient means for the propagation and rescue of alternative serotypes in an Ad5 E1-expressing cell line (such as PER.C6 or 293) was disclosed in pending U.S. provisional application (Serial No. 60/405,182, filed August 22, 2002). This method involves the incorporation of a critical E4 region into the adenovirus to be propagated. The critical E4 region is native to a virus of the same or highly similar serotype as that of the E1 gene product(s), particularly the E1B 55K region, of the complementing cell line, and comprises, in the least, nucleic acid encoding E4 Orf6.

Presently, two well-characterized adenovirus serotypes from subgroup C, Ad5 and Ad2, are the most widely used gene delivery vectors. There is a need to develop alternate Ad serotypes as gene transfer vectors since neutralizing antibodies in the general population may limit primary dosing or redosing with the same serotype. The prevalence of neutralizing antibody can vary from serotype to serotype. Neutralizing antibodies to some serotypes such as Ad5 are common, while antibodies to others are relatively rare. Alternate serotypes, furthermore, possess alternate tropisms which may lead to the elicitation of superior immune responses when used for vaccine or gene therapy purposes.

Adenovirus serotype 34, a subgroup B adenovirus, was originally isolated in 1972 and established as a recognized reference strain in 1975 (J.C. Hierholzer *et al.*, 1975 *J. Clin. Microbiol.* 1:366-376). Its antigenic relationship to 46 other human adenoviruses determined in reference horse antisera has been discussed; J.C. Hierholzer *et al.*, 1991 *Arch. Virol.* 121:179-197. Partial sequence information is available for Ad34. There have been several disclosures relating to Ad34 hexon sequences. The complete sequence of Ad34 hexon with some 5' and 3'



flanking sequence (3358 bp) was deposited in GenBank (Accession No. AB052911) by Mukouyama. A partial sequence of Ad34 hexon (1449 bp) was disclosed in Takeuchi *et al.*, 1999 *J. Clin. Microbiol.* 37:3392-3394, and GenBank (Accession No. AB018426). A partial sequence of Ad34 hexon (253 bp) was disclosed in Allard *et al.*, 2001 *J. Clin. Microbiol.* 39: 498-505, and deposited in GenBank (Accession No. AF161573). Perera and Cardoso deposited two partial sequences of Ad34 hexon (571bp and 301 bp) with GenBank (Accession Nos. AJ272610 and AJ250786). Sequence for the Ad34 fiber gene was deposited by Arun, Mukouyama and Inada with GenBank (Accession No. AB073168). The sequence of the virus associated RNA region (VA RNA1 & 2) for Ad34 (162 bp) was disclosed by Kidd *et al.*, 1995 *Virology* 207:32-45, and GenBank (Accession No. U10677). Moreover, the sequence of the virus associated RNA region for Ad34 and partial sequence for the pre-terminal protein and 52/55K proteins (354 bp) was disclosed in Ma & Matthews, 1996 *J. Virol.* 70: 5083-99, and GenBank (Accession No. U52571). Adhikary, Mukouyama and Inada disclosed the sequence of the Ad34 genes for L4 100kDa, L4pVIII, E3 12.3kDa, E3 14.9kDa, E3 gp18.5kDa, E3 20.3kDa, E3 20.5kDa, E3 10.2kDa, E3 15.2kDa1, E3 15.2kDa2, and partial fiber sequence (4828 bp) and deposited the sequence with GenBank (Accession No. AB079724). The sequence of the right end of the viral genome (1038 bp) was disclosed in Chen & Horwitz, 1990 *Virology* 179:567-75, and GenBank (Accession No. M62712).

The fields of vaccines and gene therapy would greatly benefit from additional knowledge concerning alternative adenoviral serotypes, particularly those serotypes such as Ad34 which are not well represented in the human population. Of particular interest are recombinant adenoviral vectors based on alternative adenoviral serotypes, and means of obtaining such recombinant adenoviral vectors. This need in the art is met with the disclosure of the present application related to recombinant adenoviral vectors based on adenoviral serotype 34.

#### SUMMARY OF THE INVENTION

The present invention relates to recombinant, replication-deficient adenovirus vectors of serotype 34, a rare adenoviral serotype, and methods for generating the recombinant adenovirus based on the alternative serotype. Additionally, means of employing the recombinant adenovirus for the delivery and expression of exogenous genes are provided. The invention, thus, encompasses recombinant, replication-defective adenoviral vectors of serotype 34 which comprise one or more transgenes operatively linked to regulatory sequences which promote effective expression of the respective transgene(s). Host administration of such recombinant



adenovirus serotype 34 vectors, whether administered alone or in a combined modality and/or prime boost regimen, results in the efficient expression of the incorporated transgene and effectively induces an immune response capable of specifically recognizing the particular antigen administered (*e.g.*, HIV). Furthermore, the recombinant virus should evade pre-existing immunity to adenovirus serotypes which are more commonly encountered in the human population (*e.g.*, Ad5 and Ad2). The disclosed methods, thus, present an enhanced means for inducing an immune response against a particular antigen of interest (*e.g.*, HIV). Accordingly, the resultant immune response should offer a prophylactic advantage to previously uninfected individuals and/or provide a therapeutic effect by reducing viral load levels within an infected individual, thus prolonging the asymptomatic phase of infection.

#### BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 illustrates the homologous recombination scheme utilized to recover pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

Figure 2 illustrates the homologous recombination scheme utilized to recover pMRKAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

Figures 3A-1 to 3A-9 illustrate a nucleic acid sequence for wild-type adenovirus serotype 34 (SEQ ID NO: 1). The ATCC product number for Ad34 is VR-716.

Figure 4 illustrates the time course of SEAP expression using MRKAd5 and Ad34 vectors in rhesus macaques. Data represent cohort geometric means.

Figure 5 illustrates, in tabular format, T cell responses induced using MRKAd5 and Ad34 vectors expressing HIV-1 gag. Data are expressed in numbers of spot-forming cells per million PBMC (SFC/10<sup>6</sup> PBMC). "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

Figure 6 illustrates, in tabular format, the levels of CD4<sup>+</sup> and CD8<sup>+</sup> Gag-specific T cells in Ad34-immunized macaques at week 12. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

Figure 7 illustrates the nucleic acid sequence (SEQ ID NO: 3) of the optimized human HIV-1 gag open reading frame.

Figure 8 illustrates the nucleic acid sequence encoding the gag expression cassette (SEQ ID NO: 4). The various regions of the figure are as follows: (1) a first underlined segment of nucleic acid sequence encoding the immediate early gene promoter region from human cytomegalovirus; (2) a first segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; (3) a region in caps which contains the

coding sequence of HIV-1 gag; (4) a second segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; and (5) a second underlined segment, this segment containing nucleic acid sequence encoding a bovine growth hormone polyadenylation signal sequence.

5 Figure 9 illustrates the nucleic acid sequence encoding the SEAP expression cassette (SEQ ID NO: 5). The various regions of the figure are as follows: (1) a first underlined segment of nucleic acid sequence encoding the immediate early gene promoter region from human cytomegalovirus; (2) a first segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; (3) a region in caps which  
10 contains the coding sequence of the human placental SEAP gene; (4) a second segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; and (5) a second underlined segment, this segment containing nucleic acid sequence encoding a bovine growth hormone polyadenylation signal sequence.

15 Figures 10A-1 to 10A-47 illustrate the nucleotide sequence of the pMRKAd5HIV-1gag vector (SEQ ID NO: 6 [coding] and SEQ ID NO: 7 [non-coding]).

Figures 11A-1 to 11A-10 illustrate a nucleic acid sequence for wild-type adenovirus serotype 35 (SEQ ID NO: 13). The ATCC product number for Ad35 is VR-718.

20 Figure 12 illustrates, in tabular format, T cell responses induced using a heterologous Ad34 prime/Ad35 boost regimen in macaques. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

Figure 13 illustrates, in tabular format, the levels of CD4+ and CD8+ Gag-specific T cells in Ad34 primed/Ad35 boosted macaques at week 28. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.



## DETAILED DESCRIPTION OF THE INVENTION

Rare adenoviral serotypes possess an inherent advantage over the more commonly exploited adenoviral serotypes (for instance, adenoviral serotypes 2 and 5) since preexisting immunity is unlikely to limit their efficient delivery and expression of exogenous genes to their target site. Different adenoviral serotypes also exhibit distinct tropisms by reason of their varying capsid structure and, thus, present the potential for targeting different tissues and possibly leading to the elicitation of superior immune responses when used for vaccine or gene therapy purposes. These rare adenoviral serotypes when rendered replication-defective, however, can be difficult to propagate and rescue in currently available adenoviral propagation cell lines.

Applicants have recently managed to successfully rescue and propagate one such rare, replication-defective alternative serotype, adenovirus serotype 34, a subgroup B adenovirus, and herein demonstrate the effective functioning of the adenovirus in the delivery and expression of exogenous transgenes.

Accordingly, the present invention relates to a recombinant adenoviral vector of serotype 34 suitable for use in gene therapy or vaccination protocols. The nucleic acid sequence for wild-type adenovirus serotype 34 (SEQ ID NO: 1) is illustrated in Figures 3A-1 to 3A-9, although any functional homologue or different strain of adenovirus serotype 34 can be utilized in accordance with the methods of the present invention, as one of ordinary skill in the art will appreciate. Ad34 sequence has been noted to differ in a few regions. The following sites are just a sampling of sequence variation that can be found in Ad34: (1) around base pair 10640 of SEQ ID NO: 1, a series of thirteen ("13") rather than twelve ("12") "T"s follow the sequence gtgagtctcta (SEQ ID NO: 8); (2) around base pair 15372 of SEQ ID NO: 1, a series of fifteen ("15") or seventeen ("17") rather than sixteen ("16") "A"s follow the sequence ccgcactttct (SEQ ID NO: 9); (3) around base pair 17325 of SEQ ID NO: 1, a series of thirteen ("13") rather than twelve ("12") "A"s follow the sequence attgacattgg (SEQ ID NO: 10); and (4) around base pair 25717 of SEQ ID NO: 1, the sequence cagtctggagga (SEQ ID NO: 11) following the sequence ggagga (SEQ ID NO: 12) is deleted. Adenovirus serotypes have been distinguished through a number of art-appreciated biological, chemical, immunological and structural criteria which include hemagglutination properties of rat and rhesus monkey erythrocytes, DNA homology, restriction enzyme cleavage patterns, percentage G+C content and oncogenicity; Straus, *supra*; Horwitz, *supra*. A given serotype can be identified by a number of methods including restriction mapping of viral DNA; analyzing the mobility of viral DNA; analyzing the mobility of virion polypeptides on SDS-polyacrylamide gels following electrophoresis; comparison of sequence



information to known sequence particularly from capsid genes (*e.g.*, hexon) which contain sequences that define a serotype; and comparing a sequence with reference sera for a particular serotype available from the ATCC. Classification of adenovirus serotypes by SDS-PAGE has been discussed in Wadell *et al.*, 1980 *Ann. N.Y. Acad. Sci.* 354:16-42. Classification of  
5 adenovirus serotypes by restriction mapping has been discussed in Wadell *et al.*, *Current Topics in Microbiology and Immunology* 110:191-220. Adenovirus serotype 34, a subgroup B adenovirus, was originally isolated in 1972 and was established as a recognized reference strain in 1975 (J.C. Hierholzer *et al.*, 1975 *J. Clin. Microbiol.* 1:366-376). Its antigenic relationship to  
10 46 other human adenoviruses determined in reference horse antisera has been discussed in the art; J.C. Hierholzer *et al.*, 1991 *Arch. Virol.* 121:179-197.

Adenovirus serotype 34 vectors in accordance with the present invention are at least partially deleted in E1 and devoid (or essentially devoid) of E1 activity, rendering the vector incapable of replication in the intended host. Preferably, the E1 region is completely deleted or inactivated. The adenoviruses may contain additional deletions in E3, and other early  
15 regions, albeit in situations where E2 and/or E4 is deleted, E2 and/or E4 complementing cell lines may be required to generate recombinant, replication-defective adenoviral vectors.

Adenoviral vectors of use in the methods of the present invention can be constructed using well known techniques, such as those reviewed in Hitt *et al.*, 1997 "Human  
20 Adenovirus Vectors for Gene Transfer into Mammalian Cells" *Advances in Pharmacology* 40:137-206, which is hereby incorporated by reference. Often, a plasmid or shuttle vector containing the heterologous nucleic acid of interest is generated which comprises sequence homologous to the specific adenovirus of interest. The shuttle vector and viral DNA or second plasmid containing the cloned viral DNA are then co-transfected into a host cell where  
25 homologous recombination occurs and results in the incorporation of the heterologous nucleic acid into the viral nucleic acid. Preferred shuttle vectors and cloned viral genomes contain adenoviral and plasmid portions. For shuttle vectors used in the construction of replication-defective vectors, the adenoviral portion typically contains non-functional or deleted E1 and E3 regions and the gene expression cassette, flanked by convenient restriction sites. The plasmid portion of the shuttle vector typically contains an antibiotic resistance marker under the  
30 transcriptional control of a prokaryotic promoter. Ampicillin resistance genes, neomycin resistance genes and other pharmaceutically acceptable antibiotic resistance markers may be used. To aid in the high level production of the nucleic acid by fermentation in prokaryotic organisms, it is advantageous for the shuttle vector to contain a prokaryotic origin of replication and be of high copy number. A number of commercially available prokaryotic cloning vectors



provide these benefits. Non-essential DNA sequences are, preferably removed. It is also preferable that the vectors not be able to replicate in eukaryotic cells. This minimizes the risk of integration of nucleic acid vaccine sequences into the recipients' genome. Tissue-specific promoters or enhancers may be used whenever it is desirable to limit expression of the nucleic acid to a particular tissue type.

Homologous recombination of the shuttle vector and wild-type adenovirus 34 viral DNA (Ad34 backbone vector) results in the generation of adenoviral pre-plasmids (see, for instance, pAd34ΔE1ΔE4Ad5Orf6, pMRKAd34ΔE1ΔE4Ad5Orf6, pAd34ΔE1gagΔE4Ad5Orf6, and pAd34ΔE1SEAPΔE4Ad5Orf6). Upon linearization, the pre-plasmids are capable of replication in PER.C6<sup>®</sup> cells or alternative E1-complementing cell lines. Infected cells and media can then be harvested once viral replication is complete.

A packaging cell will generally be needed in order to produce sufficient amount of adenovirus. The packaging cell should contain elements which are necessary for the production of the specific adenovirus of interest. It is preferable that the packaging cell and the vector not contain overlapping elements which could lead to replication competent virus by recombination. Specific examples of cells which are suitable for the propagation of recombinant Ad34 E1-deleted vectors express the early region 1 (E1) of adenovirus 34 or another group B serotype. Alternatively, propagation cell lines can be used which express adenoviral E1 and E4 regions (particularly, E4 open reading frame 6 ("ORF6")) which are derived from the same serotype but different subgroup than Ad34 (*e.g.*, Ad5 E1 and E4); *see, e.g.*, Abrahamsen *et al.*, 1997 *J. Virol.* 8946-8951, and U.S. Patent No. 5,849,561. Additionally, a cell line could be used that expresses E1B from Ad34 in addition to (1) E1A or (2) E1A and E1B from a serotype of a different subgroup. In copending U.S. provisional application serial no. 60/405,182, filed August 22, 2002, a strategy was disclosed for the efficient propagation and rescue of alternative adenoviral serotypes. The method is based on incorporating, into the genome of the adenovirus vector, an E4 region (or portion thereof including E4 ORF6) of the same or highly similar serotype as that of the E1 gene product(s), particularly E1B, being expressed by the complementing cell line. Examples 1-4 demonstrate the viability of such a method through the incorporation of an Ad5E4 region and its propagation in PER.C6 cells (which cells express Ad5E1). The wildtype adenovirus serotype 5 sequence is known and described in the art; *see* Chroboczek *et al.*, 1992 *J. Virol.* 186:280, which is hereby incorporated by reference. Placement of the E4 region or ORF6-containing portion is not critical. The critical step is making sure that either a promoter is supplied or the gene is strategically placed so that it runs off a promoter native to the vector (*e.g.*, such as the E4 promoter). The native E4 region of the vector can be



replaced, deleted or left intact. This method is, thus, suitable for use in the propagation and rescue of the adenoviral vectors of the present invention.

Typically, propagation cells are human cells derived from the retina or kidney, although any cell line capable of expressing the appropriate E1 and/or E4 region(s) can be  
5 utilized in the present invention. Embryonal cells such as amniocytes have been shown to be particularly suited for the generation of E1 complementing cell lines. Several cell lines are available. These include but are not limited to the known cell lines PER.C6 (ECACC deposit number 96022940), 911, 293, and E1 A549.

The present invention encompasses methods for producing a recombinant,  
10 replication-defective adenovirus of serotype 34 in an adenoviral E1-complementing cell line, comprising transfecting a recombinant, replication-defective adenoviral vector of serotype 34 in an adenoviral E1-complementing cell and allowing for the production of viral particles. The viral particles so produced form another aspect of the present invention. Host cells comprising the recombinant, replication-defective adenoviral serotype 34 vectors of the present invention  
15 form yet another aspect of the present invention; host cells being defined as a population of cells not including a transgenic human being. Recombinant, replication-defective adenovirus harvested in accordance with the methods of the present invention are encompassed herein as well. This harvested material may be purified, formulated and stored prior to host administration.

20 Adenoviral vectors in accordance with the present invention are very well suited to effectuate expression of desired proteins, especially in situations where an individual's immune response effectively prevents administration or readministration via the more commonly employed adenoviral serotypes. Accordingly, specific embodiments of the present invention are recombinant, replication-defective adenoviral vectors of serotype 34 which comprise a  
25 heterologous nucleic acid of interest. The nucleic acid of interest can be a gene, or a functional part of a gene. The nucleic acid can be DNA and/or RNA, can be double or single stranded, and can exist in the form of an expression cassette. The nucleic acid can be inserted in an E1 parallel (transcribed 5' to 3') or anti-parallel (transcribed in a 3' to 5' direction relative to the vector backbone) orientation. The nucleic acid can be codon-optimized for expression in the desired  
30 host (*e.g.*, a mammalian host). The heterologous nucleic acid can be in the form of an expression cassette. A gene expression cassette will typically contain (a) nucleic acid encoding a protein or antigen of interest; (b) a heterologous promoter operatively linked to the nucleic acid encoding the protein; and (c) a transcription termination signal.



In specific embodiments, the heterologous promoter is recognized by an eukaryotic RNA polymerase. One example of a promoter suitable for use in the present invention is the immediate early human cytomegalovirus promoter (Chapman *et al.*, 1991 *Nucl. Acids Res.* 19:3979-3986). Further examples of promoters that can be used in the present invention are the strong immunoglobulin promoter, the EF1 alpha promoter, the murine CMV promoter, the Rous Sarcoma Virus promoter, the SV40 early/late promoters and the beta actin promoter, albeit those of skill in the art can appreciate that any promoter capable of effecting expression in the intended host can be used in accordance with the methods of the present invention. The promoter may comprise a regulatable sequence such as the Tet operator sequence. Sequences such as these that offer the potential for regulation of transcription and expression are useful in instances where repression of gene transcription is sought. The adenoviral gene expression cassette may comprise a transcription termination sequence; specific embodiments of which are the bovine growth hormone termination/polyadenylation signal (bGHpA) or the short synthetic polyA signal (SPA) of 50 nucleotides in length defined as follows: AATAAAAGATCTTTATTTTCATTAGATCTGTGTGTT-GGTTTTTTGTGTG (SEQ ID NO:2). A leader or signal peptide may also be incorporated into the transgene. In specific embodiments, the leader is derived from the tissue-specific plasminogen activator protein, tPA.

Heterologous nucleic acids of interest are genes (or their functional counterparts) which encode immunogenic and/or therapeutic proteins. Preferred therapeutic proteins are those which elicit some measurable therapeutic benefit in the individual host upon administration. Preferred immunogenic proteins are any proteins which are capable of eliciting an immune response in an individual. Applicants have exemplified the delivery of a representative immunogenic protein (HIV gag) in the present specification in non-human primates (rhesus macaques), albeit any gene encoding a therapeutic or immunogenic protein can be used in accordance with the methods disclosed herein. The adenovirus serotype 34 vectors were found to induce significant levels of gag-specific T cells; Figure 5. Moreover, the results indicated that immunization with the disclosed vectors was able to elicit both HIV-specific CD4+ and CD8+ T cells; Figure 6.

An aspect of the present invention, therefore, relates to adenovirus serotype 34-based vectors carrying an HIV transgene. In these embodiments, nucleic acid encoding any HIV antigen may be utilized (specific examples of which include gag, pol, nef, gp160, gp41, gp120, tat, and rev, including derivatives of the aforementioned genes). The embodiments exemplified herein employ nucleic acid encoding a codon-optimized p55 gag antigen; *see* Figure 7 (SEQ ID NO: 3). Codon-optimized HIV-1 env genes are disclosed in PCT International Applications



PCT/US97/02294 and PCT/US97/10517, published August 28, 1997 (WO 97/31115) and December 24, 1997, respectively. Codon-optimized HIV-1 pol genes are disclosed in U.S. Application Serial No. 09/745,221, filed December 21, 2000 and PCT International Application PCT/US00/34724, also filed December 21, 2000. Codon-optimized HIV-1 nef genes are  
5 disclosed in U.S. Application Serial No. 09/738,782, filed December 15, 2000 and PCT International Application PCT/US00/34162, also filed December 15, 2000.

In this specific embodiment of a recombinant, replication-defective Ad34 vector comprising an HIV-1 gene, the gene may be derived from HIV-1 strain CAM-1; Myers et al, eds. "Human Retroviruses and AIDS: 1995, IIA3-IIA19, which is hereby incorporated by reference.  
10 This gene closely resembles the consensus amino acid sequence for the clade B (North American/European) sequence. HIV gene sequence(s) may be based on various clades of HIV-1; specific examples of which are Clades B and C. Sequences for genes of many HIV strains are publicly available from GenBank and primary, field isolates of HIV are available from the National Institute of Allergy and Infectious Diseases (NIAID) which has contracted with Quality  
15 Biological (Gaithersburg, MD) to make these strains available. Strains are also available from the World Health Organization (WHO), Geneva Switzerland. It is well within the purview of the skilled artisan to choose an appropriate nucleotide sequence which encodes a specific HIV antigen, or immunologically relevant portion or modification thereof. "Immunologically relevant" as defined herein means (1) with regard to a viral antigen, that the protein is capable,  
20 upon administration, of eliciting a measurable immune response within an individual sufficient to retard the propagation and/or spread of the virus and/or to reduce the viral load present within the individual; or (2) with regards to a nucleotide sequence, that the sequence is capable of encoding for a protein capable of the above.

The present invention encompasses methods for (1) effectuating a therapeutic  
25 response in an individual and (2) generating an immune response (including a cellular-mediated immune response) comprising administering to an individual a recombinant adenovirus serotype 34 vector in accordance with the present invention. One aspect of the present invention are methods for generating an enhanced immune response against one or more antigens (bacterial, viral (*e.g.*, HIV), or other (*e.g.*, cancer)) which comprise the administration of a recombinant  
30 adenovirus serotype 34 vehicle expressing the antigen of interest. Administration of recombinant Ad34 vectors in this manner provides for improved cellular-mediated immune responses, particularly where there is pre-existing immunity in a given host to the more well-represented adenovirus serotypes (*e.g.*, Ad2 and Ad5). An effect of the improved vaccine administration methods should be a lower transmission rate to (or occurrence rate in) previously



uninfected individuals (*i.e.*, prophylactic applications) and/or a reduction in the levels of virus/bacteria/foreign agent within an infected individual (*i.e.*, therapeutic applications). As relates to HIV indications, an effect of the improved vaccine administration methods should be a lower transmission rate to previously uninfected individuals (*i.e.*, prophylactic applications) and/or a reduction in the levels of viral loads within an infected individual (*i.e.*, therapeutic applications) so as to prolong the asymptomatic phase of HIV infection. Administration, intracellular delivery and expression of the recombinant Ad34 vectors elicits a host CTL and Th response.

Accordingly, the present invention relates to methodology regarding administration of the recombinant Ad34 viral vectors (or immunogenic compositions thereof, herein termed vaccines) to provide effective immunoprophylaxis, to prevent establishment of an infection following exposure to the viral (for instance, HIV), bacterial or other agent, or as a post-infection therapeutic vaccine to mitigate infection to result in the establishment of a lower virus/bacteria/other load with beneficial long term consequences.

The recombinant adenovirus serotype 34 vectors of the present invention may be administered alone, or as part of a prime/boost administration regimen. A priming dose(s) of at least one antigen (*e.g.*, an HIV antigen) is first delivered with a recombinant adenoviral vector. This dose effectively primes the immune response so that, upon subsequent identification of the antigen(s) in the circulating immune system, the immune response is capable of immediately recognizing and responding to the antigen(s) within the host. The priming dose(s) is then followed with a boosting dose comprising a recombinant adenoviral vector containing at least one gene encoding the antigen. A mixed modality prime and boost inoculation scheme will result in an enhanced immune response, particularly where there is pre-existing anti-vector immunity. Prime-boost administrations typically involve priming the subject (by viral vector, plasmid, protein, *etc.*) at least one time, allowing a predetermined length of time to pass, and then boosting (by viral vector, plasmid, protein, *etc.*). Multiple primings, typically 1-4, are usually employed, although more may be used. The length of time between priming and boost may typically vary from about four months to a year, albeit other time frames may be used as one of ordinary skill in the art will appreciate.

In addition to a single protein or antigen of interest being delivered by the recombinant, replication-defective adenovirus serotype 34 vectors of the present invention, two or more proteins or antigens can be delivered either via separate vehicles or delivered *via* the same vehicle. Multiple genes/functional equivalents may be ligated into a proper shuttle plasmid for generation of a pre-adenoviral plasmid comprising multiple open reading frames. Open



reading frames for the multiple genes/functional equivalents can be operatively linked to distinct promoters and transcription termination sequences. In other embodiments, the open reading frames may be operatively linked to a single promoter, with the open reading frames operatively linked by an internal ribosome entry sequence (IRES; as disclosed in WO 95/24485), or suitable alternative allowing for transcription of the multiple open reading frames to run off of a single promoter. In certain embodiments, the open reading frames may be fused together by stepwise PCR or suitable alternative methodology for fusing together two open reading frames. Due to consideration must be given, however, to the effective packaging limitations of the viral vehicle. Adenovirus type 5, for instance, has been shown to exhibit an upper cloning capacity limit of approximately 105% of the wildtype Ad5 sequence.

Prime-boost regimens can employ different adenoviral serotypes. One example of such a protocol would be a priming dose(s) comprising a recombinant adenoviral vector of a first serotype followed by a boosting dose comprising a recombinant adenoviral vector of a second and different serotype; *see*, for instance, Example 6 and Figures 12 and 13. Therein, a cohort of monkeys was given two doses of an Ad34-based HIV gag vector at weeks 0 and 4, and boosted at week 24 with an Ad35-based HIV gag vector. Administration of the Ad35-based vector resulted in about a 3-fold enhancement in T cell responses when compared to the levels at the time of the booster. In an alternative embodiment, the priming dose can comprise a mixture of separate adenoviral vehicles each comprising a gene encoding for a different protein/antigen. In such a case, the boosting dose would also comprise a mixture of vectors each comprising a gene encoding for a separate protein/antigen, provided that the boosting dose(s) administers recombinant viral vectors comprising genetic material encoding for the same or similar set of antigens that were delivered in the priming dose(s). These multiple gene/vector administration modalities can further be combined. It is further within the scope of the present invention to embark on combined modality regimes which include multiple but distinct components from a specific antigen.

Compositions, including vaccine compositions, comprising the adenoviral vectors of the present invention are an important aspect of the present invention. These compositions can be administered to mammalian hosts, preferably human hosts, in either a prophylactic or therapeutic setting. Potential hosts/vaccinees include but are not limited to primates and especially humans and non-human primates, and include any non-human mammal of commercial or domestic veterinary importance. Compositions comprising recombinant adenoviral serotype 34 vectors may be administered alone or in combination with other viral- or non-viral-based DNA/protein vaccines. They also may be administered as part of a broader treatment regimen.



The present invention encompasses those situations as well where the disclosed recombinant adenoviral serotype 34 vectors are administered in conjunction with other therapies; for example, HAART therapy (in the case of a recombinant HIV vector).

Compositions comprising the recombinant viral vectors may contain  
5 physiologically acceptable components, such as buffer, normal saline or phosphate buffered saline, sucrose, other salts and polysorbate. In certain embodiments, the formulation has: 2.5-10 mM TRIS buffer, preferably about 5 mM TRIS buffer; 25-100 mM NaCl, preferably about 75 mM NaCl; 2.5-10% sucrose, preferably about 5% sucrose; 0.01 -2 mM MgCl<sub>2</sub>; and 0.001%-  
10 0.01% polysorbate 80 (plant derived). The pH should range from about 7.0-9.0, preferably about 8.0. One skilled in the art will appreciate that other conventional vaccine excipients may also be used in the formulation. In specific embodiments, the formulation contains 5mM TRIS, 75 mM NaCl, 5% sucrose, 1mM MgCl<sub>2</sub>, 0.005% polysorbate 80 at pH 8.0. This has a pH and divalent cation composition which is near the optimum for Ad5 and Ad6 stability and minimizes the potential for adsorption of virus to a glass surface. It does not cause tissue irritation upon  
15 intramuscular injection. It is preferably frozen until use.

The amount of viral particles in the vaccine composition to be introduced into a vaccine recipient will depend on the strength of the transcriptional and translational promoters used and on the immunogenicity of the expressed gene product. In general, an immunologically or prophylactically effective dose of  $1 \times 10^7$  to  $1 \times 10^{12}$  particles and preferably about  $1 \times 10^{10}$  to  
20  $1 \times 10^{11}$  particles is administered directly into muscle tissue. Subcutaneous injection, intradermal introduction, impression through the skin, and other modes of administration such as intraperitoneal, intravenous, or inhalation delivery are also contemplated. Parenteral administration, such as intravenous, intramuscular, subcutaneous or other means of administration of interleukin-12 protein, concurrently with or subsequent to parenteral  
25 introduction of the vaccine compositions of this invention is also advantageous.

The following non-limiting Examples are presented to better illustrate the workings of the invention.

### 30 *Example 1*

#### Construction of pAd34ΔE1ΔE4Ad5Orf6

To generate an E1- Ad34 based vector that can propagate in existing group C/Ad5 E1 complementing cell lines (293, PER.C6), Ad5 Orf6 was inserted in place of the native E4 region. To construct the Ad34 pre-Adenovirus plasmid, advantage was taken of the sequence



homology between Ad34 and Ad35. Cotransformation of BJ 5183 bacteria with purified wild-type Ad34 viral DNA and the appropriately constructed Ad35 ITR cassette resulted in the circularization of the viral genome by homologous recombination. The construction of the pre-Ad plasmid based on Ad34, is outlined below:

5                   To construct pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6 (An Ad34 pre-Ad plasmid containing an E1 deletion and an E4 deletion substituted with Ad5 Orf6), we utilized an Ad35 ITR cassette. We anticipated that sequence homology between Ad34 and Ad35 would allow homologous recombination to occur. The Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31913 and bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end  
10 of the Ad35 genome (*see* Figures 11A-1 to 11A-10) separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. The four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad35 bp 31913 and 34419 generating pNEBAd35-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used  
15 commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique *Swa* I restriction site located in the deletion and an E4 deletion from Ad35 bp 31914 to 34418 into which Ad5 Orf6 was introduced in an E4 parallel  
20 orientation. In this construct Ad5Orf6 expression is driven by the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 31913 and bp 3403 to 3886) in the ITR cassette provided regions of homology with the purified Ad34 viral DNA in which bacterial recombination could occur following cotransformation into BJ 5183 bacteria (Figure 1). The ITR cassette was also designed to contain unique restriction enzyme sites (*Pme*I) located at the end of the viral ITR's  
25 so that digestion would release the recombinant Ad34 genome from the plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

### *Example 2*

#### 30 Rescue of pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6 into Virus

In order to determine if pre-adenovirus plasmid pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6, could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmid was digested with *Pme* I and transfected into T-25 flasks of PER.C6 cells using the calcium phosphate co-precipitation technique (Cell Pfect Transfection Kit, Amersham Pharmacia



Biotech Inc). *PmeI* digestion releases the viral genome from plasmid sequences allowing viral replication to occur after cell entry. Viral cytopathic effect (CPE), indicating that virus replication and amplification was occurring was observed following transfection. When CPE was complete, approximately 7-10 days post transfection, the infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Approximately 1 ml of the cell lysate was used to infect a T-225 flask of PER.C6 cells at 80-90% confluence. Once CPE was reached, infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Clarified cell lysates were then used to infect 2-layer NUNC cell factories of PER.C6 cells. Following complete CPE, the virus was purified by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the rescued viruses, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with *HindIII* and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products of the corresponding pre-Adenovirus plasmid (that had been digested with *PmeI/HindIII* prior to labeling) from which they were derived. The expected sizes were observed, indicating that the viruses had been successfully rescued.

### 20 *Example 3*

#### Insertion of an Expression Cassette into pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6

In order to introduce a gag or SEAP expression cassette (*see* Figures 8 and 9, respectively) into the E1 region of pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6, bacterial recombination was again used. A gag expression cassette consisting of the following: 1) the immediate early gene promoter from human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence, was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2 (a precursor to the Ad35 ITR cassettes described above), generating pNEBAd35CMVgagBGHpA. pNEBAd35-2 contains Ad35 sequences from the left end of the genome (bp 4 to 456 and bp 3403 to 3886) with a unique *SwaI* site between bp 456 and 3403 at the position of the deletion. The gag expression cassette was obtained from a previously constructed shuttle plasmid by *EcoRI* digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the *SwaI* site in pNEBAd35-2. This cloning step resulted in the gag expression cassette being inserted into the



E1 deletion between bp 456 and 3403 in the E1 parallel orientation. The shuttle vector containing the gag transgene was digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad35 bp 4 to 456 and bp 3403 to 3886 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6, linearized in the E1 region by digestion with *Swa* I, resulted in the generation of the Ad34 gag-containing pre-Adenovirus plasmid pAd34 $\Delta$ E1gag $\Delta$ E4Ad5Orf6 by homologous recombination. Potential clones were screened by restriction analysis.

A similar strategy was used to generate Ad34 pre-Ad plasmids containing a SEAP expression cassette. In this case a SEAP expression cassette consisting of: 1) the immediate early gene promoter from human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2, generating pNEBAd35CMVSEAPBGHpA. The SEAP expression cassette was obtained from a previously constructed shuttle plasmid by *Eco*RI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the *Swa*I site in pNEBAd35-2. The transgene was then recombined into the pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6, generating pAd34 $\Delta$ E1SEAP $\Delta$ E4Ad5Orf6 as described above for the gag transgene.

All pre-Ad plasmids were rescued into virus and expanded to prepare CsCl purified stocks as described above.

#### *Example 4*

##### Construction of pMRKAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6

To construct an Ad34 pre-Ad plasmid that was composed entirely of Ad34 sequences, an Ad34 ITR cassette was generated. The Ad34 ITR cassette was constructed containing sequences from the right (bp 31584 to 31895 and bp 34409 to 34772) and left (bp 4 to 456 and bp 3402 to 3885) end of the Ad34 genome (*see* Figures 3A-1 to 3A-9) separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd34-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad34 bp 31895 and 34409 generating pNEBAd34-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a



deletion of E1 sequences from Ad34 bp 457 to 3401 with a unique *Swa* I restriction site located in the deletion and an E4 deletion from Ad34 bp 31896 to 34408 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct Ad5Orf6 expression is driven by the Ad34 E4 promoter. The Ad34 sequences (bp 31584 to 31895 and bp 3402 to 3885) in the ITR cassette provided regions of homology with the purified Ad34 viral DNA in which bacterial recombination could occur following cotransformation into BJ 5183 bacteria (Figure 2). The ITR cassette was also designed to contain unique restriction enzyme sites (*Pme*I) located at the end of the viral ITR's so that digestion would release the recombinant Ad34 genome from the plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pMRKAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

### *Example 5*

#### *In Vivo* Studies

##### A. Immunization

Cohorts of 3 rhesus macaques were given single intramuscular injections of one of the two vectors: (1)  $10^{11}$  vp MRKAd5-SEAP (in MRKAd vector backbone of Figures 10A-1 to 10A-45 disclosed in PCT/US01/28861, published March 21, 2002); and (2)  $10^{11}$  vp Ad34 $\Delta$ E1SEAP $\Delta$ E4Ad5Orf6. Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the *Guide for Care and Use of Laboratory Animals*, Institute of Laboratory Animal Resources, National Research Council.

##### B. SEAP Assay

Serum samples were analyzed for circulating human secreted alkaline phosphatase (SEAP) levels using TROPIX phospho-light chemiluminescent kit (Applied Biosystems Inc). Duplicate 5  $\mu$ L aliquots of each serum were mixed with 45  $\mu$ L of kit-supplied dilution buffer in a 96-well white DYNEX plate. Serially diluted solutions of a human placental alkaline phosphatase (Catalog no. M5905, Sigma, St. Louis, MO) in 10% naïve monkey serum served to provide the standard curve. Endogenous SEAP activity in the samples was inactivated by heating the well for 30 minutes at 65 °C. Enzymatic SEAP activities in the samples were



determined following the procedures described in the kit. Chemiluminescence readings (in relative light units) were recorded using DYNEX luminometer. RLU readings were converted to ng/mL SEAP using a log-log regression analyses.

#### C. ELISPOT Assay

5                   The IFN- $\gamma$  ELISPOT assays for rhesus macaques were conducted following a previously described protocol (Allen *et al.*, 2001 *J. Virol.* 75(2):738-749), with some modifications. For antigen-specific stimulation, a peptide pool was prepared from 20-aa peptides that encompass the entire HIV-1 gag sequence with 10-aa overlaps (Synpep Corp., Dublin, CA). To each well, 50  $\mu$ L of  $2-4 \times 10^5$  peripheral blood mononuclear cells (PBMCs)  
10 were added; the cells were counted using Beckman Coulter Z2 particle analyzer with a lower size cut-off set at 80 femtoliters ("fL"). Either 50  $\mu$ L of media or the gag peptide pool at 8  $\mu$ g/mL concentration per peptide were added to the PBMC. The samples were incubated at 37°C, 5% CO<sub>2</sub> for 20-24 hrs. Spots were developed accordingly and the plates were processed using custom-built imager and automatic counting subroutine based on the ImagePro platform  
15 (Silver Spring, MD); the counts were normalized to  $10^6$  cell input.

#### D. Intracellular Cytokine Staining (ICS)

To 1 ml of  $2 \times 10^6$  PBMC/mL in complete RPMI media (in 17x100mm round bottom polypropylene tubes (Sarstedt, Newton, NC)), anti-hCD28 (clone L293, Becton-Dickinson) and anti-hCD49d (clone L25, Becton-Dickinson) monoclonal antibodies were added  
20 to a final concentration of 1  $\mu$ g/mL. For gag-specific stimulation, 10  $\mu$ L of the peptide pool (at 0.4 mg/mL per peptide) were added. The tubes were incubated at 37 °C for 1 hr., after which 20  $\mu$ L of 5 mg/mL of brefeldin A (Sigma) were added. The cells were incubated for 16 hr at 37 °C, 5% CO<sub>2</sub>, 90% humidity. 4 mL cold PBS/2%FBS were added to each tube and the cells were pelleted for 10 min at 1200 rpm. The cells were re-suspended in PBS/2%FBS and stained (30  
25 min, 4 °C) for surface markers using several fluorescent-tagged mAbs: 20  $\mu$ L per tube anti-hCD3-APC, clone FN-18 (Biosource); 20  $\mu$ L anti-hCD8-PerCP, clone SK1 (Becton Dickinson); and 20  $\mu$ L anti-hCD4-PE, clone SK3 (Becton Dickinson). Sample handling from this stage was conducted in the dark. The cells were washed and incubated in 750  $\mu$ L 1xFACS Perm buffer (Becton Dickinson) for 10 min at room temperature. The cells were pelleted and re-suspended in  
30 PBS/2%FBS and 0.1  $\mu$ g of FITC-anti-hIFN- $\gamma$ , clone MD-1 (Biosource) was added. After 30 min incubation, the cells were washed and re-suspended in PBS. Samples were analyzed using all four color channels of the Becton Dickinson FACSCalibur instrument. To analyze the data, the low side- and forward-scatter lymphocyte population was initially gated; a common fluorescence



cut-off for cytokine-positive events was used for both CD4<sup>+</sup> and CD8<sup>+</sup> populations, and for both mock and gag-peptide reaction tubes of a sample.

#### E. Results

Expression: Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 4. Results indicate that the peak levels of SEAP protein produced by the alternative adenovirus serotype were lower than but were within 3-fold of that of MRKAd5 at the same high dose level of 10<sup>11</sup> vp (Figure 4). The levels of SEAP in the serum dropped dramatically after day 10 and were close to background as early as day 15. These observations strongly indicate that the Ad34-based vector is efficient in expressing a transgene following intramuscular administration in a primate.

Immunogenicity: Vaccine-induced T cell responses against HIV-1 gag were quantified using IFN-gamma ELISPOT assay against a pool of 20-aa peptides that encompassed the entire protein sequence. The results are shown in Figure 5; they are expressed as the number of spot-forming cells (SFC) per million peripheral blood mononuclear cells (PBMCs) that responded to the peptide pool or the mock (no peptide) control.

Immunization with gag-expressing Ad34 vector induced detectable levels of circulating gag-specific T cells immediately after a single dose of the vector. The responses improved following a second dose given at wk 4. Overall, the responses to the Ad34-based vector were slightly lower than those induced by the same dose of MRKAd5-gag. The results strongly indicate the Ad34-based vector can prime effectively for HIV-specific T cell responses.

IFN- $\gamma$  ICS analyses of the PBMC from the Ad34-immunized animals revealed that the vector can induce detectable levels of both CD4<sup>+</sup> and CD8<sup>+</sup> HIV-specific T cells (Figure 6).

#### 25 *Example 6*

##### Heterologous Immunization

Cohorts of 3 monkeys were immunized (at wks 0, 4) with 10<sup>11</sup> vp Ad34 $\Delta$ E1 gag $\Delta$ E4Ad5Orf6 followed by a booster at week 24 with 10<sup>10</sup> vp Ad35 $\Delta$ E1 gag $\Delta$ E4Ad5Orf6. Vaccine-induced T cell responses against HIV-1 gag were quantified using IFN-gamma ELISPOT assay against a pool of 20-aa peptides that encompassed the entire protein sequence. The results are shown in Figure 12; they are expressed as the number of spot-forming cells (SFC) per million peripheral blood mononuclear cells (PBMCs) that responded to the peptide pool or the mock (no peptide) control.



Immunization with gag-expressing Ad34 vector induced detectable levels of circulating gag-specific T cells that decreased to between 94-139 SFC/ $10^6$  PBMC at the time of the boost. Heterologous immunization with an Ad35-based HIV vector resulted in as much as a 3-fold increase in T cell responses.

- 5 IFN- $\gamma$  ICS analyses of the PBMCs from the Ad34 primed/Ad35 boosted animals at week 28 revealed that the vector can induce detectable levels of both CD4<sup>+</sup> and CD8<sup>+</sup> HIV-specific T cells (Figure 13).



## WHAT IS CLAIMED IS:

1. A recombinant adenoviral vector of serotype 34 which is at least partially deleted in E1 and devoid of E1 activity.
- 5
2. A population of cells comprising the recombinant adenoviral vector of claim 1.
3. A method for producing recombinant, replication-defective adenovirus particles comprising:
- 10 (a) transfecting a recombinant adenoviral vector of claim 1 into a population of cells; and
- (b) harvesting the resultant recombinant, replication-defective adenovirus.
- 15 4. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 3.
5. A composition comprising purified recombinant adenovirus particles in accordance with claim 4.
- 20 6. A composition in accordance with claim 5 which comprises a physiologically acceptable carrier.
7. A recombinant adenoviral vector of serotype 34 which is at least partially deleted in E1 and devoid of E1 activity which comprises heterologous nucleic acid.
- 25 8. A population of cells comprising the recombinant adenoviral vector of claim 7.
9. A method for producing recombinant, replication-defective adenovirus particles comprising:
- 30 (a) transfecting a recombinant adenoviral vector of claim 7 into a population of cells; and
- (b) harvesting the resultant recombinant, replication-defective adenovirus.



10. A recombinant vector in accordance with claim 7 wherein the vector comprises a gene expression cassette comprising:
- (a) a nucleic acid encoding a protein;
  - 5 (b) a heterologous promoter operatively linked to the nucleic acid encoding the protein; and
  - (c) a transcription termination sequence.
11. A recombinant vector in accordance with claim 10 wherein the gene  
10 expression cassette is inserted into the E1 region.
12. A recombinant vector in accordance with claim 7 wherein the heterologous nucleic acid comprises codons optimized for expression in a human host.
13. A recombinant vector in accordance with claim 7 which comprises  
15 heterologous nucleic acid in the E1 deletion.
14. A recombinant vector in accordance with claim 7 which is at least partially deleted in E3.  
20
15. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 9.
16. A composition comprising purified recombinant adenovirus particles in  
25 accordance with claim 9.
17. A composition in accordance with claim 16 which comprises a physiologically acceptable carrier.
18. A method for effecting the delivery and expression of heterologous  
30 nucleic acid comprising administering the composition of claim 16 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.



19. A method in accordance with claim 18 wherein the composition is preceded or followed by administration of heterologous nucleic acid with an adenovirus of a different serotype.

5 20. A composition in accordance with claim 16 wherein the heterologous nucleic acid encodes an HIV antigen.

21. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 20.

10 22. A composition in accordance with claim 21 wherein the HIV antigen is HIV-1 gag or immunologically relevant modification thereof.

23. A composition in accordance with claim 21 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.

24. A composition in accordance with claim 21 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.

20 25. A recombinant adenoviral vector of serotype 34 which is at least partially deleted in E1 and devoid of E1 activity which comprises an HIV-1 gene.

26. A population of cells comprising the recombinant adenoviral vector of claim 25.

25 27. A method for producing recombinant, replication-defective adenovirus particles comprising:

(a) transfecting a recombinant adenoviral vector of claim 25 into a population of cells; and

30 (b) harvesting the resultant recombinant, replication-defective adenovirus.

28. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 27.



29. A composition comprising purified recombinant adenovirus particles in accordance with claim 28.

5 30. A composition in accordance with claim 29 which comprises a physiologically acceptable carrier.

10 31. A method for effecting the delivery and expression of the HIV-1 gene comprising administering the composition of claim 30 prior or subsequent to administration of the HIV-1 gene with the same or different vector.

32. A method in accordance with claim 31 wherein the composition is preceded or followed by administration of the HIV-1 gene with an adenovirus of a different serotype.

15 33. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 29.

20 34. A composition in accordance with claim 29 wherein the HIV antigen is HIV-1 gag or immunologically relevant modification thereof.

35. A composition in accordance with claim 29 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.

25 36. A composition in accordance with claim 29 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.

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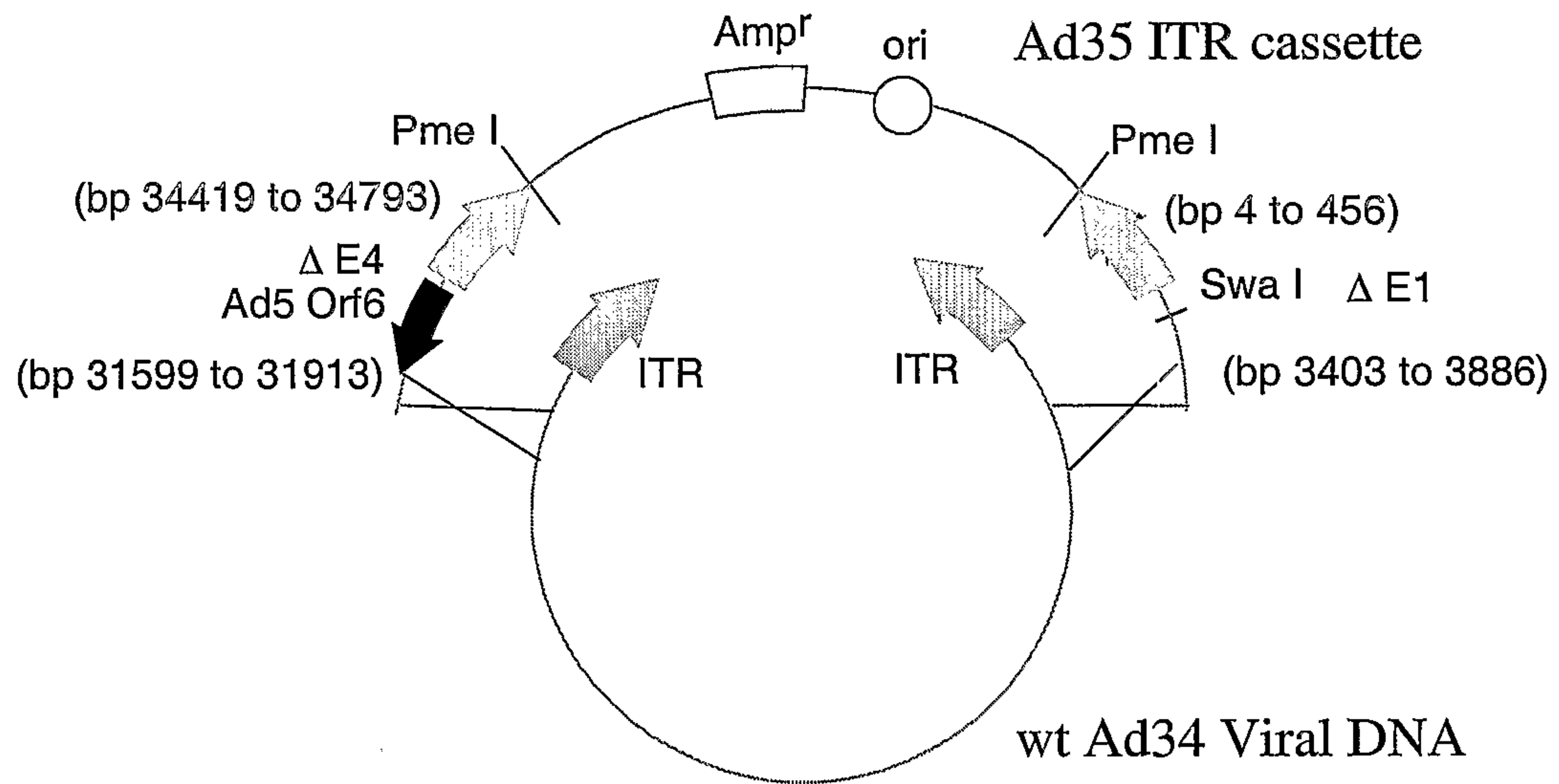


FIG. 1



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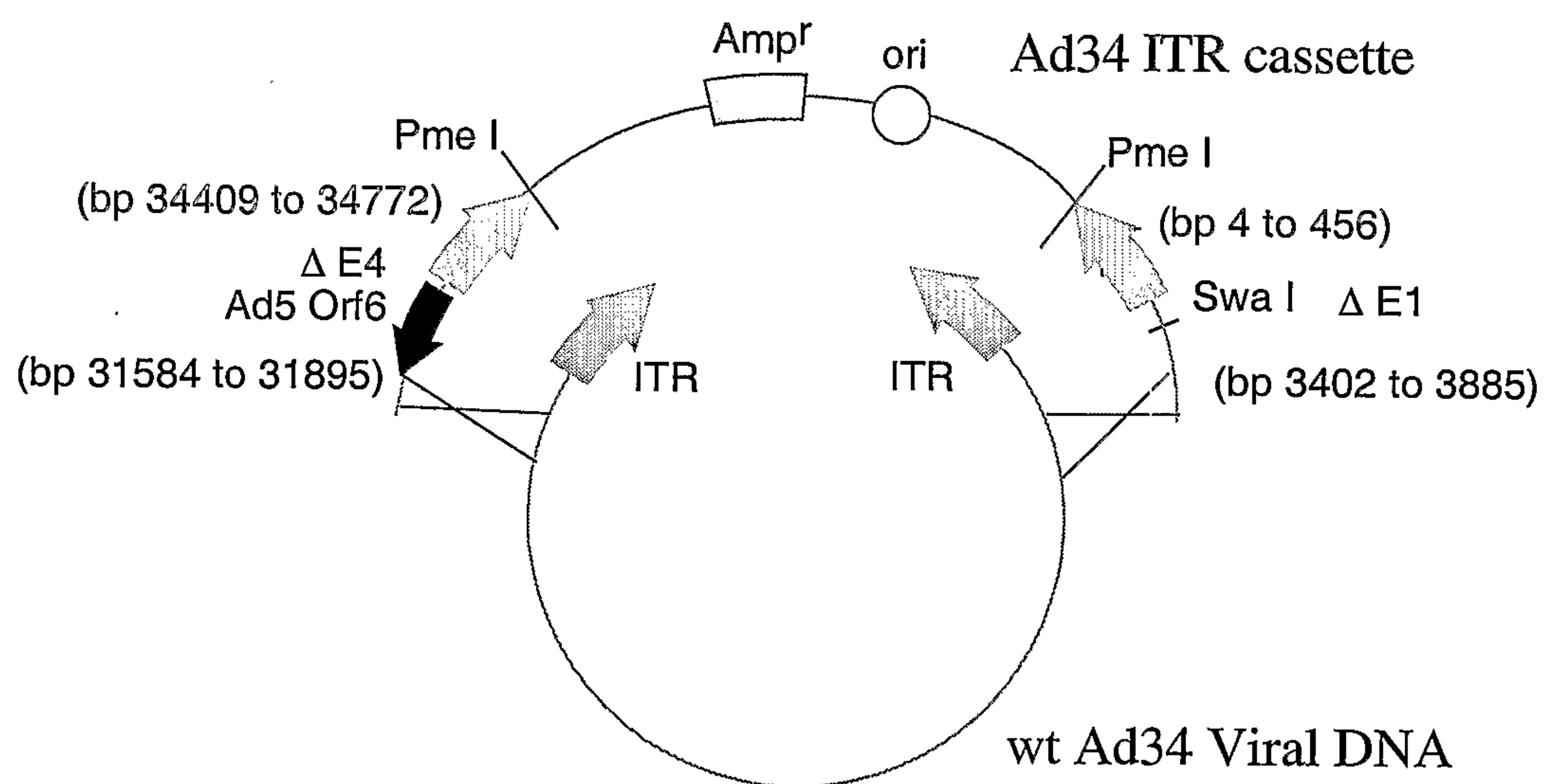


FIG. 2



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1 catcatcaat aatatacctt atagatggaa tgggtgccaat atgtaaataga ggtgatttta  
 61 aaaattgtgg ggtgtgtggt gattggctgt ggggttaacg gctaaacggg gcggcgcggc  
 121 cgtgggaaaa tgacgttttg tgggggtgga gtttttttgc aagttgtcgc gggaaatgtg  
 181 acgcataaaa aggctttttt tctcacggaa ctactgactt tccccacggg atttaacagg  
 241 aatgaggta gttttgaccg gatgcaagtg aaaattgctg atttgccgcg gaaaactgaa  
 301 tgaggaagtg tttttctgaa taatgtggta tttatggcag ggtggagtat ttgttcaggg  
 361 ccaggtagac tttgacccat tacgtggagg tttcgattac cgtgtttttt acctgaattt  
 421 ccgcgtaccg tgtcaaagtc ttctgttttt acgtagggtg cagctgatcg ctacggattt  
 481 tatacctcag ggtttgtgtc aagaggccac tcttgagtgc cagcgagaag agttttctcc  
 541 tctgcgccgg cagttaata ataaaaaat gagagatttg cgatttctgc ctcaggaaat  
 601 aatttctgct gagactggaa atgaaatact ggagcttgtg gtgcacgcc tgatgggaga  
 661 cgatccggag ccacctgtgc agctttttga gcctcctacg cttcaggaac tgtatgattt  
 721 agaggtagag ggatcggagg attctaataga ggaagctgtg aatggctttt ttaccgattc  
 781 tatgctttta gctgctaata aaggattaga attagatccg cctttggaca ctttcgatac  
 841 tccaggggtg attgtggaaa gcggtacagg tgtaagaaaa ttacctgatt tgggttccgt  
 901 ggactgtgat ttgcactgct atgaagacgg gtttcctccg agtgatgagg aggaccatga  
 961 aaaggagcag tctatgcaga ctgcagcggg tgagggagtg aaggctgcca gtgttggtt  
 1021 tcagttggat tgccccggagc ttctggaca tggctgtaag tcttgtgaat ttcacaggaa  
 1081 aaatactgga gtaaaggaac tgttatgttc gctttgttat atgagagcgc actgccactt  
 1141 tatttacagt aagtgtgttt aagttaaaat ttaaaggaat atgctgtttt tcacatgat  
 1201 attgagtggg agttttgtgc ttcttattat aggtcctgtg tctgatgctg atgagtcacc  
 1261 atctcctgat tctactacct cacctcctga gattcaagca cctgttcctg tggacgtgcg  
 1321 caagcccatt cctgtgaagc ttaagcctgg gaaacgtcca gcagtggaaa aacttgagga  
 1381 cttgttacag ggtggggacg gacctttgga cttgagtaca cggaaacggc caagacaata  
 1441 agtgttccat atccgtgttt acttaaggtg acgtcaatat ttgtgtgaga gtgcaatgta  
 1501 ataaaaatat gttaactggt cactggtttt tattgctttt tgggcgggga ctcaggata  
 1561 taagtagaag cagacctgta tggttagctc ataggagctg gctttcatcc atggaggtt  
 1621 gggccatttt ggaagacctt agaaagacta ggcaactggt agaggacgct tcggacggag  
 1681 tctccggttt ttggagattc tggttcgcta gtgaattagc tagggtagtt tttaggataa  
 1741 aacaggacta taaagaagaa tttgaaaagt tgttggtaga ttgcccagga ctttttgaag  
 1801 ctcttaattt gggccatcaa gttcacttta aagaaaaagt tttatcagtt ttagactttt  
 1861 caaccccagg tagaactgcc gctgctgtgg cttttcttac tttatatta gataaatgga  
 1921 tcccgcagac tcatttcagc aggggatac tttttgattt cgtagccaca gcattgtgga  
 1981 gaacatggaa ggttcgcaag atgaggacaa tcttaggta ctggccagtg cagcctttgg  
 2041 gtgtagcggg aatcctgagg catccaccgg tcatgccagc ggttctggag gaggaacagc  
 2101 aagaggacaa cccgagagcc ggcctggacc ctccagtgga ggaggcggag tagctgactt  
 2161 gtctcctgaa ctgcaacggg tgcttactgg atctacgtcc actggacggg ataggggctg  
 2221 taagagggag agggcatcta gtggtactga tgctagatct gagttggctt taagtttaat  
 2281 gagtcgcaga cgtcctgaaa ccatttgggtg gcatgaggtc cagaaagagg gaagggatga  
 2341 agtttctgta ttgcaggaga aatattcact ggaacagggtg aaaacatggt ggttggagcc  
 2401 tgaggatgat tgggaggtgg ccattaataaa ttatgccaaag atagctttga ggcctgataa  
 2461 acagtataag attactagac ggattaatat ccggaatgct tgttacatat ctggaaatgg  
 2521 ggctgaggtg gtaatagata ctcaagacaa ggcagttatt agatgctgca tgatggatat  
 2581 gtggcctgga gtagtcggta tggaaagcagt aacttttgta aatgttaagt ttaggggaga  
 2641 tggttataat ggaatagtgt ttatggccaa taccaaactt atattgcatg gttgtagctt  
 2701 ttttggtttc aacaatacct gtgtagatgc ctggggacag gttagtgtac ggggatgtag  
 2761 tttctatgcy tgttggattg ccacagctgg cagaaccaag agtcaattgt ctctgaagaa  
 2821 atgcatatc caaagatgta acctgggcat tctgaatgaa ggcgaagcaa gggctccgca  
 2881 ctgcgcttct acagatactg gatgttttat ttttaattaag ggcaatgcca gcgtaaagca  
 2941 taacatgatt tgccggtgctt ccgatgagag gccttatcaa atgctcactt gtgccggtgg  
 3001 gcattgtaat atgctggcta ctgtgcatat tgtttcccat caacgcaaaa aatggcctgt  
 3061 ttttgatcac aatgtgttga ccaagtgtac catgcatgca ggtgggctga gaggaatgtt  
 3121 tatgccttac cagtgttaaca tgaatcatgt gaaagtgttg ttggaaccag atgccttttc  
 3181 cagaatgagc ctaacaggaa tctttgacat gaacatgcaa atctggaaga tcctgaggta  
 3241 tgatgatagc agatcgaggg tgccgcgatg cgaatgcgga ggcaagcatg ccaggtcca  
 3301 gccggtgtgt gtagatgtga ctgaagatct gagaccggat catttgggta ttgcccgcac  
 3361 tggagcagag ttccgatcca gtggagaaga aactgactaa ggtgagtatt gggaaaactt  
 3421 ggggtggggg tttcagatgg acagattgag taaaaatttg ttttttctgt ctttcagctg  
 3481 tcatgagtgg aaacgcttct ttaaggggg gagtcttcag cccttatctg acagggcgtc  
 3541 tcccacctcg ggcaggagtt cgtcagaatg ttatgggatc tactgtggat ggaagaccgc  
 3601 tccaaccgc caattcttca acgctgacct atgctacttt aagttcttca cctttggacg  
 3661 cagctgcagc cgccgcgcc gcctctgttg ccgctaacac tgtgcttggga atgggttact  
 3721 atggaagtat cgtggctaata tccacttct ctaataacc ttctaccctg actcaggaca  
 3781 agttacttgt ctttttggcc cagctggagg ctttgacca acgtctgggt gaactttatc  
 3841 agcaggtggc cgagttgcga gtacaaactg agtctgctgt cggcacggca aagtctaaat

FIG. 3A-1



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3901 aaaaaaaaaat tccacaatca atgaataaat aaacgagcct gttggttgatt taaaatcaag  
 3961 tgttttttatt tcattttttcg cgcacgggat gccctagacc accgatctcg atcattgaga  
 4021 acacggtgga tttttttccag aatcctatag aggtggggatt gaatgttttag atacatgggc  
 4081 attaggccat ctttgggggtg gagatagctc cattgaaggg attcatgctc cggggtagtg  
 4141 ttgtaaatca cccagtcata acaaggctgc agtgcacatggt gttgcacaa atcttttaga  
 4201 agtaggctga ttgccacaga taagcccttg gtgtaggtgt ttacaaaccg gttgagctgg  
 4261 gaggggtgca ttcgggggtga aattatgtgc attttggatt ggatttttaa gttggcaata  
 4321 ttgccgcaa gatctcgtct tgggttcacg ttatgaagga ccaccaagac ggtgtatccg  
 4381 gtacatttag gaaatttatc gtgtagcttg gatggaaaag cgtggaaaaa tttggagaca  
 4441 cccttggtgc ctccgagatt tccatgcac tcatccatga taatagcaat ggggccgtgg  
 4501 gcagcagcgc gggcaaacac gttccgtggg tctgacacat catagttagt ttccctgagt  
 4561 aatcatcat aagccatttt aatgaatttg gggcggagag tacccgattg gggtagaat  
 4621 gttccttcgg gccccggagc atagttcccc tcacagattt gcatttccca agctttcagt  
 4681 tccgatggtg gaatcatgct cacctggggg gctatgaaga acaccgtttc tggggcgggg  
 4741 gtgattagtt gggatgatag caagtctctg agcaattgag atttgccaca tccgggtggg  
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 4861 tctcgaagca agggggccac ctctgttcac atttccctta catgcatatt tccccgacc  
 4921 aatccatta ggaggcgtc tctcctagt gatagaagt cttgtagtga gaaaagt  
 4981 ttcagcgggt ttagaccgtc agccatgggc attttggaga gagtttgctg caaagtct  
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 5281 acttctgtcg cttgggcgcc tgtatgtcgg ccaagtagca gtttaccatg agttcgtagt  
 5341 tgagcgcctc ggctgcgtgg cctttggcgc ggagcttacc tttggaagt tctttgcata  
 5401 ccgggcagta taggcatttc agcgcataca gcttggggcgc aaggaaaatg gattctgggg  
 5461 agtatgcac tgccgcgcag gaggcgcaaa cagtttcaca tccaccagc caggtaaat  
 5521 ccggttcatt ggggtcaaaa acaagttttc cgccatattt tttgatgcgt tctttacct  
 5581 tgggtctccat gagttcgtgt cctcgttgag tgacaaacag gctgtccgta tccccgtaga  
 5641 ctgattttac aggcctcttc tccagtggag tgcctcggtc ttcttcgtac aggaactctg  
 5701 accactctga taaaaaggcg cgcgtccagg ccagcaciaa ggaggctatg tgggaggggt  
 5761 agcgatcgtt gtcaaccagg ggtccacct tttccaaagt atgcaaacac atgtcacct  
 5821 cttcaacatc caggaatgtg attggcttgt aggtgtattt cacgtgacct ggggtccccg  
 5881 ctgggggggt ataaaagggg gcggttcttt gctcttctc actgtcttcc ggatcgtgt  
 5941 ccaggaacgt cagctgttgg ggtaggtatt cctctcga ggcgggcatg acctctgcac  
 6001 tcaggttgtc agtttctaag aacgaggagg atttgatatt gacagtgccg gttgagatgc  
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 6181 tcttttctt gtccgcgcgc tctttggcag cgatgttgag ttggacatac tgcggtgcta  
 6241 ggcacttcca ttcggggaag atagttgtca attcatctgg cacgattctc acctgccc  
 6301 ctcgattatg caaggtaatt aaatccacac tgggtggccac ctgcctcga aggggttctg  
 6361 tgggtccaaca gagcctacct cctttcctag aacagaaagg ggaagtggg tctagcataa  
 6421 gttcatcggg agggctctga tccatggtaa agattcccgg aagtaaatcc ttatcaaat  
 6481 agctgatggg agtggggctc tctaaggcca tttgccattc tcgagctgcc agtgcacgct  
 6541 catatgggtt aaggggactg ccccagggca tgggatgggt gagtgcagag gcatacatgc  
 6601 cacagatgtc atagacgtag atgggatcct caaagatgcc tatatagggt ggatagcatc  
 6661 gccccctct gatacttctt cgacatagc catatagttc atgtgatggc gctagcaacc  
 6721 ccggacccaa gttggtgcca ttgggttttt ctgttctgta gacaatctgg cgaaagatgg  
 6781 cgtgagaatt ggaagagatg gtgggtcttt gaaaaatgtt gaaatgggca tgaggtagac  
 6841 ctacagagtc tctgacaaag tgggcataag attcttgaag cttgggtacc agttcggcgg  
 6901 tgacaagtac gtctagggcg cagtagtcaa gtgtttcttg aatgatgtca taacctgggt  
 6961 ggtttttctt ttcccacagt tcgcggttga gaaggtattc ttcgcatcc ttccagtact  
 7021 cttctagcgg aaaccctct ttgtctgcac ggtatgatcc tagcatgtag aactgattaa  
 7081 ctgccttgta agggcagcag ccttctctc cgggtagaga gtatgcttga gcagcttttc  
 7141 gcagcgaagc gtgagtaagg gcgaagggtg ctctgaccat gactttgaga aattggtatt  
 7201 tgaagtccat gtcgtcacag gctccctgtt cccagagttg gaagtctacc cgtttcttgt  
 7261 aggcgggggt gggcaaagcg aaagtaacat cgttgaagag aatcttaccg gctctgggca  
 7321 taaaattgcg agtgatgcgg aaaggctgtg gtacttccgc tcgattgttg atcacctggg  
 7381 cagctaggac gatctcgtcg aaaccgttga tgttgtgtcc tacgatgtat aattctatga  
 7441 aacgcggcgt gcctttgacg tgaggtagct tattgagctc atcaaagggt aggtctgtag  
 7501 ggtcagataa ggcgtagtgt tcgagagccc atctcgtgcag gtgaggattt gcatgtagga  
 7561 atgatgacca aagatccacc gccagtgtct tttgtaactg gtcccgatac tgacgaaaat  
 7621 gctggccaat tgccattttt tctggagtga cacagtagaa ggttctgggg tcttgttgcc  
 7681 atcgatccca ctttagttta atggctagat cgtgggcat gttgacgaga cgctcttctc  
 7741 ctgagagttt catgaccagc atgaaaggaa ctagtgtgtt gccaaaggac cccatccagg

FIG. 3A-2



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7801 tgtaagtttc cacatcgtag gtcaggaaga gtctttctgt gcgaggatga gagccgatcg  
7861 ggaagaactg gatttcctgc caccagttgg aggattggct gttgatgtga tggaaagtaga  
7921 agtttctgcg gcgcgccgag cattcgtgtt tgtgcttgta cagacggccg cagtagtcgc  
7981 agcgttgcac gggttgtatc tcgtgaatga gctgtacctg gcttcccttg acgagaaatt  
8041 tcagtgggaa gccgaggcct ggcgattgta tctcgtgctc ttctatatte gctgtatcgg  
8101 cctgttccatc ttctgtttcg gtggtggtea tgctgacgag cccccgcggg aggcaagtcc  
8161 agacctcggc gcgggagggg cggagctgaa ggaccagagc gcgcaggctg gagctgtcca  
8221 gagtcctgag acgctgcgga ctcaggttag taggtaggga cagaagatta acttgcattga  
8281 tcttttccag ggcgtgcggg aggttcagat ggtacttgat ttccacaggt tcgttttag  
8341 agatgtcaat ggcttgcagg gttccgtgct ctttgggccc cactaccgta cctttgtttt  
8401 ttcttttgat cgggtggggc tctcttgctt cttgcatgct cagaagcgat gacggggacg  
8461 cgcgccgggc ggaagcgggt gttccggacc cggaggcatg gctggtagtg gcacgtcggc  
8521 gccgcgcacg ggcaggttct ggtactgcgc tctgagaaga cttgcgtgcg ccaccacgcg  
8581 tcgattgacg tcttgtatct gacgtctctg ggtgaaagct accggccccg tgagcttgaa  
8641 cctgaaagag agttcaacag aatcaatttc ggtatcgta acggcagctt gtctcagtat  
8701 ttcttgtagc tcaccagagt tgtcctggta ggcgatctcc gccatgaact gctcagtttc  
8761 ttctcctga agatctccgc gaccgcctct ctcgacggtg gccgcgaggt cattggagat  
8821 acggcccatg agttgggaga atgcagtcac gccgcctcgc ttccagacgc ggctgtaaac  
8881 cacggccccc tcggagtctc ttgcccgcac caccacctga gcgaggtaa gctccacgtg  
8941 tctggtgaag accgcatagt tgcataggcg ctgaaaaagg tagttgagtg tgggtggcaat  
9001 gtgttcggcg acgaagaaat acatgatcca tcgtctcagc ggcatttcgc tgacatcgcc  
9061 cagagcttcc aagcgtcca tggcctcgta gaagtccacg gcaaaattaa aaaactggga  
9121 gtttcgcgcg gacacggta atctctctc gagaagacgg atgagttcgg ctatggtggc  
9181 ccgtacttcg cgttcgaagg ctcccgggat ctcttcttcc tcttctatct cttcttccac  
9241 taacatctct tcttcgtctt caggcggggg cggagggggc acacggcgac gtcgacggcg  
9301 cacgggcaaa cggtcgatga atcgttcaat gacctctccg cggcggcgcc gcattggttc  
9361 agtgacggcg cggccgttct cgcgcggtcg cagagtaaaa acaccgccgc gcattctctt  
9421 aaagtgggta ctgggagggt ctccgtttgg gagggagagg gcgctgatta tacattttat  
9481 taattggccc gtagggactg cgcgcagaga tctgatcgtg tcaagatcca cgggatctga  
9541 aaacctttcg acgaaagcgt ctaaccagtc acagtcacaa ggtaggctga gtacggcttc  
9601 ttgtgggccc ggggtggtat gtgttcggtc tgggtcttct gtttcttctt catctcggga  
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9781 ccaagcatca tcctgacatc tagcaagatc ttgtagtag tcttgcatga gccgttctac  
9841 gggcacttct tcctcaccgc ttctgccatg catacgtgtg agtccaaacc cgcgcattgg  
9901 ttgtaccagt gccaaagtcag ctacgactct ttcggcgagg atggcttgcg gtacttgggt  
9961 gaggggtggc tgaaagtcac caaaatccac aaagcgggtg taagccccgg tattaatggt  
10021 gtaagcacag ttggccatga ctgaccagtt aactgtctg taccaggggc gcacgagctc  
10081 ggtgtattta aggcgcgaat aggcgcgggt gtcaaagatg taatcgttgc aggtgcgcac  
10141 cagatactgg taacctataa gaaaatgcgg cgggtggttg cggtagagag gccatcgttc  
10201 tgtagctgga gcgccggggg cgaggtcttc caacataagg cggtagatag cgtagatgta  
10261 cctggacatc cagggtgattc ctgcccgggt agtagaagcc cgaggaaact cgcgtacgcg  
10321 gttccaaatg ttgcgtagcg gcatgaagta gttcattgta ggcacgggtt gaccagtggg  
10381 gcgcgcgcag tcattgatgc tctatagaca cggagaaaat gaaagcgttc agcactcga  
10441 ctccgtagcc tggaggaacg tgaacggggt gggtcgcgggt gtaccccggt tcgagacttg  
10501 tactcgagcc ggccggagcc gcggctaacg tggatttggc actcccgtct cgaccagcc  
10561 tacaaaaatc caggatagcg aatcgagtcg ttttgctggg tgccgaatgg cagggagtg  
10621 agtcctatct tttttttttg ccgctcagat gcatcccgtg ctgacacaga tgcgtcccca  
10681 acaacagccc ccctcgcagc agcagcaacc acaaaaggct gtccctgcaa ctactgcaac  
10741 tgccgctgtg agcgggtgcg gacagcccgc ctatgatctg gacttgggag agggcgaagg  
10801 actggcacgt ctaggtgccc cttcgcgccg gcggcatccg cgagttcaac tgaaaaaaga  
10861 ttctcgcgag gcgtatgtgc cccaacagaa cctatttaga gacagaagcg gcgaggagcc  
10921 ggaggagatg cgagcttccc gctttaacgc gggtcgtgag ctgctcagc gtttggacag  
10981 aagacgagtg ttgcccggag aggatctcga agttgatgaa gtgacagggg tcagtcctgc  
11041 cagggcacac gtggctgacg ccaacctgtt atcggcttac gaacagacag taaaggaaga  
11101 gcgtaatttc caaaagtctt ttaataatca tgtgcaacc ctcatggccc gcgaagaagt  
11161 cacccttggg ttgatgcatt tgtgggattt gatggaagct atcattcaga accctactag  
11221 caaacctctg accgcacagc tgtttctggg ggtgcaacac agcagagaca atgaggcttt  
11281 cagagaggcg ctgctcaaca tcaccgaacc cgaggggaga tggttgtatg atcttatcaa  
11341 cattctacag agtatcatag tgcaggagcg gagcctgggg ctggccgaga aggtggctgc  
11401 catcaattac tcggttttga gcttgggaaa gtattacgct cgcaagatct acaagactcc  
11461 atacgttccc atagacaagg aggtgaagat agatgggttc tacatgcgca tgacgtgaa  
11521 ggtgttgacc ctgagcgatg atcttggggg gtaccgcaat gacagaatgc atcgcgcggg  
11581 gagcgcagc aggagggcgc agttaagcga cagggaaact atgcacagtt tgcaagagc  
11641 tctaactgga gctggaaccg aggggtgagaa ttactttgat atgggagctg acttgcagtg

FIG. 3A-3



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11701 gcagcctagt cgcagggctc tgaacgccgc gacggcagga tgtgagcttc cttacataga  
 11761 agaggcggat gaaggcgagg aggaagaggg cgagtacttg gaagactgat ggcacaacc  
 11821 gtgttttttg ctagatggaa cagcaagcac cggatcccgc aatgcggggcg gcgctgcaga  
 11881 gccagccgtc cggcattaac tctcgggacg attggacca ggccatgcaa cgtatcatgg  
 11941 cgttgacgac tcgcaacccc gaagccttta gacagcaacc ccaggccaac cgtctatcgg  
 12001 ccatcatgga agctgtagtg ccttcccgcct ctaatcccac tcatgagaag gtcctggcca  
 12061 tcgtgaacgc gttggtggag aacaaagcta ttcgtccaga tgaggccgga ctggtataca  
 12121 acgctctctt agaacgcgtg gctcgtaca acagtagcaa tgtgcaaacc aatttggacc  
 12181 gtatgataac agatgtacgc gaagccgtgt ctcagcgcga aaggttccag cgcgatgcca  
 12241 acctgggttc gctggtggcg ttaaagtctt tcttgagtac tcagcctgct aatgtgccgc  
 12301 gtggtcaaca ggattatact aactttttaa gtgctttgag actgatggta tcagaagtac  
 12361 ctcagagcga agtatatcag tccggctctg attacttctt tcagactagc agacagggct  
 12421 tgcagacggt aaatctgagc caagctttaa aaaacctaa aggtttgtgg ggagtgcattg  
 12481 ccccggtagg agaaagagca accgtgtcta gcttgtaaac tccgaactcc cgcctattat  
 12541 tactgttggg agctccttcc accgacagcg gtagcatcga ccgtaattcc tatttggggt  
 12601 acctactaaa cctgtatcgc gaagccatag ggcaaagtca ggtggacgag cagacctatc  
 12661 aagaaattac ccaagtcagt cgcgctttgg gacaggaaga cactggcagt ttggaagcca  
 12721 ctctgaactt cttgcttacc aatcggctctc aaaagatccc tcctcaatat gctcttactg  
 12781 cggaggagga gaggatcctt agatatgtgc agcagagcgt gggattgttt ctgatgcaag  
 12841 agggggcaac tccgactgca gcactggaca tgacagcgcg aaatatggag cccagcatgt  
 12901 atgccagtaa ccgaccttcc attaacaac tgctggacta cttgcacaga gctgccgcta  
 12961 tgaactctga ttatttcacc aatgccatct taaaccgca ctggctgccc ccacctgggt  
 13021 tctacacggg cgaatatgac atgcccgacc ctaatgacgg atttctgtgg gacgacgtgg  
 13081 acagcgatgt tttttcacct ctttctgac atcgcacgtg gaaaaaggaa ggcggcgata  
 13141 gaatgcattc ttctgcatcg ctgtccgggg tcattggtgc taccgcccgt gagcccagat  
 13201 ctgcaagtcc ttttccctagt ctacctttt ctctacacag tgtacgtagc agcgaagtgg  
 13261 gtagaataag tcgcccagat ttaatgggag aagaggagta cctaaacgat tccttgctca  
 13321 gaccggcaag agaaaaaat ttcccaaca atggaataga aagtttgggt gataaaatga  
 13381 gtatagggaa gacttatgct caggatcaca gagacgagcc tgggatcatg gggactacaa  
 13441 gtatagcggc cgtagacgc cagcgcctat acagacagag gggctctgtg tgggacgatg  
 13501 aggatcggc cgatgatagc agcgtattgg acttgggtgg gagaggaagg ggcaaccctg  
 13561 ttgctcattt gcgcccctcg tgggtggta tgggtgtaaaa aaaaataaaa aagaaaaaac  
 13621 tcaccaaggc catggcgacg agcgtacgtt cgttcttctt tattatctgt gtctagtata  
 13681 atgaggcgag tcgtgctagg cggagcgggt gtgtatccgg agggctctcc tccttcgtac  
 13741 gagagcgtga tgcagcagca gcagcgcagc gcggtgatgc aatccccact ggaggctccc  
 13801 tttgtgcctc cgcgatacct ggcacctacg gagggcagaa acagcattcg ttactcggaa  
 13861 ctggcacctc agtacgatac caccaggttg tatctggtgg acaacaagtc ggcggacatt  
 13921 gcttctctga actatcagaa tgaccacagc aacttcttga ccacgggtgg gcaaaacaat  
 13981 gactttacc ctacggaagc cagcaccag accattaact ttgatgaacg atcgcgggtg  
 14041 ggcggtcagc taaaaaccat catgcatact aacatgccc acgtgaacga gtatatgttt  
 14101 agtaacaagt tcaaagcgcg tgtgatggtg tccagaaaac ctctgaggg tgttagagta  
 14161 gacgataatt atgatcataa gcaagatatt ctaaaatag agtgggtcga gtttactttg  
 14221 ccagaaggca acttttcggg cactatgact atcgacttga tgaacaatgc catcatagac  
 14281 aattacttga aagtgggagc acagaatgga gtgttggaaa gtgacattgg tgtaagtcc  
 14341 gacactagga acttcaagtt gggatgggat ccagaaacta agttgatcat gcctgggggt  
 14401 tacacctatg aggccttcca tcctgacatc gtattgctgc ctggctgagg agtgacttt  
 14461 accgaaagcc gtctgagcaa ccttcttggc attagaaaga aacacccatt ccaagagggt  
 14521 tttaagatct tgtatgagga tttagaagga ggaaatattc cagccctttt ggatgtagat  
 14581 gcttatgaga acagcaagaa agatcaaaaa gccaaaatag aagctgctgc agaagctaaa  
 14641 gcaaacatag ttgccaacga tccggtaagg gtggctaacg ctagtgaat caggggagac  
 14701 agttttgccg caacatccgt tccgactaaa gaatcattat tggatgatgt gtctcaaac  
 14761 atagagttaa aactcactat taagcctgtg gaaaaagatg gcaaaaacag aagttacaat  
 14821 gtgttggag ataaaatcaa cacggcctat cgcagttggt acctttcgta caattatggc  
 14881 gaccccgaaa aaggagtgcg ttcctggaca ttgctacca cctcagatgt cacctgcgga  
 14941 gcggagcagg tctactggtc gcttccagac atgatgcagg atcctgtcac tttccgctcc  
 15001 actagacaag tcagtaacta ccctgtggtg ggtgcagagc ttatgcccgt cttttcaaag  
 15061 agcttctaca acgaacaagc tgtgtactcc cagcagctcc gccagtccac ctgcttacg  
 15121 cacgtcttca accgctttcc tgagaaccag attttaatcc gtcgcccggc gccacaatt  
 15181 accaccgtca gtgaaaacgt tcctgctctc acagatcacg ggaccctgcc gttgcccagc  
 15241 agtatccggg gactccaacg tgtgaccgtt actgacgcca gacgcccac ctgtccctac  
 15301 gtgtacaagg cactgggcat agtcgcaccg cgcgtccttt caagccgcac tttctaaaa  
 15361 aaaaaaaaa atgtccgttc ttatctcgcc cagtaataac accggttggg gtctgcccgc  
 15421 tcccagcaag atgtacggag gcgcacgcaa acgttctacc caacatcccg tgcgtgttcg  
 15481 cgggcatttt cgcgctccat ggggtgccct caagggccgc actcgcgttc gaaccaccgt  
 15541 cgatgatgta atcgatcagg tgggtgccga cgcccgtaat tatactccta ctgcccctac

FIG. 3A-4



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15601 atctactgtg gacgcagtta ttgacagtgt agtggctgac gctcgcact atgctcgacg  
15661 taagagccgg cgaaggcgca ttgccagacg tcaccgagct accactgcca tgcgagcagc  
15721 aagagctctg ctacgaagag ctagacgcgt ggggcgaaga gccatgctta gggcggccag  
15781 acgtgcagct tcggggcgca gcgcccggcag gtcccgcagg caagcagccg ctgtcgcagc  
15841 ggcgactatt gccgacatgg cccaatcgcg aagaggcaat gtatactggg tgcgtgacgc  
15901 tgccaccggg caacgtgtac ccgtgcgcac ccgtccccct cgcacttaga agatactgag  
15961 cagtctccga tgttgtgtcc cagcggcgag gatgtccaag cgcaaataca aggaagaat  
16021 gctgcaggtt atcgcacctg aagtctacgg ccaaccgttg aaggatgaaa aaaaaccccg  
16081 caaaatcaag cgggtaaaaa aggacaaaaa agaagaggaa gatggcgatg atgggctggc  
16141 ggagtttgtg cgcgagtttg ccccacggcg acgcgtgcaa tggcgtgggc gcaaagtctg  
16201 acatgtgttg agacctggaa cttcgggtgg ctttacacc ggcgagcgtt caagcgtac  
16261 ttttaagcgt tcctatgatg aggtgtacgg ggatgatgat attcctgagc aggcagctga  
16321 ccgattaggc gagtttgctt atggcaagcg tagtagaata aatcccaagg atgaaacagt  
16381 gtccataccc ttggatcatg gaaatcccac ccctagtctt aaaccggta ctttgcagca  
16441 agtgttaccg gtaactccgc gaacaggtgt taaacgcgaa ggtgaagatt tgtatcccac  
16501 tatgcaactg atgggtgccc aacgccagaa gttggaggac gttttggaga aagtaaaagt  
16561 ggatccagat attcaacctg aggttaaagt gagaccatt aagcaggtag cgcctggctt  
16621 gggagtacaa actgtagaca ttaaaattcc cactgaaagt atggaagtgc aaactgaacc  
16681 cgcaaagcct actgccacct ccactgaagt gcaaaccggac ccattggatgc ccatgcctat  
16741 tacaactgac gccgtcggtc ccactcgaag atcccgcgaa aagtacggtc cagcaagtct  
16801 gttgatgccc aactatgtcg tacaccatc tattattcct actcctgggt accgaggcac  
16861 tcgctactat cgcagccgaa acagtacttc ccgcccgcgc cgcaagacac ctgcaaatcg  
16921 cagtcgtcgc cgtagacgca caagcaaacc gattcccggc gccctgggtgc ggcaagtgta  
16981 ccgcaatggg agtgccggaac ctttgacact gccgcgtgcg cgttaccatc ctagtatcat  
17041 cacttaatca atggtgcccg tgcctccttg cagatatggc cctcacttgt cgccttcgcg  
17101 ttcccacac tggttaccga ggaagaaact cgcgcccgtag aagagggatg ttggggcgcg  
17161 gaatgcgacg ctacaggcga cggcgtgcta tccgcaagca attgcccggg ggttttttgc  
17221 cagccttaat tccaattatc gctgctgca ttggcgcaat accaggcata gcttccgtgg  
17281 cggttcaggc ctgcgaacga cattgacatt ggaaaaaaa aaaacgtata aataaaaaat  
17341 acaatggact ctgacactcc tggactgtg actatgtttt cttagagatg gaagacatca  
17401 atttttcatc cttggctccg cgacacggca cgaagccgta catgggcacc tggagcgaca  
17461 tcggcacgag ccaactgaac gggggcgcct tcaattggag cagtatctgg agcgggctta  
17521 aaaatthtgg ctcaaccata aaaacatagc ggaacaaagc ttggaacagc agtacaggac  
17581 aggcgcttag aaataaactt aaagaccaga acttccaaca aaaagtagtc gatgggatag  
17641 cttcccggat caatggagtg gtagatttgg ctaaccaggc tgtgcagaaa aagataaaca  
17701 gtcgthtggg cccgcccgca gcaaccccag gtgaaatgca agtggaggaa gaaattcctc  
17761 cgccagaaaa acgaggcgac aagcgtccgc gtcccgatth ggaagagacg ctggtgacgc  
17821 gcgtagatga accgccttct tatgaggaag caacgaagct tggaatgcc accactagac  
17881 cgatagcccc tatggccacc ggggtgatga aaccttctca gttgcatcga cccgtcacct  
17941 tggatthtgc ccctcctcct gctgctactg ctgtaccgcg ttctaagcct gtcgctgccc  
18001 cgaaaccagt cgccgtagcc aggtcacgtc ccggggggcg tccctcgtcca aatgcacact  
18061 ggcaaaatac tctgaacagc atcgtgggtc taggcgtgca aagtgtaaaa cgcgctcgtc  
18121 gctthttaat aaatatggag tagcgttaa cttgcctatc tgtgtatatg tgtcattaca  
18181 cgccgtcaca gcatcagagg aaaaaaggaa gaggtcgtgc gtcgacgctg agttactttc  
18241 aagatggcca ccccatcgat gctgccccaa tgggcataca tgcacatcgc cggacaggat  
18301 gcttcggagt acctgagtcc gggctctggg cagttcgcgc gcgccacaga cacctacttc  
18361 aatctgggaa ataagthttag aaatcctacc gtagcgcgca ccacgatgt gaccaccgat  
18421 cgtagccagc ggctcatgth gcgcttctgt cccgttgacc gggaggacaa tacatactct  
18481 tacaagtgcc ggtacaccct ggccgtgggc gacaacagag tgctggatat ggccagcagc  
18541 ttctthtga ttaggggctg gttggacaga ggtcccagtt taaacccta ttctggtagc  
18601 gcttacaact ccctggctcc taaaggcgtc ccaatgcat ctcagthtgg ggataaggga  
18661 gttacaagca ctggcctagt ggacgacggc aatactgatg atggggaaga agccaaaaaa  
18721 gcaacataca cthtthtggaa tgctccagta aaagccgagg ctgaaatcac aaaagacgga  
18781 ttgcccgtgg gctthtggagt ttcaactgaa ggtcctaac caatctatgc tgataagctt  
18841 tatcagccag aacctcaagt gggagacgaa actthtggactg acctagacgg aaaaaccgaa  
18901 gagtatggag ggaggthtct taaacctgaa actaaaatga aacctgcta cggatcttht  
18961 gctaaacctc ctaatthttaa aggagthttag gcaaaggtht aaccaaaaga agacgatggc  
19021 actaacaaca tcgagthttag cattgacatg aacttcttht acttaagatc acaagatca  
19081 gaactcaaac ctaaaattht aatgthttag gaaaatthtgg acctggaatg tccagatact  
19141 catgthttht acaaacctgg agthttagat gctagthtctg agaccaatct tggacaacag  
19201 tctatgccc aacagacccaa ctacatthtgg ttcagagata acttcatcgg actthttagc  
19261 tataacagta ctggcaacat ggggthttag gctggccaag cgtctcagtht gaatgcagtht  
19321 gthttagctg aggcagaaa cacagaactg tctthtaccac tctthttaga ctctctgggc  
19381 gacagaacca gactthttag catgthttag caggctthtgg acagthttaga tctctgatgta  
19441 cgtgthttag aaaaatcatg thtthttagat gaactthtcca actthttagtht tccgthttag

FIG. 3A-5



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19501 ggtgtcggtc cgcgaaacaga tagttacaag gagattaagc caaatggaga ccaatctact  
 19561 tggacaaatg tagaccaac tggcagcagt gaacttgcta agggaaatcc atttgccatg  
 19621 gaaattaacc ttcaagccaa tctatggcga agtttccttt attccaatgt ggctctatat  
 19681 ctcccagact cgtacaaata caccocgtcc aatgtcactc ttccagaaaa caaaaacacc  
 19741 tacgactaca tgaacgggcg ggtgggtgccc ccatctctag tagacaccta tgtgaacatt  
 19801 ggtgccagggt ggtctctgga tgccatggac aatgtcaacc cattcaacca ccaccgtaac  
 19861 gctggcttgc gttaccgatc catgcttctg ggtaacggac gttatgtgcc tttccacata  
 19921 caagtgcctc aaaaattctt cgctgttaaa aacctgctgc ttctcccagg ctctacact  
 19981 tatgagtgga actttaggaa ggatgtaaac atggttctac agagttccct cggtaacgac  
 20041 ctacgggtag atggcgccag catcagtttt acgagcatca acctctatgc tacttttttc  
 20101 cccatggctc acaacaccgc ttccaccctt gaagccatgc tgcggaatga caccaatgat  
 20161 cagtcattca acgactacct atctgcagct aacatgctct accccattcc tgccaatgca  
 20221 accaatattc ccatttccat tccttctcgc aactgggocg ctttcagagg ctggtcattt  
 20281 accagactga aaaccaaga aactccctct ttggggctct gatttgacc ctacttcgtc  
 20341 tattctgggt ctattcccta cctggatggt acctctacc tgaaccacac ttttaagaag  
 20401 gtttccatca tgtttgactc ttcagtgagc tggcctggaa atgacagggt actatctcct  
 20461 aacgaatttg aaataaagcg cactgtggat ggcgaaggct acaacgtagc ccaatgcaac  
 20521 atgaccaag actggttctt ggtacagatg ctgcaccaact acaacatcgg ctatcagggc  
 20581 ttctacattc cagaaggata caaagatcgc atgtattcat ttttcagaaa cttccagccc  
 20641 atgagcaggc aggtggttga tgaggatcaat tacaagact tcaaggccgt cgccataccc  
 20701 taccaacaca acaactctgg ctttgtgggt tacatggctc cgaccatgcy tcaaggtaa  
 20761 ccctatcccg ctaactatcc ctatccactc attggaacaa ctgccgtaa tagtgttacg  
 20821 cagaaaaagt tcttgtgtga cagaaccatg tggcgcatac cgttctcaag caacttcatg  
 20881 tctatgggag cccttacaga cttgggacag aacatgctct atgccaactc agctcatgct  
 20941 ctggacatga cctttgaggt ggatcccatg gatgagcca cctgcttta tcttctcttc  
 21001 gaagttttcg acgtggtcag agtgcacag ccacaccgcy gcatcatcga ggcagctac  
 21061 ctgcgtacac cgttctcggc cggtaacgct accacgtaag aagcttcttg cttcttgcaa  
 21121 acagcagcty caaccatggc ctgcggatcc caaacggct ccagcagagca agagctcaga  
 21181 gccattgtcc aagacctggg ttgcggacca tattttttgg gaacctttga taagcgttc  
 21241 ccgggggtca tggccccga taagctcgc tgtgccattg taaatacggc cggacgtgag  
 21301 acggggggag agcactggtt ggctttcgggt tggaaaccac gttctaacac ctgctacctt  
 21361 tttgatcctt ttggattctc ggatgatcgt ctcaaacaga tttaccagtt tgaatatgag  
 21421 ggtctcctgc gccgcagcgc tcttgctacc aaggaccggt gtattacgct ggaaaaatct  
 21481 acccagaccg tgcagggccc ccgttctgcc gcctgcggac ttttctgcty catgttctt  
 21541 catgcctttg tgcactggcc tgaccgtccc atggacggaa accccaccat gaaattgcta  
 21601 actggagtgc caaacaacat gcttcattct cctaaagtcc agcccaccct gtgtgacaat  
 21661 caaaaagcac tctaccattt tctcaatacc cattcgcctt attttctgctc tcatcgtaca  
 21721 cacatcgaiaa gggccactgc gttcagaccgt atggatgtgc aataatgatt catgtaaaca  
 21781 acgtgttcaa taaacagcac tttatttttt acatgtatcy aggctctgga ttacttattt  
 21841 atttacaagt cgaatgggtt ctgacgagaa tcagaatgac ccgcaggcag tgatacgttg  
 21901 cggaaactgat acttggggtg ccacttgaat tcgggaatca ccaacttggg aaccggtata  
 21961 tcgggcagga tgtcactcca cagctttcty gtcagctgca aagctcccag caggtcagga  
 22021 gccgaaatct tgaaatcaca attaggacca gtgctctgag cgcgagagtt gcggtacacc  
 22081 ggattgcagc actgaaacac catcagcgcag ggatgtctta cgcttgccag cacgggtggga  
 22141 tctgcaatca tgcccacatc cagatcttca gcattggcaa tgctgaacgg ggtcatcttg  
 22201 caggctctgcc taccatggc gggcaccxaa ttaggcttgt ggttacaatc gcagtgagg  
 22261 gggatcagta tcatcttggc ctgatcctgt ctgattccty gatacacggc tctcatgaaa  
 22321 gcatcatatt gcttgaaagc ctgctgggct ttactaccct cggataaaa catcccgcag  
 22381 gacctgctcy aaaactgggt agctgcgcag ccggcatcat tcacacagca gcgggcgtca  
 22441 ttggtggcta tttgcaccac acttctgccc cagcggtttt gggtgatttt ggttcgctcy  
 22501 ggattctcct tcaaggctcy ttgtccgttc tcgctggcca catccatctc gataatctgc  
 22561 tccttctgaa tcataatatt gccatgcaag cacttcagct tgccctcata atcattgcag  
 22621 ccatgaggcc acaacgcaca gcctgtacat tcccaattat ggtgggcgat ctgagaaaaa  
 22681 gaatgtatca ttccctgcag aaatcttccc atcatcgtgc tcagtgtctt gtgactagtg  
 22741 aaagttaact ggatgcctcy gtgctcctcy ttcacgtact ggtgacagat gcgcttgat  
 22801 tgttcgtgct gctcaggcat tagtttaaaa gaggttctaa gttcgttatc cagcctgtac  
 22861 ttctccatca gcagacacat cacttccatg cctttctccc aagcagacac caggggcaag  
 22921 ctaatcggat tcttaacagt gcaggcagca gctcctttag ccagagggte atctttggcg  
 22981 atcttctcaa tgcttctttt gccatccttc tcaacgatgc gcacgggcy gtagctgaaa  
 23041 cccactgcta caagttgcy ctcttctctt tcttcttctc tgtcttgact gatgtcttgc  
 23101 atggggacat gtttggctt ccttggcttc ttttctgggg gtatcggagg aggaggacty  
 23161 tcgctccgtt ccggagacag ggaggattgt gacgttctcy tcaccattac caactgacty  
 23221 tcggtagaag aacctgacc cacacggcga cagggtttc tcttctgggg cagaggtgga  
 23281 ggcgattgcy aagggtcgc gtccgaccty gaaggcgat gactggcaga acccttccg  
 23341 cgttctgggg tgtgctccct gtggcggctcy cttactgat tctcttctcy gctggccatt

FIG. 3A-6



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23401 gtgttctcct aggcagagaa acaacagaca tggaaactca gccattgctg tcaacatcgc  
 23461 cacgagtgcc atcacatctc gtcctcagcg acgaggaaaa ggagcagagc ttaagcattc  
 23521 caccgcccag tcctgccacc acctctaccc tagaagataa ggaggtcgac gcatctcatg  
 23581 acatgcagaa taaaaaagcg aaagagtctg agccagacat cgaacaagac cggggctatg  
 23641 tgacaccggg ggaacacgag gaagagttga aacgctttct agagagagag gatgaaaact  
 23701 gcccaaaaaca gcaagcggat aactatcacc aagatgctgg aaatagggat cagaacaccg  
 23761 actacctcat agggcttgac ggggaagacg cgctccttaa acatctagca agacagtcac  
 23821 tcatagtcaa ggatgcatta ttggacagaa ctgaagtgcc catcagtgtc gaagagctca  
 23881 gccgcgccca cgagcttaac ctatcttcac ctcgtactcc ccccaaactg cagccaaacg  
 23941 gcacctgcca gcaaatcct cgcttaaact tttatccagc ttttgctgtg ccagaagtac  
 24001 tggctaccta tcacatcttt tttaaaaatc aaaaaattcc agtctcctgc cgcgctaata  
 24061 gcacccgccc cgatgcccta ctcaatctgg gacctgggtc acgcttacct gatatagctt  
 24121 ccttggaaga ggttccaaag atcttcgagg gtctggggcaa taatgagact cggggccgcaa  
 24181 atgctctgca aaagggagaa aatggcatgg atgagcatca cagcgttctg gtggaattgg  
 24241 aaggcgataa tgccagactc gcagtactca agcgaagcgt cgaggtcaca cactttgcat  
 24301 accccgctgt caacctgcc cctaaagtca tgacggccgt catggaccag ttactcatta  
 24361 agcgcgcaag tcccctttca gaagacatgc atgaccaga tgccctgtgat gagggtaaac  
 24421 cagtggtcag tgatgagcag ctaaccgat ggctggggcac cgactctccc cgggatttgg  
 24481 aagagcgtcg caagcttatg atggccgtgg tgctgggttac cgtagaacta gagtgtcttc  
 24541 ggcgtttctt taccgattca gaaaccttgc gcaaactcga agagaatctg cactacactt  
 24601 ttagacacgg ctttgtgccc caggcatgca agatatctaa cgtggaactc accaacctgg  
 24661 tttcctacat gggatttctg catgagaatc gcctaggaca aagcgtgctg cacagcacc  
 24721 ttaaggggga agcccggcgt gattacatcc gcgattgtgt ttatctctac ctgtgccaca  
 24781 cgtggcaaac cggcatgggt gtatggcagc aatgtttaga agaacagAAC ctgaaagagc  
 24841 taaacaagct cttacagaaa tctcttaagg ttctgtggac agggttcgc gagcgcaccg  
 24901 tcgcttccga cctggcagac ctcatcttcc cagagcgtct cagggttact ttgCGAAACg  
 24961 gactgcctga ctttatgagc cagagcatgc ttaacaattt tcgctctttc atcctggaac  
 25021 gctccggtat cctgcccgcc acctgctgcg cactgcccctc cgactttgtg cctctcacct  
 25081 accgcgaatg cccccgccc ctatggagtc actgctacct gttccgtctg gccaactacc  
 25141 tctcctacca ctcggtatgt atcgaggatg tgagcggaga cggcttgctg gagtgtcact  
 25201 gccgctgcaa tctgtgcacg ccccaccggt ccctagcttg caacccccag ttgatgagcg  
 25261 aaaccagat aataggcacc tttgaattgc aaggccccag cagccaaggc gatgggtctt  
 25321 ctcttgggca aagtttaaaa ctgacccccg gactgtggac ctccgcctac ttgCGCAAGt  
 25381 ttgccccgga agattaccac ccctatgaaa tcaagttcta tgaggaccaa tcacagcctc  
 25441 cgaaagccga actttcggcc tgcgtcatca cccagggggc aattctggcc caattgcaag  
 25501 ccatccaaaa atcccgcCAA gaatttctac tgaaaaaggg taaggggggtc taccttgacc  
 25561 cccagaccgg cgaggaactc aacacaaggT tccctcagga tgtcccaacg acgagaaagc  
 25621 aagaagtga aggtgcagcc gccgccccca gaagatatgg aggaagattg ggacagtcag  
 25681 gcagaggaag cggaggagga ggacagtctg gaggacagtc tggaggaaga cagtttggag  
 25741 gaggaaaacg aggaggcaga ggaggtggaa gaagtaaccg ccgacaaaca gttatcctcg  
 25801 gctgcggaga caagcaacag cgctaccatc tccgctccga gtcgaggaac cgggCGGCGt  
 25861 cccagcagta gatgggacga gaccggacgc ttcccgaacc caaccagcgc ttccaagacc  
 25921 ggtaagaagg atcggcaggg atacaagtc tggcggggggc ataagaatgc catcatctcc  
 25981 tgcttgcatg agtgcggggg caacatatcc ttcacgcggc gctacttgct attccaccat  
 26041 ggggtgaact ttccgcgcaa tgttttgcac tactaccgct acctccacag cccctactat  
 26101 agccagcaaa tcccggcagt ctgcacagat aaagacagcg gcgggcaccct ccaacagaaa  
 26161 accagcagcg gcagttagaa aatacacaac aagtgcagca acaggaggat taaagattac  
 26221 agccaacgag ccagcgcaaa cccgagagtt aagaaatcgg atctttccaa ccctgtatgc  
 26281 catcttcag cagagtcggg gccaaagagca ggaactgaaa ataaaaaacc gatctctgCG  
 26341 ttcgctcacc agaagttgtt tgtatcaca gagcgaagat caacttcagc gactctcga  
 26401 ggacgcccag gctctcttca acaagtactg cgcgctgact cttaaagagt aggcagcGAC  
 26461 cgcgcttatt caaaaaaggc ggggaattaca tcatcctcga catgagtaaa gaaattccca  
 26521 cgccttacct gtggagttat cagccccaaa tgggattggc ggcaggcgcc tcccaggact  
 26581 actccaccgg catgaattgg ctcagcgccg ggccttctat gatttctcga gttaatgata  
 26641 tacgcgccca ccgaaaccaa atacttttgg aacagtcagc tcttaccacc acgccccgCC  
 26701 aacaccttaa tcccagaaat tggcccggcg ccctagtgtA ccaggaaagt cccgctccca  
 26761 cactgtatt acttcctcga gacgcccagg ccgaagtcca aatgactaat gcagggtgCG  
 26821 agttagcggg cggctccacc ctatgtcgtc acaggcctcg gcataatata aaacgcctga  
 26881 tgatcagagg ccgaggtatc cagctcaacg acgagtcggt gagctctccg cttgggtctac  
 26941 gaccagacgg aatctttcag attgcccggct gcgggagatc ttccttcacc cctcgtcagg  
 27001 ctgttctgac tttgaaagt tcgtcttcgc aacccccgctc gggcggaatc gggaccgttc  
 27061 aatttgtgga ggagtttact ccctctgtct acttcaacc cttctccgga tctcctgggc  
 27121 actaccgga cgagttcata ccgaacttcg acgcgattag cgagtcagtg gacggctacg  
 27181 attgatgtct ggtgacgCG ctgagctatc tcggctgCGa catctagacc actgcccCG  
 27241 ctttcgctgc tttgcccggg aactcattga gttcatctac ttcgaactcc ccaaggatca

FIG. 3A-7



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27301 ccctcaaggt ccggcccacg gagtgcggat tactatcgaa ggcaaaatac actctcgcct  
 27361 gcaacgaatt ttctcccagc ggcccgtgct gatcgagcga gaccagggaa acaccacggt  
 27421 ttccatctac tgcatttgta atcaccocgg attgcatgaa agcctttgct gtcttatgtg  
 27481 tactgagttt aataaaaact gaattaagac tctcctacgg actgcogett cttcaaccog  
 27541 gattttacaa ccagaagaac gaaacttttc ctgtcgtcca ggactctggt aacttcacct  
 27601 ttctactca caaactagaa gctcaacgac tacaccgctt tccagaagc atttcccta  
 27661 ctaatactac tttcaaaacc ggaggtgagc tccaaggtct tctacagaa aacccttggg  
 27721 tggaagcggg cttgttagtg cttagaattc ttgcccgttg gcttgtgatt attctttgct  
 27781 acctatacac accttgcttc actttcctag tgggtgtgtg gtattggttt aaaaaatggg  
 27841 gcccatacta gtcttgcttg ttttactttc gcttttggaa ccgggttctg ccaattacga  
 27901 tccatgtcta gacttcgacc cagaaaactg cacacttact tttgcaccog acacaagccg  
 27961 catctgtgga gttcttatta agtgcggatg ggaatgcagg tccgttgaaa ttacacacaa  
 28021 taacaaaacc tggaacaata ccttatccac cacatgggag ccaggagttc ccgagtggta  
 28081 cactgtctct gtccgaggtc ctgacggttc catccgcatt agtaacaaca ctttcatttt  
 28141 ttctgaaatg tgcgatctgg ccatgttcat gagcaaacag tattctctat ggctcctag  
 28201 caaggacaac atcgtaacgt tctccattgc ttattgcttg tgcgcttggc ttcttactgc  
 28261 tttactgtgc gtatgcatac acctgcttgt aaccactcgc atcaaaaacg ccaataacaa  
 28321 agaaaaaatg ctttaacctc tttctgttta cagacatggc ttctcttaca tctctcatat  
 28381 ttgtcagcat tgtcactgcc gctcacggac aaacagtcgt ctctatccct ctaggacata  
 28441 attacactct cataggacc ccaatcactt cagaggtcat ctggaccaaa ctgggaagcg  
 28501 ttgattactt tgatataatc tgcaacaaaa caaaaccaat aatagtaact tgcaacatac  
 28561 aaaatcttac attgattaat gttagcaaag ttacagcgg ttactattat ggttatgaca  
 28621 gatacagtag tcaatataga aattacttgg ttctgtttac ccagttaaaa accacgaaaa  
 28681 tgccaaatat ggcaaagatt cgatccgatg acaattctct agaaactttt acatctcca  
 28741 ccacaccoga cgaaaaaac atcccagatt caatgattgc aattgttgca gcggtggcag  
 28801 tgggtgatggc actaataata atatgcatgc ttttatatgc ttgtcgtac aaaaagtttc  
 28861 atcctaaaaa acaagatctc ctactaaggc ttaacattta atttcttttt atacagccat  
 28921 ggtttccact accacattcc ttatgcttac tagtcttgca actctgactt ctgctcgtc  
 28981 acacctcact gtaactatag gctcaaactg cactataaaa ggacctcaag gtggtcattgt  
 29041 cttttggtgg agaatatatg acaatggatg gtttcaaaaa ccatgtgacc aacctggtag  
 29101 atttttctgc aacggcagag acctaacat tatcaacgtg acagcaaatg acaaaggctt  
 29161 ctattatgga accgactata aaagtagttt agattataac attattgtac tgccatctac  
 29221 cactccagca ccccgcaaaa ctactttctc tagcagcagt gtcgctaaca atacaatttc  
 29281 caatccaacc tttgcccgcg ttttaaaacg cactgtgaat aattctacaa cttcacatac  
 29341 aacaatttcc acttcaacaa tcagcattat cgctgcagtg acaattggaa tatctattct  
 29401 tgtttttacc ataacctact acgctgctg ctatagaaaa gacaaacata aaggtgatcc  
 29461 attacttaga tttgatattt aatttgttct tttttttttt atttacagta tgggtgaacac  
 29521 caatcatggg acctagaaat ttcttcttca ccatactcat ttgtgcattt aatgtttgcg  
 29581 ctactttcac agcagtagcc acagcaacc cagactgtat aggagcattt gcttcctatg  
 29641 cactttttgc tttgttact tgcactctgc tatgtagcat agtctgcctg gttattaatt  
 29701 ttttccaact tctagactgg atccttgtgc gaattgccta cctgcgccac catcccgaat  
 29761 accgcaacca aaatctcgcg gcacttctta gactcatcta aaacctgca ggctatacta  
 29821 ccaatatttt tgcttctatt gcttccctac gctgtctcaa cccagctgc ctatagtact  
 29881 ccaccagaac accttagaaa atgcaaattc caacaaccgt ggtcatttct tgcttgctat  
 29941 cgagaaaaat cagaaattcc ccaaattta ataatgattg ctggaataat taatataatc  
 30001 tgttgcacca taatttcatt tttgatatac ccctatttg attttggctg gaatgctccc  
 30061 aatgcacatg atcatccaca agaccagag gaacacattc ccctacaaaa catgcaacat  
 30121 ccaatagcgc taatagatta cgaaagtga ccacaacccc cactactccc tgctattagt  
 30181 tacttcaacc taaccggcgg agatgactga aacactcacc acctccaatt ccgccgagga  
 30241 tctgctcgat atggacggcc gcgtctcaga acagcgactt gcccaactac gcatccgcca  
 30301 gcagcaggaa cgcgcggcca aagagctcag agatgtcatc caaattcacc aatgcaaaaa  
 30361 aggcataatc tgtttggtaa aacaagcca gatatcctac gagatcaccg ctactgacca  
 30421 tcgcctctct tacgaacttg gcccccacg acaaaaattt acctgcatgg tgggaatcaa  
 30481 ccccatagtt atcaccagc aaagtggaga tactaagggt tgcattcact gctcctgcca  
 30541 ttccatcgag tgcacctaca cctgctgaa gaccctatgc ggcttaagag acctgctacc  
 30601 aatgaattaa aaaatgatta ataaaaaatc acttacttga aatcagcaat aaggtctctg  
 30661 ttgaaatttt ctcccagcag cacctcactt ccctcttccc aactctggta ttctaaacc  
 30721 cgttcagcgg catactttct ccatacttta aaggggatgt caaattttag ctctctctct  
 30781 gtaccacaaa tcttcatgtc tttcttccca gatgaccaag agagtccggc tcagtgactc  
 30841 cttcaaccct gtctaccct atgaagatga aagcacctcc caacaccct ttataaacc  
 30901 agggtttatt tccccaaatg gcttcacaca aagcccagac ggagttctta ctttaaaatg  
 30961 ttaacccca ctaacaacca caggcggatc tctacagcta aaagtgggag ggggacttac  
 31021 agtggatgac actgatggta ccttacaaga aacatacgt gctacagcac ccattactaa  
 31081 aaataatcac tctgtagaac tatccattgg aatggatta gaaactcaaa acaataaact  
 31141 atgtgcaaaa ttgggaaatg ggttaaaatt taacaacggt gacatttgta taaaggatag

FIG. 3A-8



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31201 tattaacacc ttatggactg gaataaacc tccacctaac tgtcaaattg tggaaaacac  
 31261 taatacaaat gatggcaaac ttacttttagt attagtaaaa aacggagggc ttgttaatgg  
 31321 ctacgtgtct ctagttagtg tatcagacac tgtgaaccaa atgttcacac aaaagacagc  
 31381 aacatccaa ttaagattat attttgactc ttctggaaat ctattaactg atgaatcaga  
 31441 cttaaaaatt ccacttaaaa ataaatcttc tacagcgacc agtgaaactg tagccagcag  
 31501 caaagccttt atgccaagta ctacagctta tcccttcaac accactacta gggatagtga  
 31561 aaactacatt catggaatat gttactacat gactagttat gatagaagtc tatttccctt  
 31621 gaacatttct ataatgctaa acagccgtat gatttcttcc aatggtgcct atgccataca  
 31681 atttgaatgg aatctaaatg caagtgaatc tccagaaagc aacatagcta cgctgaccac  
 31741 atcccccttt ttcttttctt acattacaga agacgacaac taaaataaag ttaagtgtt  
 31801 tttatthaaa atcacaaaat tcgagtagtt attttgcctc caccttccca tttgacagaa  
 31861 tacaccaatc tctccccacg cacagcttta aacatttggg taccattaga gatagacatt  
 31921 gtttttagatt ccacattcca aacagtttca gagcgagcca atctggggtc agtgatagat  
 31981 aaaaatccat cgcgatagtc ttttaaagcg ctttcacagt ccaactgctg cggatgcaaa  
 32041 tccggagtct ggatcacggg catctggaag aagaacgatg ggaatcataa tccgaaaacg  
 32101 gtatcggacg attgtgtctc atcaaacca caagcagccg ctgtctgcgt cgctccgtgc  
 32161 aactgctggt tatgggatca ggggtccacag tgcctgaag catgatttta atagccctta  
 32221 acatcaactt tctgggtgca tgcgcgacg aacgcattct gatttctctc aaatctttgc  
 32281 agtaggtaca acacattatt acaatattgt ttaataaacc ataattaaa ggcctccagc  
 32341 caaaactcat atctgatata atcgcccctg catgaccatc ataccaagt ttaatataaa  
 32401 ttaaatgacg ttccctcaaa aacacactac ccacatacat gatctctttt ggcatgtgca  
 32461 tattaacaat ctgtctgtac catggacaac gttgggtaat catgcaacc aatataacct  
 32521 tccggaacca cactgccaac accgctcccc cagccatgca ttgaagtga cctgctgat  
 32581 tacaatgaca atgaagaacc caattctctc gaccgtgaat cacttgagaa tgaaaaatat  
 32641 ctatagtggc acaacataga cataaatgca tgcattctct cataattttt aactcctcag  
 32701 gatttagaaa catatcccag ggaataggaa gctcttgacg aacagtaaag ctggcagaac  
 32761 aaggaagacc acgaacacaa cttacactat gcatagtcac agtatcacia tctggcaaca  
 32821 gcgggtggtc ttcagtcata gaagctcggg tttcattttc ctcacaacgt ggtaactggg  
 32881 ctctggtgta aggggtgatgt ctggcgcatg atgtcgagcg tgcgcgcaac cttgtcataa  
 32941 tggagtgtct tcttgacatt ctctgatttt gtatagcaaa acgcgggcct ggcagaacac  
 33001 actcttcttc gccttctatc ctgcccctta gcgtgttccg tgtgatagtt caagtacagc  
 33061 cacactctta agttgggtcaa aagaatgctg gcttcagttg taatcaaac tccatcgcat  
 33121 ctaattgttc tgaggaaatc atccacggta gcatatgcaa atcccaacca agcaatgcaa  
 33181 ctggattgctg tttcaagcag gagaggagag ggaagagacg gaagaacct gtttaattttt  
 33241 attccaaacg atctcgcagt acttcaaatt gtagatcgcg cagatggcat ctctcgcccc  
 33301 cactgtgttg gtgaaaaagc acagctaaat caaaagaaat gcgattttca aggtgctcaa  
 33361 cgggtggcttc caacaaagcc tccacgcgca catccaagaa caaaagaata ccaaagaag  
 33421 gagcattttc taactcctca atcatcatat tacattcctg caccattccc agataatttt  
 33481 cagctttcca gccttgaatt attcgtgtca gttcttgtgg taaatccaat ccacacatta  
 33541 caaacaggtc ccggagggcg cctccacca ccattcttaa acacaccctc ataatgacaa  
 33601 aatatcttgc tctgtgtca cctgtagcga attgagaatg gcaacatcaa ttgacatgcc  
 33661 cttggctcta agttcttctt taagttctag ttgtaaaaac tctctcatat tatcaccaaa  
 33721 ctgcttagcc agaagcccc cgggaacaag agcaggggac gctacagtgc agtacaagcg  
 33781 cagacctccc caattggctc cagcaaaaac aagattggaa taagcatatt gggaccgccc  
 33841 agtaatatca tcgaagttgc tggaaatata atcaggcaga gtttcttcta aaaattgaat  
 33901 aaaagaaaaa tttgcaaaa aaacattcaa aacctctggg atgcaaattgc aataggttac  
 33961 cgcgctgcgc tccaacattg ttagttttga attagtctgc aaaaataaaa aaaaaaaca  
 34021 gcgtcatatc atagtagcct gacgaacagg tggataaatc agtctttcca tcacaagaca  
 34081 agccacaggg tctccagctc gaccctcgta aaacctgtca tggtgattaa acaacagcac  
 34141 cgaaagtcc tcgcggtgac cagcatgat aattcttgat gaagcataca atccagacat  
 34201 gttagcatca gttaacgaga aaaaacagcc aacatagcct ttgggtataa ttatgcttaa  
 34261 tcgtaagtat agcaaagcca cccctcgcgg atacaaagta aaaggcacag gagaataaaa  
 34321 aatataatta tttctctgct gctgttcagg caacgtcgc cccggctcct ctaaatacac  
 34381 atacaaagcc tcatcagcca tggcttacca gacaaagtac agcgggacag cacaagctct  
 34441 aaagtcactc tccaacctct ccacaatata tatacacaag ccctaaactg acgtaatggg  
 34501 agtaaagtgt aaaaaatccc gccaaacca acacacacc cgaaactgcg tcaccagggg  
 34561 aaagtacagt ttcacttccg caatcccaac aagcgtcact tctcttttct cacggtagct  
 34621 cacatcccat taacttgcaa cgtcattttc ccacggccgc gccgccccgt ttagccgtta  
 34681 accccacagc caatcaccac acaccacaca attttataaa tcacctcatt tacatattgg  
 34741 caccattcca tctataaggt atattattga tgatg

FIG. 3A-9



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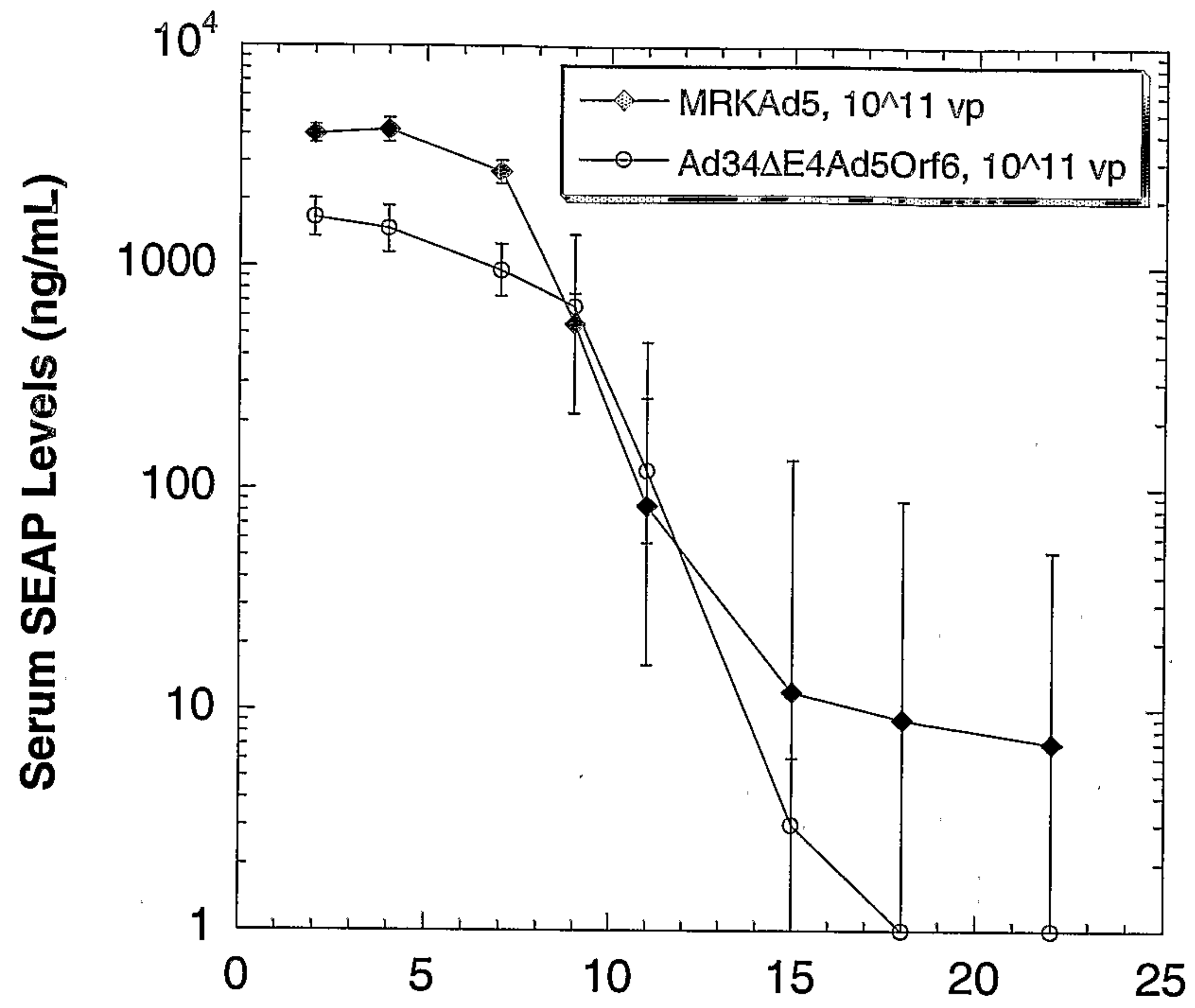


FIG. 4



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Vaccine Wk 0, 4, 24	Monkey ID	Pre		Wk 4		Wk 8		Wk 24		Wk 28		Wk 36	
		Mock	Gag <sup>a</sup>	Mock	Gag	Mock	Gag	Mock	Gag	Mock	Gag	Mock	Gag
MRKAd5gag, 10 <sup>11</sup> vp	00C018	1	5	13	1025	0	824	8	756	0	474	0	383
MRKAd5gag, 10 <sup>11</sup> vp	00C034	0	4	5	219	5	404	3	445	3	339	0	216
MRKAd5gag, 10 <sup>11</sup> vp	00C058	4	4	3	1086	0	440	4	1439	0	2338	0	940
Ad34ΔE1gagΔE4Ad5Orf6, 10 <sup>11</sup> vp	00D038	6	8	5	111	1	301	0	224	1	536	0	233
Ad34ΔE1gagΔE4Ad5Orf6, 10 <sup>11</sup> vp	00D042	6	30	4	89	4	264	1	73	0	181	0	69
Ad34ΔE1gagΔE4Ad5Orf6, 10 <sup>11</sup> vp	00D066	3	18	1	118	1	816	0	429	0	439	0	273

FIG. 5



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Vaccine	Monk ID	IFN- $\gamma$ <sup>+</sup> CD4 <sup>+</sup> CD3 <sup>+</sup> per 10 <sup>6</sup> Lymphocytes		IFN- $\gamma$ <sup>+</sup> CD8 <sup>+</sup> CD3 <sup>+</sup> per 10 <sup>6</sup> Lymphocytes	
		Mock	Gag <sup>a</sup>	Mock	Gag <sup>a</sup>
Ad34 $\Delta$ E1gag $\Delta$ E4Ad5Orf6	00D038	22	154	130	450
	00D042	32	118	96	171
	00D066	12	238	150	442

FIG. 6



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Sequence of the open reading frame for FL-gag (human codon optimized)

atgggtgctagggcttctgtgctgtctggtggtgagctggacaagtgggagaagatcaggctgaggcctggtggc  
aagaagaagtacaagctaaagcacattgtgtgggcctccagggagctggagaggtttgctgtgaaccctggcctg  
ctggagacctctgaggggtgcaggcagatcctggggccagctccagccctccctgcaaacaggctctgaggagctg  
aggtccctgtacaacacagtggctaccctgtactgtgtgcaccagaagattgatgtgaaggacaccaaggaggcc  
ctggagaagattgaggaggagcagaacaagtccaagaagaaggcccagcaggctgctgctggcacaggcaactcc  
agccaggtgtcccagaactaccccattgtgcagaacctccagggccagatggtgcaccaggccatctccccccgg  
accctgaatgcctgggtgaagggtggtggaggagaaggccttctcccctgaggtgatccccatgttctctgcctg  
tctgaggggtgccacccccaggacctgaacacctatgctgaacacagtggggggccatcaggctgccatgcagatg  
ctgaaggagaccatcaatgaggaggctgctgagtgggacaggctgcacacctgtgcacgctggccccattgcccc  
ggccagatgagggagcccaggggctctgacattgctggcaccacctccacctccaggagcagattggctggatg  
accaacaacccccccatccctgtgggggaaatctacaagaggtggatcctcctgggcctgaacaagattgtgagg  
atgtactccccacctccatcctggacatcaggcagggccccaaggagcccttcagggactatgtggacaggttc  
tacaagacctgagggctgagcaggcctcccaggaggtgaagaactggatgacagagacctgctggtgcagaat  
gccaacctgactgcaagaccatcctgaaggccctggggccctgctgccacctggaggagatgatgacagcctgc  
caggggggtggggggccctgggtcacaaggccaggggtgctggctgaggccatgtcccagggtgaccaactccgccacc  
atcatgatgcagaggggcaacttcaggaaccagaggaagacagtgaagtgcttcaactgtggcaagggtggggccac  
attgccaagaactgtagggccccaggaagaagggtgctggaagtgtggcaaggaggggccaccagatgaaggac  
tgcaatgagagggcaggccaacttctggggcaaaatctggccctcccacaaggggcaggcctggcaacttctccag  
tccaggcctgagcccacagcccctcccagggagtccttcagggttggggaggagaagaccacccccagccagaag  
caggagcccattgacaaggagctgtaccccctggcctccctgaggtccctggttggcaacgacctcctcccag  
taaaataaagcccgggcagat

FIG. 7



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1 ccattgcata cgttgtatcc atatcataat atgtacattt atattggctc atgtccaaca  
 61 ttaccgccat gttgacattg attattgact agttattaat agtaatcaat tacgggggtca  
 121 ttagttcata gcccatatat ggagttccgc gttacataac ttacggtaaa tggccccgct  
 181 ggctgaccgc ccaacgacc cgcgccattg acgtcaataa tgacgtatgt tcccatagta  
 241 acgccaatag ggactttcca ttgacgtcaa tgggtggagt atttacggta aactgccac  
 301 ttggcagtac atcaagtgta tcatatgcca agtacgccc ctattgacgt caatgacggg  
 361 aatggcccg cctggcatta tgcccagtac atgacctat gggactttcc tacttggcag  
 421 tacatctacg tattagtcac cgctattacc atgggtgatgc ggttttggca gtacatcaat  
 481 gggcgtggat agcggtttga ctacgggga tttccaagtc tccaccccat tgacgtcaat  
 541 gggagtttgt tttggcacca aatcaacgg gactttcca aatgtcgtaa caactccgcc  
 601 ccattgacgc aatggggcgg taggcgtgta cgggtggagg tctatataag cagagctcgt  
 661 ttagtgaacc gtcagatcgc ctggagacgc catccacgct gttttgacct ccatagaaga  
 721 caccgggacc gatccagcct ccgcccggcgg gaacgggtgca ttggaacgcg gattccccgt  
 781 gccaagagtg agatctaccA TGGGTGCTAG GGCTTCTGTG CTGTCTGGTG GTGAGCTGGA  
 841 CAAGTGGGAG AAGATCAGGC TGAGGCCTGG TGGCAAGAAG AAGTACAAGC TAAAGCACAT  
 901 TGTGTGGGCC TCCAGGGAGC TGGAGAGGTT TGCTGTGAAC CCTGGCCTGC TGGAGACCTC  
 961 TGAGGGGTGC AGGCAGATCC TGGGCCAGCT CCAGCCCTCC CTGCAAACAG GCTCTGAGGA  
 1021 GCTGAGGTCC CTGTACAACA CAGTGGCTAC CCTGTACTGT GTGCACCAGA AGATTGATGT  
 1081 GAAGGACACC AAGGAGGCC TGGAGAAGAT TGAGGAGGAG CAGAACAAGT CCAAGAAGAA  
 1141 GGCCAGCAG GCTGCTGCTG GCACAGGCAA CTCCAGCCAG GTGTCCCAGA ACTACCCCAT  
 1201 TGTGCAGAAC CTCCAGGGCC AGATGGTGCA CCAGGCCATC TCCCCCGGA CCCTGAATGC  
 1261 CTGGGTGAAG GTGGTGGAGG AGAAGGCCTT CTCCCCTGAG GTGATCCCCA TGTTCTCTGC  
 1321 CCTGTCTGAG GGTGCCACCC CCCAGGACCT GAACACCATG CTGAACACAG TGGGGGGCCA  
 1381 TCAGGCTGCC ATGCAGATGC TGAAGGAGAC CATCAATGAG GAGGCTGCTG AGTGGGACAG  
 1441 GCTGCATCCT GTGCACGCTG GCCCATTGC CCCCAGCCAG ATGAGGGAGC CCAGGGGCTC  
 1501 TGACATTGCT GGCACCACCT CCACCCTCCA GGAGCAGATT GGCTGGATGA CCAACAACCC  
 1561 CCCCATCCCT GTGGGGGAAA TCTACAAGAG GTGGATCATC CTGGGCCTGA ACAAGATTGT  
 1621 GAGGATGTAC TCCCCACCT CCATCCTGGA CATCAGGCAG GGCCCCAAGG AGCCCTTCAG  
 1681 GGAATATGTG GACAGGTTCT ACAAGACCCT GAGGGCTGAG CAGGCCTCCC AGGAGGTGAA  
 1741 GAACTGGATG ACAGAGACCC TGCTGGTGCA GAATGCCAAC CCTGACTGCA AGACCATCCT  
 1801 GAAGGCCCTG GGCCCTGCTG CCACCCTGGA GGAGATGATG ACAGCCTGCC AGGGGGTGGG  
 1861 GGGCCCTGGT CACAAGGCCA GGGTGCTGGC TGAGGCCATG TCCAGGTGA CCAACTCCGC  
 1921 CACCATCATG ATGCAGAGGG GCAACTTCAG GAACCAGAGG AAGACAGTGA AGTGCTTCAA  
 1981 CTGTGGCAAG GTGGGCCACA TTGCCAAGAA CTGTAGGGCC CCCAGGAAGA AGGGCTGCTG  
 2041 GAAGTGTGGC AAGGAGGGCC ACCAGATGAA GGAATGCAAT GAGAGGCAGG CCAACTTCCT  
 2101 GGGCAAATC TGGCCCTCCC ACAAGGGCAG GCCTGGCAAC TTCCTCCAGT CCAGGCCTGA  
 2161 GCCACAGCC CCTCCCAGG AGTCCTTCAG GTTTGGGGAG GAGAAGACCA CCCCAGCCA  
 2221 GAAGCAGGAG CCCATTGACA AGGAGCTGTA CCCCTGGCC TCCCTGAGGT CCCTGTTTGG  
 2281 CAACGACCCC TCCTCCAGT AAaataaagc ccgggcagat ctgatctgct gtgccttcta  
 2341 gttgccagcc atctgttgtt tgcccctccc ccgtgccttc cttgacctg gaaggtgcca  
 2401 ctcccactgt ctttccctaa taaaatgagg aaattgcatc gcattgtctg agtaggtgtc  
 2461 attctattct ggggggtggg gtggggcagc acagcaaggg ggaggattgg gaagacaata  
 2521 gcaggcatgc tggggatgcg gtgggctcta

SEQ ID NO: 2

FIG. 8



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1 ccattgcata cgttgtatcc atatcataat atgtacattt atattggctc atgtccaaca
61 ttaccgccat gttgacattg attattgact agttattaat agtaatcaat tacgggggtca
121 ttagttcata gcccatatat ggagttccgc gttacataac ttacggtaaa tggcccgcct
181 ggctgaccgc ccaacgaccc ccgccattg acgtcaataa tgacgtatgt tcccatagta
241 acgccaatag ggactttcca ttgacgtcaa tgggtggagt atttacggta aactgccac
301 ttggcagtac atcaagtgta tcatatgcca agtacgcccc ctattgacgt caatgacggg
361 aatggcccg cctggcatta tgcccagtac atgaccttat gggactttcc tacttggcag
421 tacatctacg tattagtcac cgctattacc atgggtgatgc ggttttggca gtacatcaat
481 gggcgtggat agcggtttga ctcacgggga tttccaagtc tccaccccat tgacgtcaat
541 gggagtttgt tttggcacca aatcaacgg gactttcca aatgtcgtaa caactccgcc
601 ccattgacgc aatggggcgg taggcgtgta cggtgggagg tctatataag cagagctcgt
661 ttagtgaacc gtcagatcgc ctggagacgc catccacgct gttttgacct ccatagaaga
721 caccgggacc gatccagcct ccgcccggc gaacgggtgca ttggaacgcg gattccccgt
781 gccaagagtg agatctaagt aagcttcctg cATGCTGCTG CTGCTGCTGC TGCTGGGCCT
841 GAGGCTACAG CTCTCCCTGG GCATCATCCC AGTTGAGGAG GAGAACCCGG ACTTCTGGAA
901 CCGCGAGGCA GCCGAGGCC TGGGTGCCGC CAAGAAGCTG CAGCCTGCAC AGACAGCCGC
961 CAAGAACCTC ATCATCTTCC TGGGCGATGG GATGGGGGTG TCTACGGTGA CAGCTGCCAG
1021 GATCCTAAAA GGCAGAAGA AGGACAAACT GGGCCTGAG ATACCCCTGG CCATGGACCG
1081 CTTCCCATAT GTGGCTCTGT CCAAGACATA CAATGTAGAC AAACATGTGC CAGACAGTGG
1141 AGCCACAGCC ACGGCTACC TGTGCGGGGT CAAGGGCAAC TTCCAGACCA TTGGCTTGAG
1201 TGCAGCCGCC CGCTTTAACC AGTGCAACAC GACACGCGGC AACGAGGTCA TCTCCGTGAT
1261 GAATCGGGCC AAGAAAGCAG GGAAGTCAGT GGGAGTGGTA ACCACCACAC GAGTGCAGCA
1321 CGCCTCGCCA GCCGGCACCT ACGCCACAC GGTGAACCGC AACTGGTACT CGGACGCCGA
1381 CGTGCCTGCC TCCGCCGCC AGGAGGGGTG CCAGGACATC GCTACGCAGC TCATCTCCAA
1441 CATGGACATT GACGTGATCC TAGGTGGAGG CCGAAAGTAC ATGTTTCGCA TGGGAACCC
1501 AGACCCTGAG TACCCAGATG ACTACAGCCA AGGTGGGACC AGGCTGGACG GGAAGAATCT
1561 GGTGCAGGAA TGGCTGGCGA AGCGCCAGGG TGCCCGGTAT GTGTGGAACC GCACTGAGCT
1621 CATGCAGGCT TCCCTGGACC CGTCTGTGAC CCATCTCATG GGTCTCTTTG AGCCTGGAGA
1681 CATGAAATAC GAGATCCACC GAGACTCCAC ACTGGACCCC TCCCTGATGG AGATGACAGA
1741 GGCTGCCCTG CGCCTGCTGA GCAGGAACCC CCGCGGCTTC TTCCTCTTCG TGGAGGGTGG
1801 TCGCATCGAC CATGGTCATC ATGAAAGCAG GGCTTACCGG GCACTGACTG AGACGATCAT
1861 GTTCGACGAC GCCATTGAGA GGGCGGGCCA GCTCACCAGC GAGGAGGACA CGCTGAGCCT
1921 CGTCACTGCC GACCACTCCC ACGTCTTCTC CTTCCGAGGC TACCCCTGC GAGGGAGCTC
1981 CATCTTCGGG CTGGCCCCTG GCAAGGCCCG GGACAGGAAG GCCTACACGG TCCTCCTATA
2041 CGGAAACGGT CCAGGCTATG TGCTCAAGGA CGGCGCCCGG CCGGATGTTA CCGAGAGCGA
2101 GAGCGGGAGC CCCGAGTATC GGCAGCAGTC AGCAGTGCC CTTGGACGAAG AGACCCACGC
2161 AGGCGAGGAC GTGGCGGTGT TCGCGCGCGG CCCGCAGGCG CACCTGGTTC ACGGCGTGCA
2221 GGAGCAGACC TTCATAGCGC ACGTCATGGC CTTCCGCCGC TGCCCTGGAGC CCTACACCGC
2281 CTGCGACCTG GCGCCCCCG CCGGCACCAC CGACGCCGCG CACCCGGGT AAaccgtggt
2341 ccccgcggtg cttcctctgc tggccgggac atcaggtggc ccccgctgaa ttggaatcga
2401 tcagaattca gtcgacgata tctgatcacg atctgatctg ctgtgccttc tagttgccag
2461 ccatctggtt tttgcccctc ccccgctgct tccttgacct tgggaaggtgc cactcccact
2521 gtcctttcct aataaaatga ggaaattgca tcgcattgtc tgagtaggtg tcattctatt
2581 ctgggggggtg ggggtggggca gcacagcaag ggggaggatt ggaagacaa tagcaggcat
2641 gctgggggatg cggtggggctc ta

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FIG. 9



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## PacI

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**1** TTCTTAATTA ACATCATCAA TAATATACCT TATTTTGGAT TGAAGCCAAT  
 AAGAAITTAAT TGTAGTAGTT ATTATATGGA ATAAAACCTA ACTTCGGTTA

**51** ATGATAATGA GGGGGTGGAG TTTGTGACGT GGCGCGGGGC GTGGGAACGG  
 TACTATTACT CCCCCACCTC AACACTGCA CCGCGCCCCG CACCCTTGCC

**101** GGCGGGTGAC GTAGTAGTGT GGCGGAAGTG TGATGTTGCA AGTGTGGCGG  
 CCGCCCCTG CATCATCACA CCGCCTTCAC ACTACAACGT TCACACCGCC

**151** AACACATGTA AGCGACGGAT GTGGCAAAG TGACGTTTTT GGTGTGCGCC  
 TTGTGTACAT TCGCTGCCTA CACCGTTTTT ACTGCAAAA CCACACGCGG

**201** GGTGTACACA GGAAGTGACA ATTTTCGCGC GGTTTTAGGC GGATGTTGTA  
 CCACATGTGT CCTTCACTGT TAAAAGCGCG CCAAATCCG CCTACAACAT

**251** GTAAATTTGG GCGTAACCGA GTAAGATTTG GCCATTTTCG CGGGAAAAC  
 CATTTAAACC CGCATTTGGCT CATTCTAAAC CGGTAAAAGC GCCTTTTGA

**301** GAATAAGAGG AAGTGAAATC TGAATAATTT TGTGTTACTC ATAGCGCGTA  
 CTTATTCTCC TTCACTTTAG ACTTATTAAC ACACAATGAG TATCGCGCAT

**351** ATATTTGTCT AGGGCCGCGG GGACTTTGAC CGTTTACGTG GAGACTCGCC  
 TATAACAGA TCCCGGCGCC CCTGAAACTG GCAAATGCAC CTCTGAGCGG

**401** CAGGTGTTTT TCTCAGGTGT TTTCCGCGTT CCGGGTCAA GTTGGCGTTT  
 GTCCACAAA AGAGTCCACA AAAGGCGCAA GGCCAGTTT CAACCGCAA

**451** TATTATTATA GGCGGCCGCG ATCCATTGCA TACGTTGTAT CCATATCATA  
 ATAATAATAT CCGCCGGCGC TAGGTAACGT ATGCAACATA GGTATAGTAT

**501** ATATGTACAT TTATATTGGC TCATGTCCAA CATTACCGCC ATGTTGACAT  
 TATACATGTA AATATAACCG AGTACAGGTT GTAATGGCGG TACAACGTGTA

**551** TGATTATTGA CTAGTTATTA ATAGTAATCA ATTACGGGGT CATTAGTTCA  
 ACTAATAACT GATCAATAAT TATCATTAGT TAATGCCCCA GTAATCAAGT

**601** TAGCCCATAT ATGGAGTTCC GCGTTACATA ACTTACGGTA AATGGCCCGC  
 ATCGGGTATA TACCTCAAGG CGCAATGTAT TGAATGCCAT TTACCGGGCG

**651** CTGGCTGACC GCCCAACGAC CCGGCCCAT TGACGTCAAT AATGACGTAT  
 GACCGACTGG CGGGTTGCTG GGGGCGGGTA ACTGCAGTTA TTACTGCATA

**701** GTTCCCATAG TAACGCCAAT AGGGACTTTC CATTGACGTC AATGGGTGGA  
 CAAGGGTATC ATTGCGGTTA TCCCTGAAAG GTAACCTGCAG TTACCCACCT

**751** GTATTTACGG TAAACTGCC ACTTGGCAGT ACATCAAGTG TATCATATGC  
 CATAAATGCC ATTTGACGGG TGAACCGTCA TGTAGTTCAC ATAGTATACG

FIG. 10A-1



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801 CAAGTACGCC CCCTATTGAC GTCAATGACG GTAAATGGCC CGCCTGGCAT  
 GTTCATGCGG GGGATAACTG CAGTTACTGC CATTACC GG GCGGACCGTA

851 TATGCCCAGT ACATGACCTT ATGGGACTTT CCTACTTGGC AGTACATCTA  
 ATACGGGTCA TGTACTGGAA TACCCTGAAA GGATGAACCG TCATGTAGAT

901 CGTATTAGTC ATCGCTATTA CCATGGTGAT GCGGTTTTTGG CAGTACATCA  
 GCATAATCAG TAGCGATAAT GGTACCACTA CGCCAAAACC GTCATGTAGT

951 ATGGGCGTGG ATAGCGGTTT GACTCACGGG GATTTCCAAG TCTCCACCCC  
 TACCCGCACC TATCGCCAAA CTGAGTGCCC CTAAAGGTTT AGAGGTGGGG

1001 ATTGACGTCA ATGGGAGTTT GTTTTGGCAC CAAAATCAAC GGGACTTTCC  
 TAACTGCAGT TACCCTCAAA CAAAACCGTG GTTTTAGTTG CCCTGAAAGG

1051 AAAATGTCGT AACAACTCCG CCCCATTTGAC GCAAATGGGC GGTAGGCGTG  
 TTTTACAGCA TTGTTGAGGC GGGGTAACTG CGTTTACCCG CCATCCGCAC

1101 TACGGTGGGA GGTCTATATA AGCAGAGCTC GTTTAGTGAA CCGTCAGATC  
 ATGCCACCCT CCAGATATAT TCGTCTCGAG CAAATCACTT GGCAGTCTAG

1151 GCCTGGAGAC GCCATCCACG CTGTTTTGAC CTCCATAGAA GACACCGGGA  
 CGGACCTCTG CGGTAGGTGC GACAAAACCTG GAGGTATCTT CTGTGGCCCT

1201 CCGATCCAGC CTCCGCGGCC GGAACGGTG CATTGGAACG CGGATTCCC  
 GGCTAGGTCTG GAGGCGCCGG CCCTTGCCAC GTAACCTTGC GCCTAAGGGG

1251 GTGCCAAGAG TGAGATCTAC CATGGGTGCT AGGGCTTCTG TGCTGTCTGG  
 CACGGTTCTC ACTCTAGATG GTACCCACGA TCCCGAAGAC ACGACAGACC

1301 TGGTGAGCTG GACAAGTGGG AGAAGATCAG GCTGAGGCCT GGTGGCAAGA  
 ACCACTCGAC CTGTTCAACC TCTTCTAGTC CGACTCCGGA CCACCGTTCT

1351 AGAAGTACAA GCTAAAGCAC ATTGTGTGGG CCTCCAGGGA GCTGGAGAGG  
 TCTTCATGTT CGATTTCTGT TAACACACCC GGAGGTCCCT CGACCTCTCC

1401 TTTGCTGTGA ACCCTGGCCT GCTGGAGACC TCTGAGGGGT GCAGGCAGAT  
 AAACGACACT TGGGACCGGA CGACCTCTGG AGACTCCCCA CGTCCGTCTA

1451 CCTGGGCCAG CTCCAGCCCT CCCTGCAAAC AGGCTCTGAG GAGCTGAGGT  
 GGACCCGGTC GAGGTCGGGA GGGACGTTTG TCCGAGACTC CTCGACTCCA

1501 CCCTGTACAA CACAGTGGCT ACCCTGTACT GTGTGCACCA GAAGATTGAT  
 GGGACATGTT GTGTCACCGA TGGGACATGA CACACGTGGT CTTCTAATA

1551 GTGAAGGACA CCAAGGAGGC CCTGGAGAAG ATTGAGGAGG AGCAGAACAA  
 CACTTCCTGT GGTTCCCTCC GGACCTCTTC TAACTCCTCC TCGTCTTGTT

FIG. 10A-2



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1601 GTCCAAGAAG AAGGCCAGC AGGCTGCTGC TGGCACAGGC AACTCCAGCC  
 CAGGTTCTTC TTCCGGGTCG TCCGACGACG ACCGTGTCCG TTGAGGTCGG

1651 AGGTGTCCCA GAACTACCCC ATTGTGCAGA ACCTCCAGGG CCAGATGGTG  
 TCCACAGGGT CTTGATGGGG TAACACGTCT TGGAGGTCCC GGTCTACCAC

1701 CACCAGGCCA TCTCCCCCG GACCCTGAAT GCCTGGGTGA AGGTGGTGA  
 GTGGTCCGGT AGAGGGGGG CTGGGACTTA CGGACCCACT TCCACCACCT

1751 GGAGAAGGCC TTCTCCCCTG AGGTGATCCC CATGTTCTCT GCCCTGTCTG  
 CCTCTTCCGG AAGAGGGGAC TCCACTAGGG GTACAAGAGA CGGGACAGAC

1801 AGGGTGCCAC CCCCAGGAC CTGAACACCA TGCTGAACAC AGTGGGGGGC  
 TCCCACGGTG GGGGGTCCCTG GACTTGTGGT ACGACTTGTG TCACCCCCCG

1851 CATCAGGCTG CCATGCAGAT GCTGAAGGAG ACCATCAATG AGGAGGCTGC  
 GTAGTCCGAC GGTACGTCTA CGACTTCCTC TGGTAGTTAC TCCTCCGACG

1901 TGAGTGGGAC AGGCTGCATC CTGTGCACGC TGGCCCCATT GCCCCGGCC  
 ACTCACCTG TCCGACGTAG GACACGTGCG ACCGGGGTAA CGGGGGCCGG

1951 AGATGAGGGA GCCCAGGGGC TCTGACATTG CTGGCACCAC CTCCACCCTC  
 TCTACTCCCT CGGGTCCCCG AGACTGTAAC GACCGTGGTG GAGGTGGGAG

2001 CAGGAGCAGA TTGGCTGGAT GACCAACAAC CCCCCATCC CTGTGGGGGA  
 GTCCTCGTCT AACCGACCTA CTGGTTGTTG GGGGGGTAGG GACACCCCT

2051 AATCTACAAG AGGTGGATCA TCCTGGGCCT GAACAAGATT GTGAGGATGT  
 TTAGATGTTT TCCACCTAGT AGGACCCGGA CTTGTTCTAA CACTCCTACA

2101 ACTCCCCAC CTCCATCCTG GACATCAGGC AGGGCCCCAA GGAGCCCTTC  
 TGAGGGGGTG GAGGTAGGAC CTGTAGTCCG TCCCGGGGTT CCTCGGGAAG

2151 AGGGACTATG TGGACAGGTT CTACAAGACC CTGAGGGCTG AGCAGGCCTC  
 TCCCTGATAC ACCTGTCCAA GATGTTCTGG GACTCCCGAC TCGTCCGGAG

2201 CCAGGAGGTG AAGAACTGGA TGACAGAGAC CCTGCTGGTG CAGAATGCCA  
 GGTCCCTCCAC TTCTTGACCT ACTGTCTCTG GGACGACCAC GTCTTACGGT

2251 ACCCTGACTG CAAGACCATC CTGAAGGCC TGGGCCCTGC TGCCACCCTG  
 TGGGACTGAC GTTCTGGTAG GACTTCCGGG ACCCGGGACG ACGGTGGGAC

2301 GAGGAGATGA TGACAGCCTG CCAGGGGGTG GGGGGCCCTG GTCACAAGGC  
 CTCCTCTACT ACTGTCGGAC GGTCCCCAC CCCCAGGGAC CAGTGTTCGG

2351 CAGGGTGCTG GCTGAGGCCA TGTCCCAGGT GACCAACTCC GCCACCATCA  
 GTCCACGAC CGACTCCGGT ACAGGGTCCA CTGGTTGAGG CGGTGGTAGT

FIG. 10A-3



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|      |                            |                          |                           |                           |                          |
|------|----------------------------|--------------------------|---------------------------|---------------------------|--------------------------|
| 2401 | TGATGCAGAG<br>ACTACGTCTC   | GGGCAACTTC<br>CCCGTTGAAG | AGGAACCAGA<br>TCCTTGGTCT  | GGAAGACAGT<br>CCTTCTGTCA  | GAAGTGCTTC<br>CTTCACGAAG |
| 2451 | AACTGTGGCA<br>TTGACACCGT   | AGGTGGGCCA<br>TCCACCCCGT | CATTGCCAAG<br>GTAACGGTTC  | AACTGTAGGG<br>TTGACATCCC  | CCCCCAGGAA<br>GGGGGTCCTT |
| 2501 | GAAGGGCTGC<br>CTTCCCGACG   | TGGAAGTGTG<br>ACCTTCACAC | GCAAGGAGGG<br>CGTTCCTCCC  | CCACCAGATG<br>GGTGGTCTAC  | AAGGACTGCA<br>TTCCTGACGT |
| 2551 | ATGAGAGGCA<br>TACTCTCCGT   | GGCCAACTTC<br>CCGGTTGAAG | CTGGGCAAAA<br>GACCCGTTTT  | TCTGGCCCTC<br>AGACCGGGAG  | CCACAAGGGC<br>GGTGTTCCCG |
| 2601 | AGGCCTGGCA<br>TCCGGACCGT   | ACTTCCTCCA<br>TGAAGGAGGT | GTCCAGGCCT<br>CAGGTCCGGA  | GAGCCACAG<br>CTCGGGTGTC   | CCCCTCCCGA<br>GGGGAGGGCT |
| 2651 | GGAGTCCTTC<br>CCTCAGGAAG   | AGGTTTGGGG<br>TCCAAACCCC | AGGAGAAGAC<br>TCCTCTTCTG  | CACCCCAGC<br>GTGGGGGTCG   | CAGAAGCAGG<br>GTCTTCGTCC |
| 2701 | AGCCCATTTGA<br>TCGGGTA ACT | CAAGGAGCTG<br>GTTCTCGAC  | TACCCCCTGG<br>ATGGGGGACC  | CCTCCCTGAG<br>GGAGGGACTC  | GTCCCTGTTT<br>CAGGGACAAA |
| 2751 | GGCAACGACC<br>CCGTTGCTGG   | CCTCCTCCCA<br>GGAGGAGGGT | GTAAAATAAA<br>CATTTTATTT  | GCCCAGGCAG<br>CGGGCCCGTC  | ATCTGCTGTG<br>TAGACGACAC |
| 2801 | CCTTCTAGTT<br>GGAAGATCAA   | GCCAGCCATC<br>CGGTCGGTAG | TGTTGTTTGC<br>ACAACAAACG  | CCCTCCCCCG<br>GGGAGGGGGC  | TGCCTTCCTT<br>ACGGAAGGAA |
| 2851 | GACCCTGGAA<br>CTGGGACCTT   | GGTGCCACTC<br>CCACGGTGAG | CCACTGTCCT<br>GGTGACAGGA  | TTCCTAATAA<br>AAGGATTATT  | AATGAGGAAA<br>TTACTCCTTT |
| 2901 | TTGCATCGCA<br>AACGTAGCGT   | TTGTCTGAGT<br>AACAGACTCA | AGGTGTCATT<br>TCCACAGTAA  | CTATTCTGGG<br>GATAAGACCC  | GGGTGGGGTG<br>CCCACCCAC  |
| 2951 | GGGCAGGACA<br>CCCGTCCTGT   | GCAAGGGGGA<br>CGTTCCCCCT | GGATTGGGAA<br>CCTAACCCCTT | GACAATAGCA<br>CTGTTATCGT  | GGCATGCTGG<br>CCGTACGACC |
| 3001 | GGATGCGGTG<br>CCTACGCCAC   | GGCTCTATGG<br>CCGAGATACC | CCGATCGGCG<br>GGCTAGCCGC  | CGCCGTACTG<br>GCGGCATGAC  | AAATGTGTGG<br>TTTACACACC |
| 3051 | GCGTGGCTTA<br>CGCACCGAAT   | AGGGTGGGAA<br>TCCCACCCTT | AGAATATATA<br>TCTTATATAT  | AGGTGGGGGT<br>TCCACCCCA   | CTTATGTAGT<br>GAATACATCA |
| 3101 | TTTGTATCTG<br>AAACATAGAC   | TTTTGCAGCA<br>AAAACGTCGT | GCCGCCGCCG<br>CGGCGGCGGC  | CCATGAGCAC<br>GGTACTCGTG  | CAACTCGTTT<br>GTTGAGCAAA |
| 3151 | GATGGAAGCA<br>CTACCTTCGT   | TTGTGAGCTC<br>AACACTCGAG | ATATTTGACA<br>TATAAACTGT  | ACGCGCATGC<br>TGC GCGTACG | CCCCATGGGC<br>GGGGTACCCG |

FIG. 10A-4



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3201 CGGGGTGCGT CAGAATGTGA TGGGCTCCAG CATTGATGGT CGCCCCGTCC  
 GCCCCACGCA GTCTTACACT ACCCGAGGTC GTAACCTACCA GCGGGGCAGG

3251 TGCCCGCAA CTCTACTACC TTGACCTACG AGACCGTGTC TGGAACGCCG  
 ACGGGCGTTT GAGATGATGG AACTGGATGC TCTGGCACAG ACCTTGCGGC

3301 TTGGAGACTG CAGCCTCCGC CGCCGCTTCA GCCGCTGCAG CCACCGCCCCG  
 AACCTCTGAC GTCGGAGGCG GCGGCGAAGT CGGCGACGTC GGTGGCGGGC

3351 CGGGATTGTG ACTGACTTTG CTTTCCTGAG CCCGCTTGCA AACAGTGCAG  
 GCCCTAACAC TGACTGAAAC GAAAGGACTC GGGCGAACGT TTGTCACGTC

3401 CTTCCCGTTC ATCCGCCCGC GATGACAAGT TGACGGCTCT TTTGGCACAA  
 GAAGGGCAAG TAGGCGGGCG CTACTGTTCA ACTGCCGAGA AAACCGTGTT

3451 TTGGATTCTT TGACCCGGGA ACTTAATGTC GTTTCCTCAGC AGCTGTTGGA  
 AACCTAAGAA ACTGGGCCCT TGAATTACAG CAAAGAGTCG TCGACAACCT

3501 TCTGCGCCAG CAGGTTTCTG CCCTGAAGGC TTCCTCCCCT CCAATGCGG  
 AGACGCGGTC GTCCAAAGAC GGGACTTCCG AAGGAGGGGA GGGTTACGCC

3551 TTTAAACAT AAATAAAAA CCAGACTCTG TTTGGATTTG GATCAAGCAA  
 AAATTTTGTA TTTATTTTTT GGTCTGAGAC AAACCTAAC CTAGTTCGTT

3601 GTGCTTTGCT GTCTTTATTT AGGGGTTTTG CGCGCGCGGT AGGCCCGGGA  
 CACAGAACGA CAGAAATAA TCCCCAAAAC GCGCGCGCCA TCCGGGCCCT

3651 CCAGCGGTCT CGGTCGTTGA GGGTCCTGTG TATTTTTTCC AGGACGTGGT  
 GGTCGCCAGA GCCAGCAACT CCCAGGACAC ATAAAAAAGG TCCTGCACCA

3701 AAAGGTGACT CTGGATGTTT AGATACATGG GCATAAGCCC GTCTCTGGGG  
 TTTCCACTGA GACCTACAAG TCTATGTACC CGTATTCGGG CAGAGACCCC

3751 TGGAGGTAGC ACCACTGCAG AGCTTCATGC TGCGGGGTGG TGTTGTAGAT  
 ACCTCCATCG TGGTGACGTC TCGAAGTACG ACGCCCCACC ACAACATCTA

3801 GATCCAGTCG TAGCAGGAGC GCTGGGCGTG GTGCCTAAAA ATGTCTTTCA  
 CTAGGTCAGC ATCGTCCTCG CGACCCGCAC CACGGATTTT TACAGAAAGT

3851 GTAGCAAGCT GATTGCCAGG GGCAGGCCCT TGGTGTAAGT GTTTACAAAG  
 CATCGTTCGA CTAACGGTCC CCGTCCGGGA ACCACATTCA CAAATGTTTC

3901 CGGTTAAGCT GGGATGGGTG CATACTGGG GATATGAGAT GCATCTTGGA  
 GCCAATTCGA CCCTACCCAC GTATGCACCC CTATACTCTA CGTAGAACCT

3951 CTGTATTTTT AGGTTGGCTA TGTTCCAGC CATATCCCTC CGGGGATTCA  
 GACATAAAAA TCCAACCGAT ACAAGGGTCG GTATAGGGAG GCCCCTAAGT

FIG. 10A-5



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4001 TGTTGTGCAG AACACCAGC ACAGTGTATC CGGTGCACTT GGGAAATTTG  
 ACAACACGTC TTGGTGGTCG TGTCACATAG GCCACGTGAA CCCTTTAAAC

4051 TCATGTAGCT TAGAAGGAAA TCGGTGGAAG AACTTGGAGA CGCCCTTGTG  
 AGTACATCGA ATCTTCCTTT ACGCACCTTC TTGAACCTCT GCGGGAACAC

4101 ACCTCCAAGA TTTTCCATGC ATTCGTCCAT AATGATGGCA ATGGGCCCAC  
 TGGAGGTTCT AAAAGGTACG TAAGCAGGTA TTACTIONCGT TACCCGGGTG

4151 GGGCGGCGGC CTGGGCGAAG ATATTTCTGG GATCACTAAC GTCATAGTTG  
 CCCGCCGCCG GACCCGCTTC TATAAAGACC CTAGTGATTG CAGTATCAAC

4201 TGTTCCAGGA TGAGATCGTC ATAGGCCATT TTTACAAAGC GCGGGCGGAG  
 ACAAGGTCCT ACTCTAGCAG TATCCGGTAA AAATGTTTCG CGCCCGCCTC

4251 GGTGCCAGAC TCGGGTATAA TGGTTCCATC CGGCCAGGG GCGTAGTTAC  
 CCACGGTCTG ACGCCATATT ACCAAGGTAG GCCGGGTCCC CGCATCAATG

4301 CCTCACAGAT TTGCATTTCC CACGCTTTGA GTTCAGATGG GGGGATCATG  
 GGAGTGTCTA AACGTAAAGG GTGCGAAACT CAAGTCTACC CCCCTAGTAC

4351 TCTACCTGCG GGGCGATGAA GAAAACGGTT TCCGGGGTAG GGGAGATCAG  
 AGATGGACGC CCCGCTACTT CTTTTGCCAA AGGCCCATC CCCTCTAGTC

4401 CTGGGAAGAA AGCAGGTTCC TGAGCAGCTG CGACTTACCG CAGCCGGTGG  
 GACCCTTCTT TCGTCCAAGG ACTCGTCGAC GCTGAATGGC GTCGGCCACC

4451 GCCCGTAAAT CACACCTATT ACCGGCTGCA ACTGGTAGTT AAGAGAGCTG  
 CGGGCATTTA GTGTGGATAA TGGCCGACGT TGACCATCAA TTCTCTCGAC

4501 CAGCTGCCGT CATCCCTGAG CAGGGGGGCC ACTTCGTTAA GCATGTCCCT  
 GTCGACGGCA GTAGGGACTC GTCCCCCGG TGAAGCAATT CGTACAGGGA

4551 GACTCGCATG TTTTCCCTGA CCAAATCCGC CAGAAGGCGC TCGCCGCCCA  
 CTGAGCGTAC AAAAGGGACT GGTTTAGGCG GTCTTCCGCG AGCGGCGGGT

4601 GCGATAGCAG TTCTTGCAAG GAAGCAAAGT TTTTCAACGG TTTGAGACCG  
 CGCTATCGTC AAGAACGTTC CTTCGTTTCA AAAAGTTGCC AACTCTGGC

4651 TCCGCCGTAG GCATGCTTTT GAGCGTTTGA CCAAGCAGTT CCAGGCGGTC  
 AGGCGGCATC CGTACGAAAA CTCGCAAACCT GGTTTCGTCAA GGTCCGCCAG

4701 CCACAGCTCG GTCACCTGCT CTACGGCATC TCGATCCAGC ATATCTCCTC  
 GGTGTCGAGC CAGTGGACGA GATGCCGTAG AGCTAGGTCG TATAGAGGAG

4751 GTTTCGCGGG TTGGGGCGGC TTTCGCTGTA CGGCAGTAGT CGGTGCTCGT  
 CAAAGCGCCC AACCCCGCCG AAAGCGACAT GCCGTCATCA GCCACGAGCA

FIG. 10A-6



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4801 CCAGACGGGC CAGGGTCATG TCTTTCACG GCGCAGGGT CCTCGTCAGC  
 GGTCTGCCCG GTCCAGTAC AGAAAGGTGC CCGCGTCCA GGAGCAGTCG

4851 GTAGTCTGGG TCACGGTGAA GGGGTGCGCT CCGGGCTGCG CGCTGGCCAG  
 CATCAGACCC AGTGCCACTT CCCCACGCGA GGCCCGACGC GCGACCGGTC

4901 GGTGCGCTTG AGGCTGGTCC TGCTGGTGCT GAAGCGCTGC CGGTCTTCGC  
 CCACGCGAAC TCCGACCAGG ACGACCACGA CTTGCGGACG GCCAGAAGCG

4951 CCTGCGCGTC GGCCAGGTAG CATTGACCA TGGTGTGATA GTCCAGCCCC  
 GGACGCGCAG CCGGTCCATC GTAAACTGGT ACCACAGTAT CAGGTCGGGG

5001 TCCGCGGCGT GGCCCTTGGC GCGCAGCTTG CCCTTGGAGG AGGCGCCGCA  
 AGGCGCCGCA CCGGGAACCG CGCGTCGAAC GGAACCTCC TCCGCGGCGT

5051 CGAGGGGCAG TGCAGACTTT TGAGGGCGTA GAGCTTGGGC GCGAGAAATA  
 GCTCCCCGTC ACGTCTGAAA ACTCCCGCAT CTCGAACCCG CGCTCTTTAT

5101 CCGATTCCGG GGAGTAGGCA TCCGCGCCGC AGGCCCCGCA GACGGTCTCG  
 GGCTAAGGCC CCTCATCCGT AGGCGCGGCG TCCGGGGCGT CTGCCAGAGC

5151 CATTCCACGA GCCAGGTGAG CTCTGGCCGT TCGGGGTCAA AAACCAGGTT  
 GTAAGGTGCT CGGTCCACTC GAGACCGGCA AGCCCCAGTT TTTGGTCCAA

5201 TCCCCCATGC TTTTGTGATG GTTCTTACC TCTGGTTTCC ATGAGCCGGT  
 AGGGGGTACG AAAAACTACG CAAAGAATGG AGACCAAAGG TACTCGGCCA

5251 GTCCACGCTC GGTGACGAAA AGGCTGTCCG TGTCCCCGTA TACAGACTTG  
 CAGGTGCGAG CCACTGCTTT TCCGACAGGC ACAGGGGCAT ATGTCTGAAC

5301 AGAGGCCTGT CCTCGAGCGG TGTTCCGCGG TCCTCCTCGT ATAGAACTC  
 TCTCCGGACA GGAGCTCGCC ACAAGGCGCC AGGAGGAGCA TATCTTTGAG

5351 GGACCACTCT GAGACAAAGG CTCGCGTCCA GGCCAGCACG AAGGAGGCTA  
 CCTGGTGAGA CTCTGTTTCC GAGCGCAGGT CCGGTGCTGC TTCTCCGAT

5401 AGTGGGAGGG GTAGCGGTCG TTGTCCACTA GGGGGTCCAC TCGCTCCAGG  
 TCACCCTCCC CATCGCCAGC AACAGGTGAT CCCCAGGTG AGCGAGGTCC

5451 GTGTGAAGAC ACATGTCGCC CTCTTCGGCA TCAAGGAAGG TGATTGGTTT  
 CACTTCTG TGTACAGCGG GAGAAGCCGT AGTTCCTTCC ACTAACCAA

5501 GTAGGTGTAG GCCACGTGAC CGGGTGTTCC TGAAGGGGGG CTATAAAAGG  
 CATCCACATC CGGTGCACTG GCCACAAGG ACTTCCCCC GATATTTTCC

5551 GGGTGGGGGC GCGTTCGTCC TCACTCTCTT CCGCATCGCT GTCTGCGAGG  
 CCCACCCCG CGCAAGCAGG AGTGAGAGAA GCGGTAGCGA CAGACGCTCC

FIG. 10A-7



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5601 GCCAGCTGTT GGGGTGAGTA CTCCTCTGA AAAGCGGGCA TGA CTCTGC  
 CGGTCGACAA CCCACTCAT GAGGGAGACT TTCGCCCGT ACTGAAGACG

5651 GCTAAGATTG TCAGTTTCCA AAAACGAGGA GGATTTGATA TTCACCTGGC  
 CGATTCTAAC AGTCAAAGGT TTTTGCTCCT CCTAAACTAT AAGTGGACCG

5701 CCGCGGTGAT GCCTTTGAGG GTGGCCGCAT CCATCTGGTC AGAAAAGACA  
 GGCGCCACTA CGGAACTCC CACCGGCGTA GG TAGACCAG TCTTTTCTGT

5751 ATCTTTTGT TGTCAAGCTT GGTGGCAAAC GACCCGTAGA GGGCGTTGGA  
 TAGAAAAACA ACAGTTCGAA CCACCGTTTG CTGGGCATCT CCCGCAACCT

5801 CAGCAACTTG GCGATGGAGC GCAGGGTTTG GTTTTTGTCG CGATCGGCGC  
 GTCGTTGAAC CGCTACCTCG CGTCCCAAAC CAAAACAGC GCTAGCCGCG

5851 GCTCCTTGGC CGCGATGTTT AGCTGCACGT ATTCGCGCGC AACGCACCGC  
 CGAGGAACCG GCGCTACAAA TCGACGTGCA TAAGCGCGCG TTGCGTGGCG

5901 CATTCGGGAA AGACGGTGGT GCGCTCGTCG GGCACCAGGT GCACGCGCCA  
 GTAAGCCCTT TCTGCCACCA CGCGAGCAGC CCGTGGTCCA CGTGC GCGGT

5951 ACCGCGGTTG TGCAGGGTGA CAAGGTCAAC GCTGGTGGCT ACCTCTCCGC  
 TGGCGCCAAC ACGTCCCACT GTTCCAGTTG CGACCACCGA TGGAGAGGCG

6001 GTAGGCGCTC GTTGGTCCAG CAGAGGCGGC CGCCCTTGCG CGAGCAGAAT  
 CATCCGCGAG CAACCAGGTC GTCTCCGCCG GCGGGAACGC GCTCGTCTTA

6051 GCGGGTAGGG GGTCTAGCTG CGTCTCGTCC GGGGGGTCTG CGTCCACGGT  
 CCGCCATCCC CCAGATCGAC GCAGAGCAGG CCCCCAGAC GCAGGTGCCA

6101 AAAGACCCCG GGCAGCAGGC GCGCGTCGAA GTAGTCTATC TTGCATCCTT  
 TTTCTGGGGC CCGTCGTCCG CGCGCAGCTT CATCAGATAG AACGTAGGAA

6151 GCAAGTCTAG CGCCTGCTGC CATGCGCGGG CGGCAAGCGC GCGCTCGTAT  
 CGTTCAGATC GCGGACGACG GTACGCGCCC GCCGTTGCGC CGCGAGCATA

6201 GGGTTGAGTG GGGGACCCCA TGGCATGGGG TGGGTGAGCG CGGAGGCGTA  
 CCCAACTCAC CCCCTGGGGT ACCGTACCCC ACCCACTCGC GCCTCCGCAT

6251 CATGCCGCAA ATGTCGTAAA CGTAGAGGGG CTCTCTGAGT ATTCCAAGAT  
 GTACGGCGTT TACAGCATTT GCATCTCCCC GAGAGACTCA TAAGGTTCTA

6301 ATGTAGGGTA GCATCTTCCA CCGCGGATGC TGGCGCGCAC GTAATCGTAT  
 TACATCCCAT CGTAGAAGGT GGCGCCTACG ACCGCGCGTG CATTAGCATA

6351 AGTTCGTGCG AGGGAGCGAG GAGGTCGGGA CCGAGGTTGC TACGGGCGGG  
 TCAAGCACGC TCCCTCGCTC CTCCAGCCCT GGCTCCAACG ATGCCCGCCC

FIG. 10A-8



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6401 CTGCTCTGCT CGGAAGACTA TCTGCCTGAA GATGGCATGT GAGTTGGATG  
 GACGAGACGA GCCTTCTGAT AGACGGACTT CTACCGTACA CTCAACCTAC

6451 ATATGGTTGG ACGCTGGAAG ACGTTGAAGC TGGCGTCTGT GAGACCTACC  
 TATACCAACC TCGGACCTTC TGCAACTTCC ACCGCAGACA CTCTGGATGG

6501 GCGTCACGCA CGAAGGAGGC GTAGGAGTCG CGCAGCTTGT TGACCAGCTC  
 CGCAGTGCGT GCTTCCTCCG CATCCTCAGC GCGTCGAACA ACTGGTCGAG

6551 GGCGGTGACC TGCACGTCTA GGGCGCAGTA GTCCAGGGTT TCCTTGATGA  
 CCGCCACTGG ACGTGCAGAT CCCGCGTCAT CAGGTCCCAA AGGAACTACT

6601 TGTCATACTT ATCCTGTCCC TTTTTTTTCC ACAGCTCGCG GTTGAGGACA  
 ACAGTATGAA TAGGACAGGG AAAAAAAGG TGTCGAGCGC CAACTCCTGT

6651 AACTCTTCGC GGTCTTTCCA GTACTCTTGG ATCGGAAACC CGTCGGCCTC  
 TTGAGAAGCG CCAGAAAGGT CATGAGAACC TAGCCTTTGG GCAGCCGGAG

6701 CGAACGGTAA GAGCCTAGCA TGTAGAACTG GTTGACGGCC TGGTAGGCGC  
 GCTTGCCATT CTCGGATCGT ACATCTTGAC CAACTGCCGG ACCATCCGCG

6751 AGCATCCCTT TTCTACGGGT AGCGCGTATG CCTGCGCGGC CTTCCGGAGC  
 TCGTAGGGAA AAGATGCCCA TCGCGCATAC GGACGCGCCG GAAGGCCTCG

6801 GAGGTGTGGG TGAGCGCAA GGTGTCCCTG ACCATGACTT TGAGGTACTG  
 CTCCACACCC ACTCGCGTTT CCACAGGGAC TGGTACTGAA ACTCCATGAC

6851 GTATTTGAAG TCAGTGTCGT CGCATCCGCC CTGCTCCCAG AGCAAAAAGT  
 CATAAACTTC AGTCACAGCA GCGTAGGCGG GACGAGGGTC TCGTTTTTCA

6901 CCGTGCGCTT TTTGGAACGC GGATTTGGCA GGGCGAAGGT GACATCGTTG  
 GGCACGCGAA AAACCTTGCG CCTAAACCGT CCCGCTTCCA CTGTAGCAAC

6951 AAGAGTATCT TTCCCGCGCG AGGCATAAAG TTGCGTGTGA TCGGGAAGGG  
 TTCTCATAGA AAGGGCGCGC TCCGTATTTT AACGCACACT ACGCCTTCCC

7001 TCCCGGCACC TCGGAACGGT TGTTAATTAC CTGGGCGGCG AGCACGATCT  
 AGGGCCGTGG AGCCTTGCCA ACAATTAATG GACCCGCCGC TCGTGCTAGA

7051 CGTCAAAGCC GTTGATGTTG TGGCCACAA TGTAAGTTC CAAGAAGCGC  
 GCAGTTTCGG CAACTACAAC ACCGGGTGTT ACATTTCAAG GTTCTTCGCG

7101 GGGATGCCCT TGATGGAAGG CAATTTTTTA AGTTCCTCGT AAGTGAGCTC  
 CCCTACGGGA ACTACCTTCC GTTAAAAAAT TCAAGGAGCA TCCACTCGAG

7151 TTCAGGGGAG CTGAGCCCGT GCTCTGAAAG GGCCAGTCT GCAAGATGAG  
 AAGTCCCCTC GACTCGGGCA CGAGACTTTC CCGGGTCAGA CGTTCTACTC

FIG. 10A-9



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|      |                           |                          |                          |                           |                          |
|------|---------------------------|--------------------------|--------------------------|---------------------------|--------------------------|
| 7201 | GGTTGGAAGC<br>CCAACCTTCG  | GACGAATGAG<br>CTGCTTACTC | CTCCACAGGT<br>GAGGTGTCCA | CACGGGCCAT<br>GTGCCCGGTA  | TAGCATTTGC<br>ATCGTAAACG |
| 7251 | AGGTGGTCGC<br>TCCACCAGCG  | GAAAGGTCCT<br>CTTTCAGGA  | AAACTGGCGA<br>TTTGACCGCT | CCTATGGCCA<br>GGATACCGGT  | TTTTTTCTGG<br>AAAAAAGACC |
| 7301 | GGTGATGCAG<br>CCTACTACGTC | TAGAAGGTAA<br>ATCTTCCATT | GCGGGTCTTG<br>CGCCAGAAC  | TTCCCAGCGG<br>AAGGGTCGCC  | TCCCATCCAA<br>AGGGTAGGTT |
| 7351 | GGTTCGCGGC<br>CCAAGCGCCG  | TAGGTCTCGC<br>ATCCAGAGCG | GCGGCAGTCA<br>CGCCGTCAGT | CTAGAGGCTC<br>GATCTCCGAG  | ATCTCCGCCG<br>TAGAGGCGGC |
| 7401 | AACTTCATGA<br>TTGAAGTACT  | CCAGCATGAA<br>GGTCGTACTT | GGGCACGAGC<br>CCCGTGCTCG | TGCTTCCCAA<br>ACGAAGGGTT  | AGGCCCCCAT<br>TCCGGGGGTA |
| 7451 | CCAAGTATAG<br>GGTTCATATC  | GTCTCTACAT<br>CAGAGATGTA | CGTAGGTGAC<br>GCATCCACTG | AAAGAGACGC<br>TTTCTCTGCG  | TCGGTGCGAG<br>AGCCACGCTC |
| 7501 | GATGCGAGCC<br>CTACGCTCGG  | GATCGGGAAG<br>CTAGCCCTTC | AACTGGATCT<br>TTGACCTAGA | CCCGCCACCA<br>GGCGGTGGT   | ATTGGAGGAG<br>TAACCTCCTC |
| 7551 | TGGCTATTGA<br>ACCGATAACT  | TGTGGTGAAA<br>ACACCACTTT | GTAGAAGTCC<br>CATCTTCAGG | CTGCGACGGG<br>GACGCTGCC   | CCGAACACTC<br>GGCTTGTGAG |
| 7601 | GTGCTGGCTT<br>CACGACCGAA  | TTGTAAAAAC<br>AACATTTTTG | GTGCGCAGTA<br>CACGCGTCAT | CTGGCAGCGG<br>GACCGTCGCC  | TGCACGGGCT<br>ACGTGCCCGA |
| 7651 | GTACATCCTG<br>CATGTAGGAC  | CACGAGGTTG<br>GTGCTCCAAC | ACCTGACGAC<br>TGGACTGCTG | CGCGCACAAAG<br>GCGCGTGTTT | GAAGCAGAGT<br>CTTCGTCTCA |
| 7701 | GGGAATTTGA<br>CCCTTAAACT  | GCCCCTCGCC<br>CGGGGAGCGG | TGGCGGGTTT<br>ACCGCCCAA  | GGCTGGTGGT<br>CCGACCACCA  | CTTCTACTTC<br>GAAGATGAAG |
| 7751 | GGCTGCTTGT<br>CCGACGAACA  | CCTTGACCGT<br>GGAAGTGGCA | CTGGCTGCTC<br>GACCGACGAG | GAGGGGAGTT<br>CTCCCCTCAA  | ACGGTGGATC<br>TGCCACCTAG |
| 7801 | GGACCACCAC<br>CCTGGTGGTG  | GCCGCGCGAG<br>CGGCGCGCTC | CCCAAAGTCC<br>GGGTTTCAGG | AGATGTCCGC<br>TCTACAGGCG  | GCGCGGCGGT<br>CGCGCCGCCA |
| 7851 | CGGAGCTTGA<br>GCCTCGAACT  | TGACAACATC<br>ACTGTTGTAG | GCGCAGATGG<br>CGCGTCTACC | GAGCTGTCCA<br>CTCGACAGGT  | TGGTCTGGAG<br>ACCAGACCTC |
| 7901 | CTCCCGCGGC<br>GAGGGCGCCG  | GTCAGGTCAG<br>CAGTCCAGTC | GCGGGAGCTC<br>CGCCCTCGAG | CTGCAGGTTT<br>GACGTCCAAA  | ACCTCGCATA<br>TGGAGCGTAT |
| 7951 | GACGGGTCAG<br>CTGCCAGTC   | GGCGCGGGCT<br>CCGCGCCCGA | AGATCCAGGT<br>TCTAGGTCCA | GATACCTAAT<br>CTATGGATTA  | TTCCAGGGGC<br>AAGGTCCCCG |

FIG. 10A-10



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|      |            |            |            |             |            |
|------|------------|------------|------------|-------------|------------|
| 8001 | TGGTTGGTGG | CGGCGTCGAT | GGCTTGCAAG | AGGCCGCATC  | CCCGCGGCGC |
|      | ACCAACCACC | GCCGCAGCTA | CCGAACGTTT | TCCGGCGTAG  | GGGCGCCGCG |
| 8051 | GACTACGGTA | CCGCGCGGCG | GGCGGTGGGC | CGCGGGGGTG  | TCCTTGGATG |
|      | CTGATGCCAT | GGCGCGCCGC | CCGCCACCCG | GCGCCCCCAC  | AGGAACCTAC |
| 8101 | ATGCATCTAA | AAGCGGTGAC | GCGGGCGAGC | CCCCGGAGGT  | AGGGGGGGCT |
|      | TACGTAGATT | TTCGCCACTG | CGCCCGCTCG | GGGGCCTCCA  | TCCCCCCCGA |
| 8151 | CCGGACCCGC | CGGGAGAGGG | GGCAGGGGCA | CGTCGGCGCC  | GCGCGCGGGC |
|      | GGCCTGGGCG | GCCCTCTCCC | CCGTCCCCGT | GCAGCCGCGG  | CGCGCGCCCC |
| 8201 | AGGAGCTGGT | GCTGCGCGCG | TAGGTTGCTG | GCGAACGCGA  | CGACGCGGCG |
|      | TCCTCGACCA | CGACGCGCGC | ATCCAACGAC | CGCTTGCGCT  | GCTGCGCCGC |
| 8251 | GTTGATCTCC | TGAATCTGGC | GCCTCTGCGT | GAAGACGACG  | GGCCCGGTGA |
|      | CAACTAGAGG | ACTTAGACCG | CGGAGACGCA | CTTCTGCTGC  | CCGGGCCACT |
| 8301 | GCTTGAACCT | GAAAGAGAGT | TCGACAGAAT | CAATTTTCGGT | GTCGTTGACG |
|      | CGAACTTGGA | CTTTCTCTCA | AGCTGTCTTA | GTTAAAGCCA  | CAGCAACTGC |
| 8351 | GCGGCCTGGC | GCAAAATCTC | CTGCACGTCT | CCTGAGTTGT  | CTTGATAGGC |
|      | CGCCGGACCG | CGTTTTAGAG | GACGTGCAGA | GGACTCAACA  | GAACTATCCG |
| 8401 | GATCTCGGCC | ATGAACTGCT | CGATCTCTTC | CTCCTGGAGA  | TCTCCGCGTC |
|      | CTAGAGCCGG | TACTTGACGA | GCTAGAGAAG | GAGGACCTCT  | AGAGGCGCAG |
| 8451 | CGGCTCGCTC | CACGGTGGCG | GCGAGGTCGT | TGGAAATGCG  | GGCCATGAGC |
|      | GCCGAGCGAG | GTGCCACCGC | CGCTCCAGCA | ACCTTTACGC  | CCGGTACTCG |
| 8501 | TGCGAGAAGG | CGTTGAGGCC | TCCCTCGTTC | CAGACGCGGC  | TGTAGACCAC |
|      | ACGCTCTTCC | GCAACTCCGG | AGGGAGCAAG | GTCTGCGCCG  | ACATCTGGTG |
| 8551 | GCCCCCTTCG | GCATCGCGGG | CGCGCATGAC | CACCTGCGCG  | AGATTGAGCT |
|      | CGGGGGAAGC | CGTAGCGCCC | GCGCGTACTG | GTGGACGCGC  | TCTAACTCGA |
| 8601 | CCACGTGCCG | GGCGAAGACG | GCGTAGTTTC | GCAGGCGCTG  | AAAGAGGTAG |
|      | GGTGCACGGC | CCGCTTCTGC | CGCATCAAAG | CGTCCGCGAC  | TTTCTCCATC |
| 8651 | TTGAGGGTGG | TGGCGGTGTG | TTCTGCCACG | AAGAAGTACA  | TAACCCAGCG |
|      | AACTCCCACC | ACCGCCACAC | AAGACGGTGC | TTCTTCATGT  | ATTGGGTTCG |
| 8701 | TCGCAACGTG | GATTCGTTGA | TATCCCCCAA | GGCCTCAAGG  | CGCTCCATGG |
|      | AGCGTTGCAC | CTAAGCAACT | ATAGGGGGTT | CCGGAGTTCC  | GCGAGGTACC |
| 8751 | CCTCGTAGAA | GTCCACGGCG | AAGTTGAAAA | ACTGGGAGTT  | GCGCGCCGAC |
|      | GGAGCATCTT | CAGGTGCCGC | TTCAACTTTT | TGACCCTCAA  | CGCGCGGCTG |

FIG. 10A-11



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|      |                          |                          |                          |                          |                          |
|------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|
| 8801 | ACGGTTAACT<br>TGCCAATTGA | CCTCCTCCAG<br>GGAGGAGGTC | AAGACGGATG<br>TTCTGCCTAC | AGCTCGGCGA<br>TCGAGCCGCT | CAGTGTCGCG<br>GTCACAGCGC |
| 8851 | CACCTCGCGC<br>GTGGAGCGCG | TCAAAGGCTA<br>AGTTTCCGAT | CAGGGGCCTC<br>GTCCCCGGAG | TTCTTCTTCT<br>AGAAGAAGA  | TCAATCTCCT<br>AGTTAGAGGA |
| 8901 | CTTCCATAAG<br>GAAGGTATTC | GGCCTCCCCT<br>CCGGAGGGGA | TCTTCTTCTT<br>AGAAGAAGAA | CTGGCGGCGG<br>GACCGCCGCC | TGGGGGAGGG<br>ACCCCCTCCC |
| 8951 | GGGACACGGC<br>CCCTGTGCCG | GGCGACGACG<br>CCGCTGCTGC | GCGCACCGGG<br>CGCGTGGCCC | AGGCGGTCGA<br>TCCGCCAGCT | CAAAGCGCTC<br>GTTTCGCGAG |
| 9001 | GATCATCTCC<br>CTAGTAGAGG | CCGCGGCGAC<br>GGCGCCGCTG | GGCGCATGGT<br>CCGCGTACCA | CTCGGTGACG<br>GAGCCACTGC | GCGCGGCCGT<br>CGCGCCGGCA |
| 9051 | TCTCGCGGGG<br>AGAGCGCCCC | GCGCAGTTGG<br>CGCGTCAACC | AAGACGCCGC<br>TTCTGCGGCG | CCGTCATGTC<br>GGCAGTACAG | CCGGTTATGG<br>GGCCAATACC |
| 9101 | GTTGGCGGGG<br>CAACCGCCCC | GGCTGCCATG<br>CCGACGGTAC | CGGCAGGGAT<br>GCCGTCCCTA | ACGGCGCTAA<br>TGCCGCGATT | CGATGCATCT<br>GCTACGTAGA |
| 9151 | CAACAATTGT<br>GTTGTTAACA | TGTGTAGGTA<br>ACACATCCAT | CTCCGCCGCC<br>GAGGCGGCGG | GAGGGACCTG<br>CTCCCTGGAC | AGCGAGTCCG<br>TCGCTCAGGC |
| 9201 | CATCGACCGG<br>GTAGCTGGCC | ATCGGAAAAC<br>TAGCCTTTTG | CTCTCGAGAA<br>GAGAGCTCTT | AGGCGTCTAA<br>TCCGCAGATT | CCAGTCACAG<br>GGTCAGTGTC |
| 9251 | TCGCAAGGTA<br>AGCGTTCCAT | GGCTGAGCAC<br>CCGACTCGTG | CGTGGCGGGC<br>GCACCGCCCC | GGCAGCGGGC<br>CCGTCGCCCC | GGCGGTCGGG<br>CCGCCAGCCC |
| 9301 | GTTGTTTCTG<br>CAACAAAGAC | GCGGAGGTGC<br>CGCCTCCACG | TGCTGATGAT<br>ACGACTACTA | GTAATTAAAG<br>CATTAAATTC | TAGGCGGTCT<br>ATCCGCCAGA |
| 9351 | TGAGACGGCG<br>ACTCTGCCGC | GATGGTCGAC<br>CTACCAGCTG | AGAAGCACCA<br>TCTTCGTGGT | TGTCCTTGGG<br>ACAGGAACCC | TCCGGCCTGC<br>AGGCCGGACG |
| 9401 | TGAATGCGCA<br>ACTTACGCGT | GGCGGTCGGC<br>CCGCCAGCCG | CATGCCCCAG<br>GTACGGGGTC | GCTTCGTTTT<br>CGAAGCAAAA | GACATCGGCG<br>CTGTAGCCGC |
| 9451 | CAGGTCTTTG<br>GTCCAGAAAC | TAGTAGTCTT<br>ATCATCAGAA | GCATGAGCCT<br>CGTACTCGGA | TTCTACCGGC<br>AAGATGGCCG | ACTTCTTCTT<br>TGAAGAAGAA |
| 9501 | CTCCTTCCTC<br>GAGGAAGGAG | TTGTCCTGCA<br>AACAGGACGT | TCTCTTGCA<br>AGAGAACGTA  | CTATCGCTGC<br>GATAGCGACG | GGCGGCGGCG<br>CCGCCGCCGC |
| 9551 | GAGTTTGGCC<br>CTCAAACCGG | GTAGGTGGCG<br>CATCCACCGC | CCCTCTTCCT<br>GGGAGAAGGA | CCCATGCGTG<br>GGGTACGCAC | TGACCCCGAA<br>ACTGGGGCTT |

FIG. 10A-12



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|       |                          |                          |                            |                          |                          |
|-------|--------------------------|--------------------------|----------------------------|--------------------------|--------------------------|
| 9601  | GCCCCTCATC<br>CGGGGAGTAG | GGCTGAAGCA<br>CCGACTTCGT | GGGCTAGGTC<br>CCCGATCCAG   | GGCGACAACG<br>CCGCTGTTGC | CGCTCGGCTA<br>GCGAGCCGAT |
| 9651  | ATATGGCCTG<br>TATACCGGAC | CTGCACCTGC<br>GACGTGGACG | GTGAGGGTAG<br>CACTCCCATC   | ACTGGAAGTC<br>TGACCTTCAG | ATCCATGTCC<br>TAGGTACAGG |
| 9701  | ACAAAGCGGT<br>TGTTTCGCCA | GGTATGCGCC<br>CCATACGCGG | CGTGTTGATG<br>GCACAACACTAC | GTGTAAGTGC<br>CACATTCACG | AGTTGGCCAT<br>TCAACCGGTA |
| 9751  | AACGGACCAG<br>TTGCCTGGTC | TTAACGGTCT<br>AATTGCCAGA | GGTGACCCGG<br>CCACTGGGCC   | CTGCGAGAGC<br>GACGCTCTCG | TCGGTGTACC<br>AGCCACATGG |
| 9801  | TGAGACGCGA<br>ACTCTGCGCT | GTAAGCCCTC<br>CATTCGGGAG | GAGTCAAATA<br>CTCAGTTTAT   | CGTAGTCGTT<br>GCATCAGCAA | GCAAGTCCGC<br>CGTTCAGGCG |
| 9851  | ACCAGGTACT<br>TGGTCCATGA | GGTATCCCAC<br>CCATAGGGTG | CAAAAAGTGC<br>GTTTTTCACG   | GGCGGCGGCT<br>CCGCCGCCGA | GGCGGTAGAG<br>CCGCCATCTC |
| 9901  | GGGCCAGCGT<br>CCCGGTCGCA | AGGGTGGCCG<br>TCCCACCGGC | GGGCTCCGGG<br>CCCGAGGCC    | GGCGAGATCT<br>CCGCTCTAGA | TCCAACATAA<br>AGGTTGTATT |
| 9951  | GGCGATGATA<br>CCGCTACTAT | TCCGTAGATG<br>AGGCATCTAC | TACCTGGACA<br>ATGGACCTGT   | TCCAGGTGAT<br>AGGTCCACTA | GCCGGCGGGC<br>CGGCCGCCGC |
| 10001 | GTGGTGGAGG<br>CACCACCTCC | CGCGCGGAAA<br>GCGCGCCTTT | GTCGCGGACG<br>CAGCGCCTGC   | CGGTTCCAGA<br>GCCAAGGTCT | TGTTGCGCAG<br>ACAACGCGTC |
| 10051 | CGGCAAAAAG<br>GCCGTTTTTC | TGCTCCATGG<br>ACGAGGTACC | TCGGGACGCT<br>AGCCCTGCGA   | CTGGCCGGTC<br>GACCGGCCAG | AGGCGCGCGC<br>TCCGCGCGCG |
| 10101 | AATCGTTGAC<br>TTAGCAACTG | GCTCTAGACC<br>CGAGATCTGG | GTGCAAAAGG<br>CACGTTTTCC   | AGAGCCTGTA<br>TCTCGGACAT | AGCGGGCACT<br>TCGCCCGTGA |
| 10151 | CTTCCGTGGT<br>GAAGGCACCA | CTGGTGGATA<br>GACCACCTAT | AATTCGCAAG<br>TTAAGCGTTC   | GGTATCATGG<br>CCATAGTACC | CGGACGACCG<br>GCCTGCTGGC |
| 10201 | GGGTTCGAGC<br>CCCAAGCTCG | CCCGTATCCG<br>GGGCATAGGC | GCCGTCCGCC<br>CGGCAGGCGG   | GTGATCCATG<br>CACTAGGTAC | CGGTTACCGC<br>GCCAATGGCG |
| 10251 | CCGCGTGTCG<br>GGCGCACAGC | AACCCAGGTG<br>TTGGGTCCAC | TGCGACGTCA<br>ACGCTGCAGT   | GACAACGGGG<br>CTGTTGCCCC | GAGTGCTCCT<br>CTCACGAGGA |
| 10301 | TTTGGCTTCC<br>AAACCGAAGG | TTCCAGGCGC<br>AAGGTCCGCG | GGCGGCTGCT<br>CCGCCGACGA   | GCGCTAGCTT<br>CGCGATCGAA | TTTTGGCCAC<br>AAAACCGGTG |
| 10351 | TGGCCGCGCG<br>ACCGGCGCGC | CAGCGTAAGC<br>GTCGCATTCG | GGTTAGGCTG<br>CCAATCCGAC   | GAAAGCGAAA<br>CTTTCGCTTT | GCATTAAGTG<br>CGTAATTCAC |

FIG. 10A-13



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|       |                          |                          |                          |                           |                           |
|-------|--------------------------|--------------------------|--------------------------|---------------------------|---------------------------|
| 10401 | GCTCGCTCCC<br>CGAGCGAGGG | TGTAGCCGGA<br>ACATCGGCCT | GGGTTATTTT<br>CCCAATAAAA | CCAAGGGTTG<br>GGTTCCCAAC  | AGTCGCGGGA<br>TCAGCGCCCT  |
| 10451 | CCCCCGGTTT<br>GGGGGCCAAG | GAGTCTCGGA<br>CTCAGAGCCT | CCGGCCGGAC<br>GGCCGGCCTG | TGCGGC GAAC<br>ACGCCGCTTG | GGGGGTTTGC<br>CCCCCAAACG  |
| 10501 | CTCCCCGTCA<br>GAGGGGCAGT | TGCAAGACCC<br>ACGTTCTGGG | CGCTTGCAAA<br>GCGAACGTTT | TTCCTCCGGA<br>AAGGAGGCCT  | AACAGGGACG<br>TTGTCCCTGC  |
| 10551 | AGCCCCTTTT<br>TCGGGGAAAA | TTGCTTTTCC<br>AACGAAAAGG | CAGATGCATC<br>GTCTACGTAG | CGGTGCTGCG<br>GCCACGACGC  | GCAGATGCGC<br>CGTCTACGCG  |
| 10601 | CCCCCTCCTC<br>GGGGGAGGAG | AGCAGCGGCA<br>TCGTCGCCGT | AGAGCAAGAG<br>TCTCGTTCTC | CAGCGGCAGA<br>GTCGCCGTCT  | CATGCAGGGC<br>GTACGTCCCG  |
| 10651 | ACCCTCCCCT<br>TGGGAGGGGA | CCTCCTACCG<br>GGAGGATGGC | CGTCAGGAGG<br>GCAGTCCTCC | GGCGACATCC<br>CCGCTGTAGG  | GCGGTTGACG<br>CGCCA ACTGC |
| 10701 | CGGCAGCAGA<br>GCCGTCGTCT | TGGTGATTAC<br>ACCACTAATG | GAACCCCGC<br>CTTGGGGGCG  | GGCGCCGGGC<br>CCGCGGCCCG  | CCGGCACTAC<br>GGCCGTGATG  |
| 10751 | CTGGACTTGG<br>GACCTGAACC | AGGAGGGCGA<br>TCCTCCCGCT | GGGCCTGGCG<br>CCCGGACCGC | CGGCTAGGAG<br>GCCGATCCTC  | CGCCCTCTCC<br>GCGGGAGAGG  |
| 10801 | TGAGCGGCAC<br>ACTCGCCGTG | CCAAGGGTGC<br>GGTTCCACG  | AGCTGAAGCG<br>TCGACTTCGC | TGATACGCGT<br>ACTATGCGCA  | GAGGCGTACG<br>CTCCGCATGC  |
| 10851 | TGCCGCGGCA<br>ACGGCGCCGT | GAACCTGTTT<br>CTTGGACAAA | CGCGACCGCG<br>GCGCTGGCGC | AGGGAGAGGA<br>TCCCTCTCCT  | GCCCGAGGAG<br>CGGGCTCCTC  |
| 10901 | ATGCGGGATC<br>TACGCCCTAG | GAAAGTTCCA<br>CTTTCAAGGT | CGCAGGGCGC<br>GCGTCCCGCG | GAGCTGCGGC<br>CTCGACGCCG  | ATGGCCTGAA<br>TACCGGACTT  |
| 10951 | TCGCGAGCGG<br>AGCGCTCGCC | TTGCTGCGCG<br>AACGACGCGC | AGGAGGACTT<br>TCCTCCTGAA | TGAGCCCGAC<br>ACTCGGGCTG  | GCGCGAACCG<br>CGCGCTTGGC  |
| 11001 | GGATTAGTCC<br>CCTAATCAGG | CGCGCGCGCA<br>GCGCGCGCGT | CACGTGGCGG<br>GTGCACCGCC | CCGCCGACCT<br>GGCGGCTGGA  | GGTAACCGCA<br>CCATTGGCGT  |
| 11051 | TACGAGCAGA<br>ATGCTCGTCT | CGGTGAACCA<br>GCCACTTGGT | GGAGATTAAC<br>CCTCTAATTG | TTTCAAAAAA<br>AAAGTTTTTT  | GCTTTAACAA<br>CGAAATTGTT  |
| 11101 | CCACGTGCGT<br>GGTGCACGCA | ACGCTTGTGG<br>TGCGAACACC | CGCGCGAGGA<br>GCGCGCTCCT | GGTGGCTATA<br>CCACCGATAT  | GGACTGATGC<br>CCTGACTACG  |
| 11151 | ATCTGTGGGA<br>TAGACACCCT | CTTTGTAAGC<br>GAAACATTCG | GCGCTGGAGC<br>CGCGACCTCG | AAAACCCAAA<br>TTTTGGGTTT  | TAGCAAGCCG<br>ATCGTTCGGC  |

FIG. 10A-14



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11201 CTCATGGCGC AGCTGTTCCCT TATAGTGCAG CACAGCAGGG ACAACGAGGC  
 GAGTACCGCG TCGACAAGGA ATATCACGTC GTGTCGTCCC TGTGCTCCG

11251 ATTCAGGGAT GCGCTGCTAA ACATAGTAGA GCCCGAGGGC CGCTGGCTGC  
 TAAGTCCCTA CCGGACGATT TGTATCATCT CGGGCTCCCG GCGACCGACG

11301 TCGATTTGAT AAACATCCTG CAGAGCATAG TGGTGCAGGA GCGCAGCTTG  
 AGCTAAACTA TTTGTAGGAC GTCTCGTATC ACCACGTCCT CCGGTCGAAC

11351 AGCCTGGCTG ACAAGGTGGC CGCCATCAAC TATTCCATGC TTAGCCTGGG  
 TCGGACCGAC TGTTCACCG GCGGTAGTTG ATAAGGTACG AATCGGACCC

11401 CAAGTTTAC GCCCGCAAGA TATACCATAC CCCTTACGTT CCCATAGACA  
 GTTCAAAATG CGGGCGTTCT ATATGGTATG GGAATGCAA GGGTATCTGT

11451 AGGAGGTAAA GATCGAGGGG TTCTACATGC GCATGGCGCT GAAGGTGCTT  
 TCCTCCATTT CTAGCTCCCC AAGATGTACG CGTACCGCGA CTTCACGAA

11501 ACCTTGAGCG ACGACCTGGG CGTTTATCGC AACGAGCGCA TCCACAAGGC  
 TGGAACTCGC TGCTGGACCC GCAAATAGCG TTGCTCGCGT AGGTGTTCCG

11551 CGTGAGCGTG AGCCGGCGGC GCGAGCTCAG CGACCGCGAG CTGATGCACA  
 GCACTCGCAC TCGGCCGCCG CGCTCGAGTC GCTGGCGCTC GACTACGTGT

11601 GCCTGCAAAG GGCCCTGGCT GGCACGGGCA GCGGCGATAG AGAGGCCGAG  
 CGGACGTTTC CCGGGACCGA CCGTGCCCGT CGCCGCTATC TCTCCGGCTC

11651 TCCTACTTTG ACGCGGGCGC TGACCTGCGC TGGGCCCAA GCCGACGCGC  
 AGGATGAAAC TGCGCCCGCG ACTGGACGCG ACCCGGGGTT CGGCTGCGCG

11701 CCTGGAGGCA GCTGGGGCCG GACCTGGGCT GCGGGTGGCA CCCGCGCGCG  
 GGACCTCCGT CGACCCCGGC CTGGACCCGA CCGCCACCGT GGGCGCGCGC

11751 CTGGCAACGT CGGCGGCGTG GAGGAATATG ACGAGGACGA TGAGTACGAG  
 GACCGTTGCA GCCGCCGCAC CTCCTTATAC TGCTCCTGCT ACTCATGCTC

11801 CCAGAGGACG GCGAGTACTA AGCGGTGATG TTTCTGATCA GATGATGCAA  
 GGTCTCCTGC CGCTCATGAT TCGCCACTAC AAAGACTAGT CTACTACGTT

11851 GACGCAACGG ACCCGGCGGT GCGGGCGGCG CTGCAGAGCC AGCCGTCCGG  
 CTGCGTTGCC TGGGCCGCCA CGCCCGCCGC GACGTCTCGG TCGGCAGGCC

11901 CCTTAACTCC ACGGACGACT GCGGCCAGGT CATGGACCGC ATCATGTCGC  
 GGAATTGAGG TGCCTGCTGA CCGCGGTCCA GTACCTGGCG TAGTACAGCG

11951 TGACTGCGCG CAATCCTGAC GCGTTCCGGC AGCAGCCGCA GGCCAACCGG  
 ACTGACGCGC GTTAGGACTG CGCAAGGCCG TCGTCGGCGT CCGGTTGGCC

FIG. 10A-15



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|       |                          |                           |                          |                          |                          |
|-------|--------------------------|---------------------------|--------------------------|--------------------------|--------------------------|
| 12001 | CTCTCCGCAA<br>GAGAGGCGTT | TTCTGGAAGC<br>AAGACCTTCG  | GGTGGTCCCG<br>CCACCAGGGC | GCGCGCGCAA<br>CGCGCGCGTT | ACCCACGCA<br>TGGGGTGCGT  |
| 12051 | CGAGAAGGTG<br>GCTCTTCCAC | CTGGCGATCG<br>GACCGCTAGC  | TAAACGCGCT<br>ATTTGCGCGA | GGCCGAAAAC<br>CCGGCTTTTG | AGGGCCATCC<br>TCCCGGTAGG |
| 12101 | GGCCCGACGA<br>CCGGGCTGCT | GGCCGGCCTG<br>CCGGCCGGAC  | GTCTACGACG<br>CAGATGCTGC | CGCTGCTTCA<br>GCGACGAAGT | GCGCGTGGCT<br>CGCGCACCGA |
| 12151 | CGTTACAACA<br>GCAATGTTGT | GCGGCAACGT<br>CGCCGTTGCA  | GCAGACCAAC<br>CGTCTGGTTG | CTGGACCGGC<br>GACCTGGCCG | TGGTGGGGGA<br>ACCACCCCT  |
| 12201 | TGTGCGCGAG<br>ACACGCGCTC | GCCGTGGCGC<br>CGGCACCGCG  | AGCGTGAGCG<br>TCGCACTCGC | CGCGCAGCAG<br>GCGCGTCGTC | CAGGGCAACC<br>GTCCCGTTGG |
| 12251 | TGGGCTCCAT<br>ACCCGAGGTA | GGTTGCACTA<br>CCAACGTGAT  | AACGCCTTCC<br>TTGCGGAAGG | TGAGTACACA<br>ACTCATGTGT | GCCCGCCAAC<br>CGGGCGGTTG |
| 12301 | GTGCCGCGGG<br>CACGGCGCCC | GACAGGAGGA<br>CTGTCCTCCT  | CTACACCAAC<br>GATGTGGTTG | TTTGTGAGCG<br>AAACACTCGC | CACTGCGGCT<br>GTGACGCCGA |
| 12351 | AATGGTGACT<br>TTACCACTGA | GAGACACCGC<br>CTCTGTGGCG  | AAAGTGAGGT<br>TTTCACTCCA | GTACCAGTCT<br>CATGGTCAGA | GGGCCAGACT<br>CCCGGTCTGA |
| 12401 | ATTTTTTCCA<br>TAAAAAAGGT | GACCAGTAGA<br>CTGGTCATCT  | CAAGGCCTGC<br>GTTCCGGACG | AGACCGTAAA<br>TCTGGCATT  | CCTGAGCCAG<br>GGACTCGGTC |
| 12451 | GCTTTCAAAA<br>CGAAAGTTTT | ACTTGCAGGG<br>TGAACGTCCC  | GCTGTGGGGG<br>CGACACCCCC | GTGCGGGCTC<br>CACGCCCGAG | CCACAGGCGA<br>GGTGTCCGCT |
| 12501 | CCGCGCGACC<br>GGCGCGCTGG | GTGTCTAGCT<br>CACAGATCGA  | TGCTGACGCC<br>ACGACTGCGG | CAACTCGCGC<br>GTTGAGCGCG | CTGTTGCTGC<br>GACAACGACG |
| 12551 | TGCTAATAGC<br>ACGATTATCG | GCCCTTCACG<br>CGGGAAGTGC  | GACAGTGGCA<br>CTGTCACCGT | GCGTGTCCCG<br>CGCACAGGGC | GGACACATAC<br>CCTGTGTATG |
| 12601 | CTAGGTCACT<br>GATCCAGTGA | TGCTGACACT<br>ACGACTGTGA  | GTACCGCGAG<br>CATGGCGCTC | GCCATAGGTC<br>CGGTATCCAG | AGGCGCATGT<br>TCCGCGTACA |
| 12651 | GGACGAGCAT<br>CCTGCTCGTA | ACTTTCCAGG<br>TGAAAGGTCC  | AGATTACAAG<br>TCTAATGTTC | TGTCAGCCGC<br>ACAGTCGGCG | GCGCTGGGGC<br>CGCGACCCCG |
| 12701 | AGGAGGACAC<br>TCCTCCTGTG | GGGCAGCCTG<br>CCCGTCGGAC  | GAGGCAACCC<br>CTCCGTTGGG | TAAACTACCT<br>ATTTGATGGA | GCTGACCAAC<br>CGACTGGTTG |
| 12751 | CGGCGGCAGA<br>GCCGCCGTCT | AGATCCCCCTC<br>TCTAGGGGAG | GTTGCACAGT<br>CAACGTGTCA | TTAAACAGCG<br>AATTTGTGCG | AGGAGGAGCG<br>TCCTCCTCGC |

FIG. 10A-16



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12801 CATTTTGCGC TACGTGCAGC AGAGCGTGAG CCTTAACCTG ATGCGCGACG  
 GTAAAACGCG ATGCACGTCG TCTCGCACTC GGAATTGGAC TACGCGCTGC

12851 GGGTAACGCC CAGCGTGGCG CTGGACATGA CCGCGCGCAA CATGGAACCG  
 CCCATTGCGG GTCGCACCGC GACCTGTACT GCGCGCGCCTT GTACCTTGGC

12901 GGCATGTATG CCTCAAACCG GCCGTTTATC AACCGCCTAA TGGACTACTT  
 CCGTACATAC GGAGTTTGGC CGGCAAATAG TTGGCGGATT ACCTGATGAA

12951 GCATCGCGCG GCCGCCGTGA ACCCCGAGTA TTTCACCAAT GCCATCTTGA  
 CGTAGCGCGC CGGCGGGCACT TGGGGCTCAT AAAGTGGTTA CGGTAGAACT

13001 ACCCGCACTG GCTACCGCCC CCTGGTTTCT ACACCGGGGG ATTCGAGGTG  
 TGGGCGTGAC CGATGGCGGG GGACCAAAGA TGTGGCCCCC TAAGCTCCAC

13051 CCCGAGGGTA ACGATGGATT CCTCTGGGAC GACATAGACG ACAGCGTGTT  
 GGGCTCCCAT TGCTACCTAA GGAGACCCTG CTGTATCTGC TGTCGCACAA

13101 TTCCCCGCAA CCGCAGACCC TGCTAGAGTT GCAACAGCGC GAGCAGGCAG  
 AAGGGGCGTT GCGTCTGGG ACGATCTCAA CGTTGTCGCG CTCGTCCGTC

13151 AGGCGGCGCT GCGAAAGGAA AGCTTCCGCA GGCCAAGCAG CTTGTCCGAT  
 TCCGCCGCGA CGCTTTCCTT TCGAAGGCGT CCGGTTCGTC GAACAGGCTA

13201 CTAGGCGCTG CGGCCCCGCG GTCAGATGCT AGTAGCCCAT TTCCAAGCTT  
 GATCCGCGAC GCCGGGGCGC CAGTCTACGA TCATCGGGTA AAGGTTGAA

13251 GATAGGGTCT CTTACCAGCA CTCGCACCAC CCGCCCGCGC CTGCTGGGCG  
 CTATCCCAGA GAATGGTCGT GAGCGTGGTG GCGGGGCGCG GACGACCCGC

13301 AGGAGGAGTA CCTAAACAAC TCGCTGCTGC AGCCGCAGCG CGAAAAAAC  
 TCCTCCTCAT GGATTTGTTG AGCGACGACG TCGGCGTCGC GCTTTTTTTG

13351 CTGCCTCCGG CATTTCCCAA CAACGGGATA GAGAGCCTAG TGGACAAGAT  
 GACGGAGGCC GTAAAGGGTT GTTGCCCTAT CTCTCGGATC ACCTGTTCTA

13401 GAGTAGATGG AAGACGTACG CGCAGGAGCA CAGGGACGTG CCAGGCCCGC  
 CTCATCTACC TTCTGCATGC GCGTCCCTCGT GTCCCTGCAC GGTCCGGGCG

13451 GCCCGCCCAC CCGTCGTCAA AGGCACGACC GTCAGCGGGG TCTGGTGTGG  
 CGGGCGGGTG GGCAGCAGTT TCCGTGCTGG CAGTCGCCCC AGACCACACC

13501 GAGGACGATG ACTCGGCAGA CGACAGCAGC GTCCTGGATT TGGGAGGGAG  
 CTCCTGCTAC TGAGCCGTCT GCTGTCGTCG CAGGACCTAA ACCCTCCCTC

13551 TGGCAACCCG TTTGCGCACC TTCGCCCCAG GCTGGGGAGA ATGTTTTTAAA  
 ACCGTTGGGC AAACGCGTGG AAGCGGGGTC CGACCCCTCT TACAAAATTT

FIG. 10A-17



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|       |             |            |            |             |            |
|-------|-------------|------------|------------|-------------|------------|
| 13601 | AAAAAAAAAA  | GCATGATGCA | AAATAAAAAA | CTCACCAAGG  | CCATGGCACC |
|       | TTTTTTTTTT  | CGTACTACGT | TTTATTTTTT | GAGTGGTTCC  | GGTACCGTGG |
| 13651 | GAGCGTTGGT  | TTTCTTGAT  | TCCCCTTAGT | ATGCGGCGCG  | CGCCGATGTA |
|       | CTCGCAACCA  | AAAGAACATA | AGGGGAATCA | TACGCCGCGC  | GCCGCTACAT |
| 13701 | TGAGGAAGGT  | CCTCCTCCCT | CCTACGAGAG | TGTGGTGAGC  | GCGGCGCCAG |
|       | ACTCCTTCCA  | GGAGGAGGGA | GGATGCTCTC | ACACCACTCG  | CGCCGCGGTC |
| 13751 | TGGCGGCGGC  | GCTGGGTTCT | CCCTTCGATG | CTCCCCTGGA  | CCCGCCGTTT |
|       | ACCGCCGCCG  | CGACCCAAGA | GGGAAGCTAC | GAGGGGACCT  | GGGCGGCAAA |
| 13801 | GTGCCTCCGC  | GGTACCTGCG | GCCTACCGGG | GGGAGAAACA  | GCATCCGTTA |
|       | CACGGAGGCG  | CCATGGACGC | CGGATGGCCC | CCCTCTTTGT  | CGTAGGCAAT |
| 13851 | CTCTGAGTTG  | GCACCCCTAT | TCGACACCAC | CCGTGTGTAC  | CTGGTGGACA |
|       | GAGACTCAAC  | CGTGGGGATA | AGCTGTGGTG | GGCACACATG  | GACCACCTGT |
| 13901 | ACAAGTCAAC  | GGATGTGGCA | TCCCTGAACT | ACCAGAACGA  | CCACAGCAAC |
|       | TGTTCAAGTTG | CCTACACCGT | AGGGACTTGA | TGGTCTTGCT  | GGTGTGTTG  |
| 13951 | TTTCTGACCA  | CGGTCATTCA | AAACAATGAC | TACAGCCCGG  | GGGAGGCAAG |
|       | AAAGACTGGT  | GCCAGTAAGT | TTTGTTACTG | ATGTCGGGCC  | CCCTCCGTTT |
| 14001 | CACACAGACC  | ATCAATCTTG | ACGACCGGTC | GCACTGGGGC  | GGCGACCTGA |
|       | GTGTGTCTGG  | TAGTTAGAAC | TGCTGGCCAG | CGTGACCCCG  | CCGCTGGACT |
| 14051 | AAACCATCCT  | GCATACCAAC | ATGCCAAATG | TGAACGAGTT  | CATGTTTACC |
|       | TTTGGTAGGA  | CGTATGGTTG | TACGGTTTAC | ACTTGCTCAA  | GTACAAATGG |
| 14101 | AATAAGTTTA  | AGGCGCGGGT | GATGGTGTCG | CGCTTGCCTA  | CTAAGGACAA |
|       | TTATTCAAAT  | TCCGCGCCCA | CTACCACAGC | GCGAACGGAT  | GATTCCTGTT |
| 14151 | TCAGGTGGAG  | CTGAAATACG | AGTGGGTGGA | G TTCACGCTG | CCCGAGGGCA |
|       | AGTCCACCTC  | GACTTTATGC | TCACCCACCT | CAAGTGCGAC  | GGGCTCCCGT |
| 14201 | ACTACTCCGA  | GACCATGACC | ATAGACCTTA | TGAACAACGC  | GATCGTGGAG |
|       | TGATGAGGCT  | CTGGTACTGG | TATCTGGAAT | ACTTGTTGCG  | CTAGCACCTC |
| 14251 | CACTACTTGA  | AAGTGGGCAG | ACAGAACGGG | GTTCTGGAAA  | GCGACATCGG |
|       | GTGATGAACT  | TTCACCCGTC | TGTCTTGCCC | CAAGACCTTT  | CGCTGTAGCC |
| 14301 | GGTAAAGTTT  | GACACCCGCA | ACTTCAGACT | GGGGTTTGAC  | CCCGTCACTG |
|       | CCATTTCAA   | CTGTGGGCGT | TGAAGTCTGA | CCCCAAACTG  | GGGCAGTGAC |
| 14351 | GTCTTGTCAT  | GCCTGGGGTA | TATACAAACG | AAGCCTTCCA  | TCCAGACATC |
|       | CAGAACAGTA  | CGGACCCCAT | ATATGTTTGC | TTCGGAAGGT  | AGGTCTGTAG |

FIG. 10A-18



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|       |                          |                          |                          |                          |                          |
|-------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|
| 14401 | ATTTTGCTGC<br>TAAAACGACG | CAGGATGCGG<br>GTCCTACGCC | GGTGGACTTC<br>CCACCTGAAG | ACCCACAGCC<br>TGGGTGTCGG | GCCTGAGCAA<br>CGGACTCGTT |
| 14451 | CTTGTTGGGC<br>GAACAACCCG | ATCCGCAAGC<br>TAGGCGTTCC | GGCAACCCTT<br>CCGTTGGGAA | CCAGGAGGGC<br>GGTCCTCCCG | TTTAGGATCA<br>AAATCCTAGT |
| 14501 | CCTACGATGA<br>GGATGCTACT | TCTGGAGGGT<br>AGACCTCCCA | GGTAACATTC<br>CCATTGTAAG | CCGCACTGTT<br>GGCGTGACAA | GGATGTGGAC<br>CCTACACCTG |
| 14551 | GCCTACCAGG<br>CGGATGGTCC | CGAGCTTGAA<br>GCTCGAACTT | AGATGACACC<br>TCTACTGTGG | GAACAGGGCG<br>CTTGTCCCGC | GGGGTGGCGC<br>CCCCACCGCG |
| 14601 | AGGCGGCAGC<br>TCCGCCGTCG | AACAGCAGTG<br>TTGTCGTCAC | GCAGCGGCGC<br>CGTCGCCGCG | GGAAGAGAAC<br>CCTTCTCTTG | TCCAACGCGG<br>AGGTTGCGCC |
| 14651 | CAGCCGCGGC<br>GTCGGCGCCG | AATGCAGCCG<br>TTACGTCGGC | GTGGAGGACA<br>CACCTCCTGT | TGAACGATCA<br>ACTTGCTAGT | TGCCATTCGC<br>ACGGTAAGCG |
| 14701 | GGCGACACCT<br>CCGCTGTGGA | TTGCCACACG<br>AACGGTGTGC | GGCTGAGGAG<br>CCGACTCCTC | AAGCGCGCTG<br>TTCGCGCGAC | AGGCCGAAGC<br>TCCGGCTTCG |
| 14751 | AGCGGCCGAA<br>TCGCCGGCTT | GCTGCCGCCC<br>CGACGGCGGG | CCGCTGCGCA<br>GGCGACGCGT | ACCCGAGGTC<br>TGGGCTCCAG | GAGAAGCCTC<br>CTCTTCGGAG |
| 14801 | AGAAGAAACC<br>TCTTCTTTGG | GGTGATCAAA<br>CCACTAGTTT | CCCCTGACAG<br>GGGGACTGTC | AGGACAGCAA<br>TCCTGTCGTT | GAAACGCAGT<br>CTTTGCGTCA |
| 14851 | TACAACCTAA<br>ATGTTGGATT | TAAGCAATGA<br>ATTCGTTACT | CAGCACCTTC<br>GTCGTGGAAG | ACCCAGTACC<br>TGGGTCATGG | GCAGCTGGTA<br>CGTCGACCAT |
| 14901 | CCTTGCATAC<br>GGAACGTATG | AACTACGGCG<br>TTGATGCCGC | ACCCTCAGAC<br>TGGGAGTCTG | CGGAATCCGC<br>GCCTTAGGCG | TCATGGACCC<br>AGTACCTGGG |
| 14951 | TGCTTTGCAC<br>ACGAAACGTG | TCCTGACGTA<br>AGGACTGCAT | ACCTGCGGCT<br>TGGACGCCGA | CGGAGCAGGT<br>GCCTCGTCCA | CTACTGGTCG<br>GATGACCAGC |
| 15001 | TTGCCAGACA<br>AACGGTCTGT | TGATGCAAGA<br>ACTACGTTCT | CCCCGTGACC<br>GGGGCACTGG | TTCCGCTCCA<br>AAGGCGAGGT | CGCGCCAGAT<br>GCGCGGTCTA |
| 15051 | CAGCAACTTT<br>GTCGTTGAAA | CCGGTGGTGG<br>GGCCACCACC | GCGCCGAGCT<br>CGCGGCTCGA | GTTGCCCGTG<br>CAACGGGCAC | CACTCCAAGA<br>GTGAGGTTCT |
| 15101 | GCTTCTACAA<br>CGAAGATGTT | CGACCAGGCC<br>GCTGGTCCGG | GTCTACTCCC<br>CAGATGAGGG | AACTCATCCG<br>TTGAGTAGGC | CCAGTTTACC<br>GGTCAAATGG |
| 15151 | TCTCTGACCC<br>AGAGACTGGG | ACGTGTTCAA<br>TGCACAAGTT | TCGCTTTCCC<br>AGCGAAAGGG | GAGAACCAGA<br>CTCTTGGTCT | TTTTGGCGCG<br>AAAACCGCGC |

FIG. 10A-19



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15201 CCCGCCAGCC CCCACCATCA CCACCGTCAG TGAAAACGTT CCTGCTCTCA  
 GGGCGGTCGG GGGTGGTAGT GGTGGCAGTC ACTTTTGCAA GGACGAGAGT

15251 CAGATCACGG GACGCTACCG CTGCGCAACA GCATCGGAGG AGTCCAGCGA  
 GTCTAGTGCC CTGCGATGGC GACGCGTTGT CGTAGCCTCC TCAGGTGGCT

15301 GTGACCATTA CTGACGCCAG ACGCCGCACC TGCCCCCTACG TTTACAAGGC  
 CACTGGTAAT GACTGCGGTC TGCGGCGTGG ACGGGGATGC AAATGTTCCG

15351 CCTGGGCATA GTCTCGCCGC GCGTCCTATC GAGCCGCACT TTTTGAGCAA  
 GGACCCGTAT CAGAGCGGCG CGCAGGATAG CTCGGCGTGA AAAACTCGTT

15401 GCATGTCCAT CCTTATATCG CCCAGCAATA ACACAGGCTG GGGCCTGCGC  
 CGTACAGGTA GGAATATAGC GGGTCGTTAT TGTGTCCGAC CCCGGACGCG

15451 TTCCAAGCA AGATGTTTGG CGGGGCAAG AAGCGCTCCG ACCAACACCC  
 AAGGGTTCGT TCTACAAACC GCCCCGGTTC TTCGCGAGGC TGGTTGTGGG

15501 AGTGCGCGTG CGCGGGCACT ACCGCGCGCC CTGGGGCGCG CACAAACGCG  
 TCACGCGCAC GCGCCCGTGA TGGCGCGCGG GACCCCGCGC GTGTTTGCGC

15551 GCCGCACTGG GCGCACCACC GTCGATGACG CCATCGACGC GGTGGTGGAG  
 CGGCGTGACC CGCGTGGTGG CAGCTACTGC GGTAGCTGCG CCACCACCTC

15601 GAGGCGCGCA ACTACACGCC CACGCCGCCA CCAGTGTCCA CAGTGGACGC  
 CTCCGCGCGT TGATGTGCGG GTGCGGGCGGT GGTCACAGGT GTCACCTGCG

15651 GGCCATTCAG ACCGTGGTGC GCGGAGCCCG GCGCTATGCT AAAATGAAGA  
 CCGGTAAGTC TGGCACCACG CGCCTCGGGC CGCGATACGA TTTTACTTCT

15701 GACGGCGGAG GCGCGTAGCA CGTCGCCACC GCCGCCGACC CGGCACTGCC  
 CTGCCGCCTC CGCGCATCGT GCAGCGGTGG CGGCGGCTGG GCCGTGACGG

15751 GCCCAACGCG CGGCGGGCGG CCTGCTTAAC CGCGCACGTC GCACCGGCCG  
 CGGGTTGCGC GCCGCCGCCG GGACGAATTG GCGCGTGACG CGTGGCCGGC

15801 ACGGGCGGCC ATGCGGGCCG CTCGAAGGCT GGCCGCGGGT ATTGTCACTG  
 TGCCCGCCGG TACGCCCGGC GAGCTTCCGA CCGGCGCCA TAACAGTGAC

15851 TGCCCCCAG GTCCAGGCGA CGAGCGGCCG CCGCAGCAGC CGCGGCCATT  
 ACGGGGGGTC CAGGTCCGCT GCTCGCCGGC GCGTTCGTCG GCGCCGGTAA

15901 AGTGCTATGA CTCAGGGTCG CAGGGGCAAC GTGTATTGGG TCGCGACTC  
 TCACGATACT GAGTCCAGC GTCCCCGTTG CACATAACCC ACGCGCTGAG

15951 GGTTAGCGGC CTGCGCGTGC CCGTGCGCAC CCGCCCCCG CGCAACTAGA  
 CCAATCGCCG GACGCGCACG GGCACGCGTG GCGGGGGGGC GCGTTGATCT

FIG. 10A-20



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|       |            |            |            |            |            |
|-------|------------|------------|------------|------------|------------|
| 16001 | TTGCAAGAAA | AAACTACTTA | GACTCGTACT | GTTGTATGTA | TCCAGCGGCG |
|       | AACGTTCTTT | TTTGATGAAT | CTGAGCATGA | CAACATACAT | AGGTCGCCGC |
| 16051 | GCGGCGCGCA | ACGAAGCTAT | GTCCAAGCGC | AAAATCAAAG | AAGAGATGCT |
|       | CGCCGCGCGT | TGCTTCGATA | CAGGTTCCGC | TTTTAGTTTC | TTCTCTACGA |
| 16101 | CCAGGTCATC | GCGCCGGAGA | TCTATGGCCC | CCCGAAGAAG | GAAGAGCAGG |
|       | GGTCCAGTAG | CGCGGCCTCT | AGATACCGGG | GGGCTTCTTC | CTTCTCGTCC |
| 16151 | ATTACAAGCC | CCGAAAGCTA | AAGCGGGTCA | AAAAGAAAAA | GAAAGATGAT |
|       | TAATGTTCGG | GGCTTTCGAT | TTCGCCCAGT | TTTTCTTTTT | CTTCTACTA  |
| 16201 | GATGATGAAC | TTGACGACGA | GGTGGAACTG | CTGCACGCTA | CCGCGCCCAG |
|       | CTACTACTTG | AACTGCTGCT | CCACCTTGAC | GACGTGCGAT | GGCGCGGGTC |
| 16251 | GCGACGGGTA | CAGTGGAAG  | GTCGACGCGT | AAAACGTGTT | TTGCGACCCG |
|       | CGCTGCCCAT | GTCACCTTTC | CAGCTGCGCA | TTTTGCACAA | AACGCTGGGC |
| 16301 | GCACCACCGT | AGTCTTTACG | CCCGGTGAGC | GCTCCACCCG | CACCTACAAG |
|       | CGTGGTGGCA | TCAGAAATGC | GGGCCACTCG | CGAGGTGGGC | GTGGATGTTC |
| 16351 | CGCGTGTATG | ATGAGGTGTA | CGGCGACGAG | GACCTGCTTG | AGCAGGCCAA |
|       | GCGCACATAC | TACTCCACAT | GCCGCTGCTC | CTGGACGAAC | TCGTCCGGTT |
| 16401 | CGAGCGCCTC | GGGGAGTTTG | CCTACGGAAA | GCGGCATAAG | GACATGCTGG |
|       | GCTCGCGGAG | CCCCTCAAAC | GGATGCCTTT | CGCCGTATTC | CTGTACGACC |
| 16451 | CGTTGCCGCT | GGACGAGGGC | AACCCAACAC | CTAGCCTAAA | GCCCGTAACA |
|       | GCAACGGCGA | CCTGCTCCCG | TTGGGTTGTG | GATCGGATTT | CGGGCATTGT |
| 16501 | CTGCAGCAGG | TGCTGCCCGC | GCTTGCACCG | TCCGAAGAAA | AGCGCGGCCT |
|       | GACGTGCTCC | ACGACGGGCG | CGAACGTGGC | AGGCTTCTTT | TCGCGCCGGA |
| 16551 | AAAGCGCGAG | TCTGGTGACT | TGGCACCCAC | CGTGCAGCTG | ATGGTACCCA |
|       | TTTCGCGCTC | AGACCACTGA | ACCGTGGGTG | GCACGTCGAC | TACCATGGGT |
| 16601 | AGCGCCAGCG | ACTGGAAGAT | GTCTTGGAAA | AAATGACCGT | GGAACCTGGG |
|       | TCGCGGTTCG | TGACCTTCTA | CAGAACCTTT | TTTACTGGCA | CCTTGGACCC |
| 16651 | CTGGAGCCCG | AGGTCCGCGT | GCGGCCAATC | AAGCAGGTGG | CGCCGGGACT |
|       | GACCTCGGGC | TCCAGGCGCA | CGCCGGTTAG | TTGTTCCACC | GCGGCCCTGA |
| 16701 | GGGCGTGCAG | ACCGTGGACG | TTCAGATACC | CACTACCAGT | AGCACCAGTA |
|       | CCCGCACGTC | TGGCACCTGC | AAGTCTATGG | GTGATGGTCA | TCGTGGTCAT |
| 16751 | TTGCCACCGC | CACAGAGGGC | ATGGAGACAC | AAACGTCCCC | GGTTGCCTCA |
|       | AACGGTGGCG | GTGTCTCCCG | TACCTCTGTG | TTTGCAGGGG | CCAACGGAGT |

FIG. 10A-21



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|       |                          |                           |                          |                          |                           |
|-------|--------------------------|---------------------------|--------------------------|--------------------------|---------------------------|
| 16801 | GCGGTGGCGG<br>CGCCACCGCC | ATGCCGCGGT<br>TACGGCGCCA  | GCAGGCGGTC<br>CGTCCGCCAG | GCTGCGGCCG<br>CGACGCCGGC | CGTCCAAGAC<br>GCAGGTTCTG  |
| 16851 | CTCTACGGAG<br>GAGATGCCTC | GTGCAAACGG<br>CACGTTTGCC  | ACCCGTGGAT<br>TGGGCACCTA | GTTTCGCGTT<br>CAAAGCGCAA | TCAGCCCCCC<br>AGTCGGGGGG  |
| 16901 | GGCGCCCGCG<br>CCGCGGGCGC | CCGTTGAGG<br>GGCAAGCTCC   | AAGTACGGCG<br>TTCATGCCGC | CCGCCAGCGC<br>GGCGGTGCGC | GCTACTGCCC<br>CGATGACGGG  |
| 16951 | GAATATGCCC<br>CTTATACGGG | TACATCCTTC<br>ATGTAGGAAG  | CATTGCGCCT<br>GTAACGCGGA | ACCCCCGGCT<br>TGGGGGCCGA | ATCGTGGCTA<br>TAGCACCGAT  |
| 17001 | CACCTACCGC<br>GTGGATGGCG | CCCAGAAGAC<br>GGGTCTTCTG  | GAGCAACTAC<br>CTCGTTGATG | CCGACGCCGA<br>GGCTGCGGCT | ACCACCACTG<br>TGGTGGTGAC  |
| 17051 | GAACCCGCCG<br>CTTGGGCGGC | CCGCCGTCGC<br>GGCGGCAGCG  | CGTCGCCAGC<br>GCAGCGGTCG | CCGTGCTGGC<br>GGCACGACCG | CCCGATTTC<br>GGGCTAAAGG   |
| 17101 | GTGCGCAGGG<br>CACGCGTCCC | TGGCTCGCGA<br>ACCGAGCGCT  | AGGAGGCAGG<br>TCCTCCGTCC | ACCCTGGTGC<br>TGGGACCACG | TGCCAACAGC<br>ACGGTTGTG   |
| 17151 | GCGCTACCAC<br>CGCGATGGTG | CCCAGCATCG<br>GGGTCGTAGC  | TTTAAAAGCC<br>AAATTTTCGG | GGTCTTTGTG<br>CCAGAAACAC | GTTCTTGCAG<br>CAAGAACGTC  |
| 17201 | ATATGGCCCT<br>TATACCGGGA | CACCTGCCGC<br>GTGGACGGCG  | CTCCGTTTCC<br>GAGGCAAAGG | CGGTGCCGGG<br>GCCACGGCCC | ATTCGAGGA<br>TAAGGCTCCT   |
| 17251 | AGAATGCACC<br>TCTTACGTGG | GTAGGAGGGG<br>CATCCTCCCC  | CATGGCCGGC<br>GTACCGGCCG | CACGGCCTGA<br>GTGCCGGACT | CGGGCGGCAT<br>GCCCCCGTA   |
| 17301 | GCGTCGTGCG<br>CGCAGCACGC | CACCACCGGC<br>GTGGTGGCCG  | GGCGGCGCGC<br>CCGCCGCGCG | GTCGCACCGT<br>CAGCGTGGCA | CGCATGCGCG<br>GCGTACGCGC  |
| 17351 | GCGGTATCCT<br>CGCCATAGGA | GCCCCTCCTT<br>CGGGGAGGAA  | ATTCCACTGA<br>TAAGGTGACT | TCGCCGCGGC<br>AGCGGCGCCG | GATTGGCGCC<br>CTAACC GCGG |
| 17401 | GTGCCCGGAA<br>CACGGGCCTT | TTGCATCCGT<br>AACGTAGGCA  | GGCCTTGCAG<br>CCGGAACGTC | GCGCAGAGAC<br>CGCGTCTCTG | ACTGATTAAA<br>TGACTAATTT  |
| 17451 | AACAAGTTGC<br>TTGTTCAACG | ATGTGGAAA<br>TACACCTTTT   | ATCAAATAA<br>TAGTTTTATT  | AAAGTCTGGA<br>TTTCAGACCT | CTCTCACGCT<br>GAGAGTGC GA |
| 17501 | CGCTTGGTCC<br>GCGAACCAGG | TGTA ACTATT<br>ACATTGATAA | TTGTAGAATC<br>AACATCTTAC | GAAGACATCA<br>CTTCTGTAGT | ACTTTGCGTC<br>TGAAACGCAG  |
| 17551 | TCTGGCCCCG<br>AGACCGGGGC | CGACACGGCT<br>GCTGTGCCGA  | CGCGCCCGTT<br>GCGCGGGCAA | CATGGGAAAC<br>GTACCCTTTG | TGGCAAGATA<br>ACCGTTCTAT  |

FIG. 10A-22



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|       |                          |                           |                          |                          |                           |
|-------|--------------------------|---------------------------|--------------------------|--------------------------|---------------------------|
| 17601 | TCGGCACCAG<br>AGCCGTGGTC | CAATATGAGC<br>GTTATACTCG  | GGTGGCGCCT<br>CCACCGCGGA | TCAGCTGGGG<br>AGTCGACCCC | CTCGCTGTGG<br>GAGCGACACC  |
| 17651 | AGCGGCATTA<br>TCGCCGTAAT | AAAATTTTCGG<br>TTTTAAAGCC | TTCCACCGTT<br>AAGGTGGCAA | AAGAACTATG<br>TTCTTGATAC | GCAGCAAGGC<br>CGTCGTTCCG  |
| 17701 | CTGGAACAGC<br>GACCTTGTCG | AGCACAGGCC<br>TCGTGTCCGG  | AGATGCTGAG<br>TCTACGACTC | GGATAAGTTG<br>CCTATTCAAC | AAAGAGCAAA<br>TTTCTCGTTT  |
| 17751 | ATTTCCAACA<br>TAAAGGTTGT | AAAGGTGGTA<br>TTTCCACCAT  | GATGGCCTGG<br>CTACCGGACC | CCTCTGGCAT<br>GGAGACCGTA | TAGCGGGGTG<br>ATCGCCCCAC  |
| 17801 | GTGGACCTGG<br>CACCTGGACC | CCAACCAGGC<br>GGTTGGTCCG  | AGTGCAAAAT<br>TCACGTTTTA | AAGATTAACA<br>TTCTAATTGT | GTAAGCTTGA<br>CATTCGAACT  |
| 17851 | TCCCCGCCCT<br>AGGGGCGGGA | CCCGTAGAGG<br>GGGCATCTCC  | AGCCTCCACC<br>TCGGAGGTGG | GGCCGTGGAG<br>CCGGCACCTC | ACAGTGTCTC<br>TGTCACAGAG  |
| 17901 | CAGAGGGGCG<br>GTCTCCCCGC | TGGCGAAAAG<br>ACCGCTTTTC  | CGTCCGCGCC<br>GCAGGCGCGG | CCGACAGGGA<br>GGCTGTCCCT | AGAAACTCTG<br>TCTTTGAGAC  |
| 17951 | GTGACGCAAA<br>CACTGCGTTT | TAGACGAGCC<br>ATCTGCTCGG  | TCCCTCGTAC<br>AGGGAGCATG | GAGGAGGCAC<br>CTCCTCCGTG | TAAAGCAAGG<br>ATTTGTTTCC  |
| 18001 | CCTGCCACC<br>GGACGGGTGG  | ACCCGTCCA<br>TGGGCAGGGT   | TCGCGCCCAT<br>AGCGCGGGTA | GGCTACCGGA<br>CCGATGGCCT | GTGCTGGGCC<br>CACGACCCGG  |
| 18051 | AGCACACACC<br>TCGTGTGTGG | CGTAACGCTG<br>GCATTGCGAC  | GACCTGCCTC<br>CTGGACGGAG | CCCCGCCGA<br>GGGGGCGGCT  | CACCCAGCAG<br>GTGGGTCTGTC |
| 18101 | AAACCTGTGC<br>TTTGGACACG | TGCCAGGCC<br>ACGGTCCGGG   | GACCGCCGTT<br>CTGGCGGCAA | GTTGTAACCC<br>CAACATTGGG | GTCCTAGCCG<br>CAGGATCGGC  |
| 18151 | CGCGTCCCTG<br>GCGCAGGGAC | CGCCGCGCCG<br>GCGGCGCGGC  | CCAGCGGTCC<br>GGTCGCCAGG | GCGATCGTTG<br>CGCTAGCAAC | CGGCCCGTAG<br>GCCGGGCATC  |
| 18201 | CCAGTGGCAA<br>GGTCACCGTT | CTGGCAAAGC<br>GACCGTTTCG  | ACACTGAACA<br>TGTGACTTGT | GCATCGTGGG<br>CGTAGCACC  | TCTGGGGGTG<br>AGACCCCCAC  |
| 18251 | CAATCCCTGA<br>GTTAGGGACT | AGCGCCGACG<br>TCGCGGCTGC  | ATGCTTCTGA<br>TACGAAGACT | TAGCTAACGT<br>ATCGATTGCA | GTCGTATGTG<br>CAGCATAAC   |
| 18301 | TGTCATGTAT<br>ACAGTACATA | GCGTCCATGT<br>CGCAGGTACA  | CGCCGCCAGA<br>GCGGCGGTCT | GGAGCTGCTG<br>CCTCGACGAC | AGCCGCCGCG<br>TCGGCGGCGC  |
| 18351 | CGCCCGCTTT<br>GCGGGCGAAA | CCAAGATGGC<br>GGTTCTACCG  | TACCCCTTCG<br>ATGGGGAAGC | ATGATGCCGC<br>TACTACGGCG | AGTGGTCTTA<br>TCACCAGAAT  |

FIG. 10A-23



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|       |                           |                           |                          |                          |                          |
|-------|---------------------------|---------------------------|--------------------------|--------------------------|--------------------------|
| 18401 | CATGCACATC<br>GTACGTGTAG  | TCGGGCCAGG<br>AGCCCGGTCC  | ACGCCTCGGA<br>TGCGGAGCCT | GTACCTGAGC<br>CATGGACTCG | CCCGGGCTGG<br>GGGCCCGACC |
| 18451 | TGCAGTTTGC<br>ACGTCAAACG  | CCGCGCCACC<br>GGCGCGGTGG  | GAGACGTACT<br>CTCTGCATGA | TCAGCCTGAA<br>AGTCGGACTT | TAACAAGTTT<br>ATTGTTCAAA |
| 18501 | AGAAACCCCA<br>TCTTTGGGGT  | CGGTGGCGCC<br>GCCACCGCGG  | TACGCACGAC<br>ATGCGTGCTG | GTGACCACAG<br>CACTGGTGTC | ACCGGTCCCA<br>TGGCCAGGGT |
| 18551 | GCGTTTGACG<br>CGCAAACCTGC | CTGCGGTTC A<br>GACGCCAAGT | TCCCTGTGGA<br>AGGGACACCT | CCGTGAGGAT<br>GGCACTCCTA | ACTGCGTACT<br>TGACGCATGA |
| 18601 | CGTACAAGGC<br>GCATGTTCCG  | GCGGTTCACC<br>CGCCAAGTGG  | CTAGCTGTGG<br>GATCGACACC | GTGATAACCG<br>CACTATTGGC | TGTGCTGGAC<br>ACACGACCTG |
| 18651 | ATGGCTTCCA<br>TACCGAAGGT  | CGTACTTTGA<br>GCATGAAACT  | CATCCGCGGC<br>GTAGGCGCCG | GTGCTGGACA<br>CACGACCTGT | GGGGCCCTAC<br>CCCCGGGATG |
| 18701 | TTTTAAGCCC<br>AAAATTCGGG  | TACTCTGGCA<br>ATGAGACCGT  | CTGCCTACAA<br>GACGGATGTT | CGCCCTGGCT<br>GCGGGACCGA | CCCAAGGGTG<br>GGGTTCCAC  |
| 18751 | CCCCAAATCC<br>GGGGTTTAGG  | TTGCGAATGG<br>AACGCTTACC  | GATGAAGCTG<br>CTACTTCGAC | CTACTGCTCT<br>GATGACGAGA | TGAAATAAAC<br>ACTTTATTTG |
| 18801 | CTAGAAGAAG<br>GATCTTCTTC  | AGGACGATGA<br>TCCTGCTACT  | CAACGAAGAC<br>GTTGCTTCTG | GAAGTAGACG<br>CTTCATCTGC | AGCAAGCTGA<br>TCGTTCGACT |
| 18851 | GCAGCAAAAA<br>CGTCGTTTTT  | ACTCACGTAT<br>TGAGTGCATA  | TTGGGCAGGC<br>AACCCGTCCG | GCCTTATTCT<br>CGGAATAAGA | GGTATAAATA<br>CCATATTTAT |
| 18901 | TTACAAAGGA<br>AATGTTTCCT  | GGGTATTCAA<br>CCCATAGT    | ATAGGTGTCG<br>TATCCACAGC | AAGGTCAAAC<br>TTCCAGTTTG | ACCTAAATAT<br>TGGATTTATA |
| 18951 | GCCGATAAAA<br>CGGCTATTTT  | CATTTCAACC<br>GTAAAGTTGG  | TGAACCTCAA<br>ACTTGGAGTT | ATAGGAGAAT<br>TATCCTCTTA | CTCAGTGGTA<br>GAGTCACCAT |
| 19001 | CGAAACAGAA<br>GCTTTGTCTT  | ATTAATCATG<br>TAATTAGTAC  | CAGCTGGGAG<br>GTCGACCCTC | AGTCCTAAAA<br>TCAGGATTTT | AAGACTACCC<br>TTCTGATGGG |
| 19051 | CAATGAAACC<br>GTTACTTTGG  | ATGTTACGGT<br>TACAATGCCA  | TCATATGCAA<br>AGTATACGTT | AACCCACAAA<br>TTGGGTGTTT | TGAAAATGGA<br>ACTTTTACCT |
| 19101 | GGGCAAGGCA<br>CCCGTTCCGT  | TTCTTGTA<br>AAGAACATTT    | GCAACAAAAT<br>CGTTGTTTTA | GGAAAGCTAG<br>CCTTTCGATC | AAAGTCAAGT<br>TTTCAGTTCA |
| 19151 | GGAAATGCAA<br>CCTTTACGTT  | TTTTTCTCAA<br>AAAAAGAGTT  | CTACTGAGGC<br>GATGACTCCG | AGCCGCAGGC<br>TCGGCGTCCG | AATGGTGATA<br>TTACCACTAT |

FIG. 10A-24



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|       |             |              |            |             |             |
|-------|-------------|--------------|------------|-------------|-------------|
| 19201 | ACTTGACTCC  | TAAAGTGGTA   | TTGTACAGTG | AAGATGTAGA  | TATAGAAACC  |
|       | TGAACTGAGG  | ATTCACCAT    | AACATGTCAC | TTCTACATCT  | ATATCTTTGG  |
| 19251 | CCAGACACTC  | ATATTTCTTA   | CATGCCCACT | ATTAAGGAAG  | GTAACTCACG  |
|       | GGTCTGTGAG  | TATAAAGAAT   | GTACGGGTGA | TAATTCCTTC  | CATTGAGTGC  |
| 19301 | AGAACTAATG  | GGCCAACAAT   | CTATGCCCAA | CAGGCCTAAT  | TACATTGCTT  |
|       | TCTTGATTAC  | CCGGTTGTTA   | GATACGGGTT | GTCCGGATTA  | ATGTAACGAA  |
| 19351 | TTAGGGACAA  | TTTTATTGGT   | CTAATGTATT | ACAACAGCAC  | GGGTAATATG  |
|       | AATCCCTGTT  | AAAATAACCA   | GATTACATAA | TGTTGTCGTG  | CCCATTATAC  |
| 19401 | GGTGTTCTGG  | CGGGCCAAGC   | ATCGCAGTTG | AATGCTGTTG  | TAGATTTGCA  |
|       | CCACAAGACC  | GCCC GGTTTCG | TAGCGTCAAC | TTACGACAAC  | ATCTAAACGT  |
| 19451 | AGACAGAAAC  | ACAGAGCTTT   | CATACCAGCT | TTTGCTTGAT  | TCCATTGGTG  |
|       | TCTGTCCTTG  | TGTCTCGAAA   | GTATGGTCGA | AAACGAACTA  | AGGTAACCAC  |
| 19501 | ATAGAACCAG  | GTA CTTTTCT  | ATGTGGAATC | AGGCTGTTGA  | CAGCTATGAT  |
|       | TATCTTGGTC  | CATGAAAAGA   | TACACCTTAG | TCCGACAAC T | GTCGATACTA  |
| 19551 | CCAGATGTTA  | GAATTATTGA   | AAATCATGGA | ACTGAAGATG  | AACTTCCAAA  |
|       | GGTCTACAAT  | CTTAATAACT   | TTTAGTACCT | TGACTTCTAC  | TTGAAGGTTT  |
| 19601 | TTACTGCTTT  | CCACTGGGAG   | GTGTGATTAA | TACAGAGACT  | CTTACCAAGG  |
|       | AATGACGAAA  | GGTGACCCTC   | CACACTAATT | ATGTCTCTGA  | GAATGGTTCC  |
| 19651 | TAAAACCTAA  | AACAGGTCAG   | GAAAATGGAT | GGGAAAAGA   | TGCTACAGAA  |
|       | ATTTTGGATT  | TTGTCCAGTC   | CTTTTACCTA | CCCTTTTTCT  | ACGATGTCTT  |
| 19701 | TTTTTCAGATA | AAAATGAAAT   | AAGAGTTGGA | AATAATTTTG  | CCATGGAAAT  |
|       | AAAAGTCTAT  | TTTTACTTTA   | TTCTCAACCT | TTATTA AAC  | GGTACCTTTA  |
| 19751 | CAATCTAAAT  | GCCAACCTGT   | GGAGAAATTT | CCTGTACTCC  | AACATAGCGC  |
|       | GTTAGATTTA  | CGGTTGGACA   | CCTCTTTAAA | GGACATGAGG  | TTGTATCGCG  |
| 19801 | TGTATTTGCC  | CGACAAGCTA   | AAGTACAGTC | CTTCCAACGT  | AAAAATTTCT  |
|       | ACATAAACGG  | GCTGTTTCGAT  | TTCATGTCAG | GAAGGTTGCA  | TTTTTAAAGA  |
| 19851 | GATAACCCAA  | ACACCTACGA   | CTACATGAAC | AAGCGAGTGG  | TGGCTCCCGG  |
|       | CTATTGGGTT  | TGTGGATGCT   | GATGTACTTG | TTGCTCACC   | ACCGAGGGCC  |
| 19901 | GCTAGTGGAC  | TGCTACATTA   | ACCTTGGAGC | ACGCTGGTCC  | CTTGACTATA  |
|       | CGATCACCTG  | ACGATGTAAT   | TGGAACCTCG | TGCGACCAGG  | GAAC TGATAT |
| 19951 | TGGACAACGT  | CAACCCATTT   | AACCACCACC | GCAATGCTGG  | CCTGCGCTAC  |
|       | ACCTGTTGCA  | GTTGGGTAAA   | TTGGTGGTGG | CGTTACGACC  | GGACGCGATG  |

FIG. 10A-25



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|       |            |            |             |            |             |
|-------|------------|------------|-------------|------------|-------------|
| 20001 | CGCTCAATGT | TGCTGGGCAA | TGGTCGCTAT  | GTGCCCTTCC | ACATCCAGGT  |
|       | GCGAGTTACA | ACGACCCGTT | ACCAGCGATA  | CACGGGAAGG | TGTAGGTCCA  |
| 20051 | GCCTCAGAAG | TTCTTTGCCA | TTAAAAACCT  | CCTTCTCCTG | CCGGGCTCAT  |
|       | CGGAGTCTTC | AAGAAACGGT | AATTTTGGGA  | GGAAGAGGAC | GGCCCCAGTA  |
| 20101 | ACACCTACGA | GTGGAACTTC | AGGAAGGATG  | TTAACATGGT | TCTGCAGAGC  |
|       | TGTGGATGCT | CACCTTGAAG | TCCTTCCTAC  | AATTGTACCA | AGACGTCTCG  |
| 20151 | TCCCTAGGAA | ATGACCTAAG | GGTTGACGGA  | GCCAGCATTG | AGTTTGATAG  |
|       | AGGGATCCTT | TACTGGATTC | CCAACCTGCCT | CGGTCGTAAT | TCAAACCTATC |
| 20201 | CATTTGCCTT | TACGCCACCT | TCTTCCCCAT  | GGCCCACAAC | ACCGCCTCCA  |
|       | GTAAACGGAA | ATGCGGTGGA | AGAAGGGGTA  | CCGGGTGTTG | TGGCGGAGGT  |
| 20251 | CGCTTGAGGC | CATGCTTAGA | AACGACACCA  | ACGACCAGTC | CTTTAACGAC  |
|       | GCGAACTCCG | GTACGAATCT | TTGCTGTGGT  | TGCTGGTCAG | GAAATTGCTG  |
| 20301 | TATCTCTCCG | CCGCCAACAT | GCTCTACCCT  | ATACCCGCCA | ACGCTACCAA  |
|       | ATAGAGAGGC | GGCGGTGTA  | CGAGATGGGA  | TATGGGCGGT | TGCGATGGTT  |
| 20351 | CGTGCCATA  | TCCATCCCCT | CCCGCAACTG  | GGCGGCTTTC | CGCGGCTGGG  |
|       | GCACGGGTAT | AGGTAGGGGA | GGGCGTTGAC  | CCGCCGAAAG | GCGCCGACCC  |
| 20401 | CCTTCACGCG | CCTTAAGACT | AAGGAAACCC  | CATCACTGGG | CTCGGGCTAC  |
|       | GGAAGTGCGC | GGAATTCTGA | TTCTTTGGG   | GTAGTGACCC | GAGCCCGATG  |
| 20451 | GACCCTTATT | ACACCTACTC | TGGCTCTATA  | CCCTACCTAG | ATGGAACCTT  |
|       | CTGGGAATAA | TGTGGATGAG | ACCGAGATAT  | GGGATGGATC | TACCTTGGAA  |
| 20501 | TTACCTCAAC | CACACCTTTA | AGAAGGTGGC  | CATTACCTTT | GACTCTTCTG  |
|       | AATGGAGTTG | GTGTGGAAAT | TCTTCCACCG  | GTAATGGAAA | CTGAGAAGAC  |
| 20551 | TCAGCTGGCC | TGGCAATGAC | CGCCTGCTTA  | CCCCAACGA  | GTTTGAAATT  |
|       | AGTCGACCGG | ACCGTTACTG | GCGGACGAAT  | GGGGGTGCT  | CAAACCTTAA  |
| 20601 | AAGCGCTCAG | TTGACGGGGA | GGGTTACAAC  | GTTGCCCAGT | GTAACATGAC  |
|       | TTGCGGAGTC | AACTGCCCTT | CCCAATGTTG  | CAACGGGTCA | CATTGTACTG  |
| 20651 | CAAAGACTGG | TTCCTGGTAC | AAATGCTAGC  | TAACATAAAC | ATTGGCTACC  |
|       | GTTTCTGACC | AAGGACCATG | TTTACGATCG  | ATTGATATTG | TAACCGATGG  |
| 20701 | AGGGCTTCTA | TATCCCAGAG | AGCTACAAGG  | ACCGCATGTA | CTCCTTCTTT  |
|       | TCCCGAAGAT | ATAGGGTCTC | TCGATGTTCC  | TGGCGTACAT | GAGGAAGAAA  |
| 20751 | AGAAACTTCC | AGCCCATGAG | CCGTCAGGTG  | GTGGATGATA | CTAAATACAA  |
|       | TCTTTGAAGG | TCGGGTACTC | GGCAGTCCAC  | CACCTACTAT | GATTTATGTT  |

FIG. 10A-26



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|       |            |             |             |            |            |
|-------|------------|-------------|-------------|------------|------------|
| 20801 | GGACTACCAA | CAGGTGGGCA  | TCCTACACCA  | ACACAACAAC | TCTGGATTTG |
|       | CCTGATGGTT | GTCCACCCGT  | AGGATGTGGT  | TGTGTTGTTG | AGACCTAAAC |
| 20851 | TTGGCTACCT | TGCCCCCACC  | ATGCGCGAAG  | GACAGGCCTA | CCCTGCTAAC |
|       | AACCGATGGA | ACGGGGGTGG  | TACGCGCTTC  | CTGTCCGGAT | GGGACGATTG |
| 20901 | TTCCCCTATC | CGCTTATAGG  | CAAGACCGCA  | GTTGACAGCA | TTACCCAGAA |
|       | AAGGGGATAG | GCGAATATCC  | GTTCTGGCGT  | CAACTGTCGT | AATGGGTCTT |
| 20951 | AAAGTTTCTT | TGCGATCGCA  | CCCTTTGGCG  | CATCCCATTG | TCCAGTAACT |
|       | TTTCAAAGAA | ACGCTAGCGT  | GGGAAACCGC  | GTAGGGTAAG | AGGTCATTGA |
| 21001 | TTATGTCCAT | GGGCGCACTC  | ACAGACCTGG  | GCCAAAACCT | TCTCTACGCC |
|       | AATACAGGTA | CCCGCGTGAG  | TGCTCTGGACC | CGGTTTTGGA | AGAGATGCGG |
| 21051 | AACTCCGCC  | ACGCGCTAGA  | CATGACTTTT  | GAGGTGGATC | CCATGGACGA |
|       | TTGAGGCGGG | TGCGCGATCT  | GTACTGAAAA  | CTCCACCTAG | GGTACCTGCT |
| 21101 | GCCCACCCTT | CTTTATGTTT  | TGTTTGAAGT  | CTTTGACGTG | GTCCGTGTGC |
|       | CGGGTGGGAA | GAAATACAAA  | ACAAACTTCA  | GAAACTGCAC | CAGGCACACG |
| 21151 | ACCAGCCGCA | CCGCGGCGTC  | ATCGAAACCG  | TGTACCTGCG | CACGCCCTTC |
|       | TGGTCGGCGT | GGCGCCGCAG  | TAGCTTTGGC  | ACATGGACGC | GTGCGGGAAG |
| 21201 | TCGGCCGGCA | ACGCCACAAC  | ATAAAGAAGC  | AAGCAACATC | AACAACAGCT |
|       | AGCCGGCCGT | TGCGGTGTTG  | TATTTCTTCG  | TTGTTGTAG  | TTGTTGTCGA |
| 21251 | GCCGCCATGG | GCTCCAGTGA  | GCAGGAACTG  | AAAGCCATTG | TCAAAGATCT |
|       | CGGCGGTACC | CGAGGTCACT  | CGTCCTTGAC  | TTTCGGTAAC | AGTTTCTAGA |
| 21301 | TGGTTGTGGG | CCATATTTTT  | TGGGCACCTA  | TGACAAGCGC | TTTCCAGGCT |
|       | ACCAACACCC | GGTATAAAAA  | ACCCGTGGAT  | ACTGTTCGCG | AAAGGTCCGA |
| 21351 | TTGTTTCTCC | ACACAAGCTC  | GCCTGCGCCA  | TAGTCAATAC | GGCCGGTCGC |
|       | AACAAAGAGG | TGTGTTTCGAG | CGGACGCGGT  | ATCAGTTATG | CCGGCCAGCG |
| 21401 | GAGACTGGGG | GCGTACACTG  | GATGGCCTTT  | GCCTGGAACC | CGCACTCAA  |
|       | CTCTGACCCC | CGCATGTGAC  | CTACCGGAAA  | CGGACCTTGG | GCGTGAGTTT |
| 21451 | AACATGCTAC | CTCTTTGAGC  | CCTTTGGCTT  | TTCTGACCAG | CGACTCAAGC |
|       | TTGTACGATG | GAGAAACTCG  | GGAAACCGAA  | AAGACTGGTC | GCTGAGTTCG |
| 21501 | AGGTTTACCA | GTTTGAGTAC  | GAGTCACTCC  | TGCGCCGTAG | CGCCATTGCT |
|       | TCCAAATGGT | CAAACCTCATG | CTCAGTGAGG  | ACGCGGCATC | GCGGTAACGA |
| 21551 | TCTTCCCCCG | ACCGCTGTAT  | AACGCTGGAA  | AAGTCCACCC | AAAGCGTACA |
|       | AGAAGGGGGC | TGGCGACATA  | TTGCGACCTT  | TTCAGGTGGG | TTTCGCATGT |

FIG. 10A-27



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|       |             |            |            |             |            |
|-------|-------------|------------|------------|-------------|------------|
| 21601 | GGGGCCCAAC  | TCGGCCGCCT | GTGGACTATT | CTGCTGCATG  | TTTCTCCACG |
|       | CCCCGGGTTG  | AGCCGGCGGA | CACCTGATAA | GACGACGTAC  | AAAGAGGTGC |
| 21651 | CCTTTGCCAA  | CTGGCCCCAA | ACTCCCATGG | ATCACAACCC  | CACCATGAAC |
|       | GGAAACGGTT  | GACCGGGGTT | TGAGGGTACC | TAGTGTGGG   | GTGGTACTTG |
| 21701 | CTTATTACCG  | GGGTACCCAA | CTCCATGCTC | AACAGTCCCC  | AGGTACAGCC |
|       | GAATAATGGC  | CCCATGGGTT | GAGGTACGAG | TTGTCAGGGG  | TCCATGTCGG |
| 21751 | CACCCTGCGT  | CGCAACCAGG | AACAGCTCTA | CAGCTTCCTG  | GAGCGCCACT |
|       | GTGGGACGCA  | GCGTTGGTCC | TTGTGAGAT  | GTCGAAGGAC  | CTCGCGGTGA |
| 21801 | CGCCCTACTT  | CCGCAGCCAC | AGTGCGCAGA | TTAGGAGCGC  | CACTTCTTTT |
|       | GCGGGATGAA  | GGCGTCGGTG | TCACGCGTCT | AATCCTCGCG  | GTGAAGAAAA |
| 21851 | TGTCACTTGA  | AAAACATGTA | AAAATAATGT | ACTAGAGACA  | CTTTCAATAA |
|       | ACAGTGAACT  | TTTTGTACAT | TTTTATTACA | TGATCTCTGT  | GAAAGTTATT |
| 21901 | AGGCAAATGC  | TTTTATTTGT | ACACTCTCGG | GTGATTATTT  | ACCCCCACCC |
|       | TCCGTTTACG  | AAAATAAACA | TGTGAGAGCC | CACTAATAAA  | TGGGGGTGGG |
| 21951 | TTGCCGTCTG  | CGCCGTTTAA | AAATCAAAGG | GGTTCTGCCG  | CGCATCGCTA |
|       | AACGGCAGAC  | GCGGCAAATT | TTTAGTTTCC | CCAAGACGGC  | GCGTAGCGAT |
| 22001 | TGCGCCACTG  | GCAGGGACAC | GTTGCGATAC | TGGTGTTTAG  | TGCTCCACTT |
|       | ACGCGGTGAC  | CGTCCCTGTG | CAACGCTATG | ACCACAAATC  | ACGAGGTGAA |
| 22051 | AAACTCAGGC  | ACAACCATCC | GCGGCAGCTC | GGTGAAGTTT  | TCACTCCACA |
|       | TTTGAGTCCG  | TGTTGGTAGG | CGCCGTCGAG | CCACTTCAA   | AGTGAGGTGT |
| 22101 | GGCTGCGCAC  | CATCACCAAC | GCGTTTAGCA | GGTCGGGCGC  | CGATATCTTG |
|       | CCGACGCGTG  | GTAGTGGTTG | CGCAAATCGT | CCAGCCC GCG | GCTATAGAAC |
| 22151 | AAGTCGCAGT  | TGGGGCCTCC | GCCCTGCGCG | CGCGAGTTGC  | GATACACAGG |
|       | TTCAGCGTCA  | ACCCCGGAGG | CGGGACGCGC | GCGCTCAACG  | CTATGTGTCC |
| 22201 | GTTGCAGCAC  | TGGAACACTA | TCAGCGCCGG | GTGGTGCACG  | CTGGCCAGCA |
|       | CAACGTCGTG  | ACCTTGTGAT | AGTCGCGGCC | CACCACGTGC  | GACCGGTCGT |
| 22251 | CGCTCTTGTC  | GGAGATCAGA | TCCGCGTCCA | GGTCCTCCGC  | GTTGCTCAGG |
|       | GCGAGAACAG  | CCTCTAGTCT | AGGCGCAGGT | CCAGGAGGCG  | CAACGAGTCC |
| 22301 | GCGAACGGAG  | TCAACTTTGG | TAGCTGCCTT | CCCAAAAAGG  | GCGCGTGCCC |
|       | CGCTTGCCCTC | AGTTGAAACC | ATCGACGGAA | GGGTTTTTCC  | CGCGCACGGG |
| 22351 | AGGCTTTGAG  | TTGCACTCGC | ACCGTAGTGG | CATCAAAAAGG | TGACCGTGCC |
|       | TCCGAAACTC  | AACGTGAGCG | TGGCATCACC | GTAGTTTTCC  | ACTGGCACGG |

FIG. 10A-28



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|       |            |            |            |            |             |
|-------|------------|------------|------------|------------|-------------|
| 22401 | CGGTCTGGGC | GTTAGGATAC | AGCGCCTGCA | TAAAAGCCTT | GATCTGCTTA  |
|       | GCCAGACCCG | CAATCCTATG | TCGCGGACGT | ATTTTCGGAA | CTAGACGAAT  |
| 22451 | AAAGCCACCT | GAGCCTTTGC | GCCTTCAGAG | AAGAACATGC | CGCAAGACTT  |
|       | TTTCGGTGGG | CTCGGAAACG | CGGAAGTCTC | TTCTTGTACC | GCGTTCTGAA  |
| 22501 | GCCGGAAAAC | TGATTGGCCG | GACAGGCCGC | GTCGTGCACG | CAGCACCTTG  |
|       | CGGCCTTTTG | ACTAACCGGC | CTGTCCGGCG | CAGCACGTGC | GTCGTGGAAC  |
| 22551 | CGTCGGTGTT | GGAGATCTGC | ACCACATTTT | GGCCCCACCG | GTTCTTCACG  |
|       | GCAGCCACAA | CCTCTAGACG | TGGTGTAAAG | CCGGGGTGGC | CAAGAAGTGC  |
| 22601 | ATCTTGGCCT | TGCTAGACTG | CTCCTTCAGC | GCGCGCTGCC | CGTTTTTCGCT |
|       | TAGAACCGGA | ACGATCTGAC | GAGGAAGTCG | CGCGCGACGG | GCAAAAGCGA  |
| 22651 | CGTCACATCC | ATTTCAATCA | CGTGCTCCTT | ATTTATCATA | ATGCTTCCGT  |
|       | GCAGTGTAGG | TAAAGTTAGT | GCACGAGGAA | TAAATAGTAT | TACGAAGGCA  |
| 22701 | GTAGACACTT | AAGCTCGCCT | TCGATCTCAG | CGCAGCGGTG | CAGCCACAAC  |
|       | CATCTGTGAA | TTCGAGCGGA | AGCTAGAGTC | GCGTCGCCAC | GTCGGTGTTG  |
| 22751 | GCGCAGCCCG | TGGGCTCGTG | ATGCTTGTAG | GTCACCTCTG | CAAACGACTG  |
|       | CGCGTCGGGC | ACCCGAGCAC | TACGAACATC | CAGTGGAGAC | GTTTGCTGAC  |
| 22801 | CAGGTACGCC | TGCAGGAATC | GCCCCATCAT | CGTCACAAAG | GTCTTGTTGC  |
|       | GTCCATGCGG | ACGTCCTTAG | CGGGGTAGTA | GCAGTGTTTC | CAGAACAACG  |
| 22851 | TGGTGAAGGT | CAGCTGCAAC | CCGCGGTGCT | CCTCGTTCAG | CCAGGTCTTG  |
|       | ACCACTTCCA | GTCGACGTTG | GGCGCCACGA | GGAGCAAGTC | GGTCCAGAAC  |
| 22901 | CATACGGCCG | CCAGAGCTTC | CACTTGGTCA | GGCAGTAGTT | TGAAGTTCGC  |
|       | GTATGCCGGC | GGTCTCGAAG | GTGAACCAGT | CCGTCATCAA | ACTTCAAGCG  |
| 22951 | CTTTAGATCG | TTATCCACGT | GGTACTTGTC | CATCAGCGCG | CGCGCAGCCT  |
|       | GAAATCTAGC | AATAGGTGCA | CCATGAACAG | GTAGTCGCGC | GCGCGTCGGA  |
| 23001 | CCATGCCCTT | CTCCCACGCA | GACACGATCG | GCACACTCAG | CGGGTTCATC  |
|       | GGTACGGGAA | GAGGGTGCCT | CTGTGCTAGC | CGTGTGAGTC | GCCCAAGTAG  |
| 23051 | ACCGTAATTT | CACTTTCCGC | TTCGCTGGGC | TCTTCCTCTT | CCTCTTGCGT  |
|       | TGGCATTAAA | GTGAAAGGCG | AAGCGACCCG | AGAAGGAGAA | GGAGAACGCA  |
| 23101 | CCGCATACCA | CGCGCCACTG | GGTCGTCTTC | ATTCAGCCGC | CGCACTGTGC  |
|       | GGCGTATGGT | GCGCGGTGAC | CCAGCAGAAG | TAAGTCGGCG | GCGTGACACG  |
| 23151 | GCTTACCTCC | TTTGCCATGC | TTGATTAGCA | CCGGTGGGTT | GCTGAAACCC  |
|       | CGAATGGAGG | AAACGGTACG | AACTAATCGT | GGCCACCCAA | CGACTTTGGG  |

FIG. 10A-29



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|       |                           |                          |                           |                          |                          |
|-------|---------------------------|--------------------------|---------------------------|--------------------------|--------------------------|
| 23201 | ACCATTTGTA<br>TGGTAAACAT  | GCGCCACATC<br>CGCGGTGTAG | TTCTCTTTCT<br>AAGAGAAAGA  | TCCTCGCTGT<br>AGGAGCGACA | CCACGATTAC<br>GGTGCTAATG |
| 23251 | CTCTGGTGAT<br>GAGACCACTA  | GGCGGGCGCT<br>CCGCCCCCGA | CGGGCTTGGG<br>GCCCCGAACCC | AGAAGGGCGC<br>TCTTCCCCCG | TTCTTTTTCT<br>AAGAAAAAGA |
| 23301 | TCTTGGGCGC<br>AGAACCCGCG  | AATGGCCAAA<br>TTACCGGTTT | TCCGCCGCCG<br>AGGCGGCGGC  | AGGTCGATGG<br>TCCAGCTACC | CCGCGGGCTG<br>GGCGCCCGAC |
| 23351 | GGTGTGCGCG<br>CCACACGCGC  | GCACCAGCGC<br>CGTGGTCGCG | GTCTTGTGAT<br>CAGAACACTA  | GAGTCTTCCT<br>CTCAGAAGGA | CGTCCTCGGA<br>GCAGGAGCCT |
| 23401 | CTCGATACGC<br>GAGCTATGCG  | CGCCTCATCC<br>GCGGAGTAGG | GCTTTTTTGG<br>CGAAAAAACC  | GGGCGCCCGG<br>CCCGCGGGCC | GGAGGCGGGC<br>CCTCCGCCGC |
| 23451 | GCGACGGGGA<br>CGCTGCCCCCT | CGGGGACGAC<br>GCCCTGCTG  | ACGTCCTCCA<br>TGCAGGAGGT  | TGGTTGGGGG<br>ACCAACCCCC | ACGTCGCGCC<br>TGCAGCGCGG |
| 23501 | GCACCGCGTC<br>CGTGGCGCAG  | CGCGCTCGGG<br>GCGCGAGCCC | GGTGGTTTCG<br>CCACCAAAGC  | CGCTGCTCCT<br>GCGACGAGGA | CTTCCCGACT<br>GAAGGGCTGA |
| 23551 | GGCCATTTCC<br>CCGGTAAAGG  | TTCTCCTATA<br>AAGAGGATAT | GGCAGAAAAA<br>CCGTCTTTTT  | GATCATGGAG<br>CTAGTACCTC | TCAGTCGAGA<br>AGTCAGCTCT |
| 23601 | AGAAGGACAG<br>TCTTCCTGTC  | CCTAACCGCC<br>GGATTGGCGG | CCCTCTGAGT<br>GGGAGACTCA  | TCGCCACCAC<br>AGCGGTGGTG | CGCCTCCACC<br>GCGGAGGTGG |
| 23651 | GATGCCGCCA<br>CTACGGCGGT  | ACGCGCCTAC<br>TGCGCGGATG | CACCTTCCCC<br>GTGGAAGGGG  | GTCGAGGCAC<br>CAGCTCCGTG | CCCCGCTTGA<br>GGGGCGAACT |
| 23701 | GGAGGAGGAA<br>CCTCCTCCTT  | GTGATTATCG<br>CACTAATAGC | AGCAGGACCC<br>TCGTCCTGGG  | AGGTTTTGTA<br>TCCAAAACAT | AGCGAAGACG<br>TCGCTTCTGC |
| 23751 | ACGAGGACCG<br>TGCTCCTGGC  | CTCAGTACCA<br>GAGTCATGGT | ACAGAGGATA<br>TGTCTCCTAT  | AAAAGCAAGA<br>TTTTCGTTCT | CCAGGACAAC<br>GGTCCTGTTG |
| 23801 | GCAGAGGCAA<br>CGTCTCCGTT  | ACGAGGAACA<br>TGCTCCTTGT | AGTCGGGCGG<br>TCAGCCCGCC  | GGGGACGAAA<br>CCCCTGCTTT | GGCATGGCGA<br>CCGTACCGCT |
| 23851 | CTACCTAGAT<br>GATGGATCTA  | GTGGGAGACG<br>CACCTCTGCT | ACGTGCTGTT<br>TGCACGACAA  | GAAGCATCTG<br>CTTCGTAGAC | CAGCGCCAGT<br>GTCGCGGTCA |
| 23901 | GCGCCATTAT<br>CGCGGTAATA  | CTGCGACGCG<br>GACGCTGCGC | TTGCAAGAGC<br>AACGTTCTCG  | GCAGCGATGT<br>CGTCGCTACA | GCCCCTCGCC<br>CGGGGAGCGG |
| 23951 | ATAGCGGATG<br>TATCGCCTAC  | TCAGCCTTGC<br>AGTCGGAACG | CTACGAACGC<br>GATGCTTGCG  | CACCTATTCT<br>GTGGATAAGA | CACCGCGCGT<br>GTGGCGCGCA |

FIG. 10A-30



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|       |                          |                          |                          |                          |                          |
|-------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|
| 24001 | ACCCCCAAA<br>TGGGGGGTTT  | CGCCAAGAAA<br>GCGGTTCTTT | ACGGCACATG<br>TGCCGTGTAC | CGAGCCCAAC<br>GCTCGGGTTG | CCGCGCCTCA<br>GGCGCGGAGT |
| 24051 | ACTTCTACCC<br>TGAAGATGGG | CGTATTTGCC<br>GCATAAACGG | GTGCCAGAGG<br>CACGGTCTCC | TGCTTGCCAC<br>ACGAACGGTG | CTATCACATC<br>GATAGTGTAG |
| 24101 | TTTTTCCAAA<br>AAAAAGGTTT | ACTGCAAGAT<br>TGACGTTCTA | ACCCCTATCC<br>TGGGGATAGG | TGCCGTGCCA<br>ACGGCACGGT | ACCGCAGCCG<br>TGGCGTCGGC |
| 24151 | AGCGGACAAG<br>TCGCCTGTTC | CAGCTGGCCT<br>GTCGACCGGA | TGCGGCAGGG<br>ACGCCGTCCC | CGCTGTCATA<br>GCGACAGTAT | CCTGATATCG<br>GGACTATAGC |
| 24201 | CCTCGCTCAA<br>GGAGCGAGTT | CGAAGTGCCA<br>GCTTCACGGT | AAAATCTTTG<br>TTTTAGAAAC | AGGGTCTTGG<br>TCCCAGAACC | ACGCGACGAG<br>TGCGCTGCTC |
| 24251 | AAGCGCGCGG<br>TTCGCGCGCC | CAAACGCTCT<br>GTTTGCGAGA | GCAACAGGAA<br>CGTTGTCCTT | AACAGCGAAA<br>TTGTCGCTTT | ATGAAAGTCA<br>TACTTTCAGT |
| 24301 | CTCTGGAGTG<br>GAGACCTCAC | TTGGTGGAAC<br>AACCACCTTG | TCGAGGGTGA<br>AGCTCCCACT | CAACGCGCGC<br>GTTGCGCGCG | CTAGCCGTAC<br>GATCGGCATG |
| 24351 | TAAAACGCAG<br>ATTTTGCATC | CATCGAGGTC<br>GTAGCTCCAG | ACCCACTTTG<br>TGGGTGAAAC | CCTACCCGGC<br>GGATGGGCCG | ACTTAACCTA<br>TGAATTGGAT |
| 24401 | CCCCCAAGG<br>GGGGGGTTCC  | TCATGAGCAC<br>AGTACTCGTG | AGTCATGAGT<br>TCAGTACTCA | GAGCTGATCG<br>CTCGACTAGC | TGCGCCGTGC<br>ACGCGGCACG |
| 24451 | GCAGCCCCTG<br>CGTCGGGGAC | GAGAGGGATG<br>CTCTCCCTAC | CAAATTTGCA<br>GTTTAAACGT | AGAACAAACA<br>TCTTGTTTGT | GAGGAGGGCC<br>CTCCTCCCGG |
| 24501 | TACCCGCAGT<br>ATGGGCGTCA | TGGCGACGAG<br>ACCGCTGCTC | CAGCTAGCGC<br>GTCGATCGCG | GCTGGCTTCA<br>CGACCGAAGT | AACGCGCGAG<br>TTGCGCGCTC |
| 24551 | CCTGCCGACT<br>GGACGGCTGA | TGGAGGAGCG<br>ACCTCCTCGC | ACGCAAATA<br>TGCGTTTGAT  | ATGATGGCCG<br>TACTACCGGC | CAGTGCTCGT<br>GTCACGAGCA |
| 24601 | TACCGTGGAG<br>ATGGCACCTC | CTTGAGTGCA<br>GAACTCACGT | TGCAGCGGTT<br>ACGTCGCCAA | CTTTGCTGAC<br>GAAACGACTG | CCGGAGATGC<br>GGCCTCTACG |
| 24651 | AGCGCAAGCT<br>TCGCGTTCGA | AGAGGAAACA<br>TCTCCTTTGT | TTGCACTACA<br>AACGTGATGT | CCTTTCGACA<br>GGAAAGCTGT | GGGCTACGTA<br>CCCGATGCAT |
| 24701 | CGCCAGGCCT<br>GCGGTCCGGA | GCAAGATCTC<br>CGTTCTAGAG | CAACGTGGAG<br>GTTGCACCTC | CTCTGCAACC<br>GAGACGTTGG | TGGTCTCCTA<br>ACCAGAGGAT |
| 24751 | CCTTGGAATT<br>GGAACCTTAA | TTGCACGAAA<br>AACGTGCTTT | ACCGCCTTGG<br>TGGCGGAACC | GCAAAACGTG<br>CGTTTTGCAC | CTTCATTCCA<br>GAAGTAAGGT |

FIG. 10A-31



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24801 CGCTCAAGGG CGAGGCGCGC CGCGACTACG TCCGCGACTG CGTTTACTTA  
 GCGAGTCCC GCTCCGCGCG GCGCTGATGC AGGCGCTGAC GCAAATGAAT

24851 TTTCTATGCT ACACCTGGCA GACGGCCATG GCGGTTTGGC AGCAGTGCTT  
 AAAGATACGA TGTGGACCGT CTGCCGGTAC CCGCAAACCG TCGTCACGAA

24901 GGAGGAGTGC AACCTCAAGG AGCTGCAGAA ACTGCTAAAG CAAAACCTGA  
 CCTCCTCACG TTGGAGTTCC TCGACGTCTT TGACGATTTT GTTTTGAACT

24951 AGGACCTATG GACGGCCTTC AACGAGCGCT CCGTGGCCGC GCACCTGGCG  
 TCCTGGATAC CTGCCGGAAG TTGCTCGCGA GGCACCGGCG CGTGGACCGC

25001 GACATCATT TCCCCGAACG CCTGCTTAAA ACCCTGCAAC AGGGTCTGCC  
 CTGTAGTAAA AGGGGCTTGC GGACGAATTT TGGGACGTTG TCCAGACGG

25051 AGACTTCACC AGTCAAAGCA TGTTCAGAA CTTTAGGAAC TTTATCCTAG  
 TCTGAAGTGG TCAGTTTCGT ACAACGTCTT GAAATCCTTG AAATAGGATC

25101 AGCGCTCAGG AATCTTGCCC GCCACCTGCT GTGCACTTCC TAGCGACTTT  
 TCGCGAGTCC TTAGAACGGG CGGTGGACGA CACGTGAAGG ATCGCTGAAA

25151 GTGCCCATTA AGTACCGCGA ATGCCCTCCG CCGCTTTGGG GCCACTGCTA  
 CACGGGTAAT TCATGGCGCT TACGGGAGGC GGCGAAACCC CGGTGACGAT

25201 CCTTCTGCAG CTAGCCAACCT ACCTTGCCTA CCACTCTGAC ATAATGGAAG  
 GGAAGACGTC GATCGGTTGA TGGAACGGAT GGTGAGACTG TATTACCTTC

25251 ACGTGAGCGG TGACGGTCTA CTGGAGTGTC ACTGTCGCTG CAACCTATGC  
 TGCACTCGCC ACTGCCAGAT GACCTCACAG TGACAGCGAC GTTGGATACG

25301 ACCCCGCACC GCTCCCTGGT TTGCAATTCG CAGCTGCTTA ACGAAAGTCA  
 TGGGGCGTGG CGAGGGACCA AACGTTAAGC GTCGACGAAT TGCTTTCAGT

25351 AATTATCGGT ACCTTTGAGC TGCAGGGTCC CTCGCCTGAC GAAAAGTCCG  
 TTAATAGCCA TGGAAACTCG ACGTCCCAGG GAGCGGACTG CTTTTTCAGGC

25401 CGGCTCCGGG GTTGAAACTC ACTCCGGGGC TGTGGACGTC GGCTTACCTT  
 GCCGAGGCC CAACTTTGAG TGAGGCCCCG ACACCTGCAG CCGAATGGAA

25451 CGCAAATTTG TACCTGAGGA CTACCACGCC CACGAGATTA GGTTCTACGA  
 GCGTTTAAAC ATGGACTCCT GATGGTGCGG GTGCTCTAAT CCAAGATGCT

25501 AGACCAATCC CGCCCGCCTA ATGCGGAGCT TACCGCCTGC GTCATTACCC  
 TCTGGTTAGG GCGGGCGGAT TACGCCTCGA ATGGCGGACG CAGTAATGGG

25551 AGGGCCACAT TCTTGGCCAA TTGCAAGCCA TCAACAAAGC CCGCCAAGAG  
 TCCCGGTGTA AGAACCAGGT AACGTTCCGGT AGTTGTTTCG GCGGGTCTC

FIG. 10A-32



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25601 TTTCTGCTAC GAAAGGGACG GGGGGTTTAC TTGGACCCCC AGTCCGGCGA  
 AAAGACGATG CTTTCCCTGC CCCCAAATG AACCTGGGGG TCAGGCCGCT

25651 GGAGCTCAAC CCAATCCCCC CGCCGCCGCA GCCCTATCAG CAGCAGCCGC  
 CCTCGAGTTG GGTTAGGGGG GCGGCCGCGT CGGGATAGTC GTCGTCGGCG

25701 GGGCCCTTGC TTCCCAGGAT GGCACCCAAA AAGAAGCTGC AGCTGCCGCC  
 CCCGGGAACG AAGGGTCCTA CCGTGGGTTT TTCTTCGACG TCGACGGCGG

25751 GCCACCCACG GACGAGGAGG AATACTGGGA CAGTCAGGCA GAGGAGGTTT  
 CGGTGGGTGC CTGCTCCTCC TTATGACCCT GTCAGTCCGT CTCCTCCAAA

25801 TGGACGAGGA GGAGGAGGAC ATGATGGAAG ACTGGGAGAG CCTAGACGAG  
 ACCTGCTCCT CCTCCTCCTG TACTACCTTC TGACCCTCTC GGATCTGCTC

25851 GAAGCTTCCG AGGTCGAAGA GGTGTCAGAC GAAACACCGT CACCCTCGGT  
 CTTCGAAGGC TCCAGCTTCT CCACAGTCTG CTTTGTGGCA GTGGGAGCCA

25901 CGCATTCCCC TCGCCGGCGC CCCAGAAATC GGCAACCGGT TCCAGCATGG  
 GCGTAAGGGG AGCGGCCGCG GGGTCTTTAG CCGTTGGCCA AGGTCTGACC

25951 CTACAACCTC CGCTCCTCAG GCGCCGCCGG CACTGCCCGT TCGCCGACCC  
 GATGTTGGAG GCGAGGAGTC CCGGCCGCC GTGACGGGCA AGCGGCTGGG

26001 AACCGTAGAT GGGACACCAC TGGAAACCAGG GCCGGTAAGT CCAAGCAGCC  
 TTGGCATCTA CCCTGTGGTG ACCTTGGTCC CGGCCATTCA GGTTCGTCGG

26051 GCCGCCGTTA GCCCAAGAGC AACAAACAGCG CCAAGGCTAC CGCTCATGGC  
 CGGCGGCAAT CGGGTTCTCG TTGTTGTCGC GGTTCGATG GCGAGTACCG

26101 GCGGGCACAA GAACGCCATA GTTGCTTGCT TGCAAGACTG TGGGGGCAAC  
 CGCCCGTGTT CTTGCGGTAT CAACGAACGA ACGTTCTGAC ACCCCCGTTG

26151 ATCTCCTTCG CCCGCCGCTT TCTTCTCTAC CATCACGGCG TGGCCTTCCC  
 TAGAGGAAGC GGGCGGCGAA AGAAGAGATG GTAGTGCCGC ACCGGAAGGG

26201 CCGTAACATC CTGCATTACT ACCGTCATCT CTACAGCCA TACTGCACCG  
 GGCATTGTAG GACGTAATGA TGGCAGTAGA GATGTCGGGT ATGACGTGGC

26251 GCGGCAGCGG CAGCAACAGC AGCGGCCACA CAGAAGCAA GGCACCGGA  
 CGCCGTCGCC GTCGTTGTCG TCGCCGGTGT GTCTTCGTTT CCGCTGGCCT

26301 TAGCAAGACT CTGACAAAGC CCAAGAAATC CACAGCCGCG GCAGCAGCAG  
 ATCGTTCTGA GACTGTTTCG GGTTCCTTAG GTGTCGCCGC CGTCGTCGTC

26351 GAGGAGGAGC GCTGCGTCTG GCGCCCAACG AACCCGTATC GACCCGCGAG  
 CTCCTCCTCG CGACGCAGAC CGCGGGTTGC TTGGGCATAG CTGGGCGCTC

FIG. 10A-33



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|       |            |            |            |            |             |
|-------|------------|------------|------------|------------|-------------|
| 26401 | CTTAGAAACA | GGATTTTTCC | CACTCTGTAT | GCTATATTTT | AACAGAGCAG  |
|       | GAATCTTTGT | CCTAAAAAGG | GTGAGACATA | CGATATAAAG | TTGTCTCGTC  |
| 26451 | GGCCAAGAA  | CAAGAGCTGA | AAATAAAAAA | CAGGTCTCTG | CGATCCCTCA  |
|       | CCCGGTTCTT | GTTCTCGACT | TTTATTTTTT | GTCCAGAGAC | GCTAGGGAGT  |
| 26501 | CCCGCAGCTG | CCTGTATCAC | AAAAGCGAAG | ATCAGCTTCG | GCGCACGCTG  |
|       | GGCGTTCGAC | GGACATAGTG | TTTTCGCTTC | TAGTCGAAGC | CGCGTGCGAC  |
| 26551 | GAAGACGCGG | AGGCTCTCTT | CAGTAAATAC | TGCGCGCTGA | CTCTTAAGGA  |
|       | CTTCTGCGCC | TCCGAGAGAA | GTCATTTATG | ACGCGCGACT | GAGAATTCCT  |
| 26601 | CTAGTTTCGC | GCCCTTTCTC | AAATTTAAGC | GCGAAACTA  | CGTCATCTCC  |
|       | GATCAAAGCG | CGGGAAAGAG | TTTAAATTCG | CGCTTTTGAT | GCAGTAGAGG  |
| 26651 | AGCGGCCACA | CCCGGCGCCA | GCACCTGTTG | TCAGCGCCAT | TATGAGCAAG  |
|       | TCGCCGGTGT | GGGCCGCGGT | CGTGGACAAC | AGTCGCGGTA | ATACTCGTTC  |
| 26701 | GAAATTCCCA | CGCCCTACAT | GTGGAGTTAC | CAGCCACAAA | TGGGACTTGC  |
|       | CTTTAAGGGT | GCGGGATGTA | CACCTCAATG | GTCGGTGTTT | ACCCTGAACG  |
| 26751 | GGCTGGAGCT | GCCCAAGACT | ACTCAACCCG | AATAAACTAC | ATGAGCGCGG  |
|       | CCGACCTCGA | CGGGTTCTGA | TGAGTTGGGC | TTATTTGATG | TACTCGCGCC  |
| 26801 | GACCCACAT  | GATATCCCGG | GTCAACGGAA | TACGCGCCCA | CCGAAACCGA  |
|       | CTGGGGTGTA | CTATAGGGCC | CAGTTGCCTT | ATGCGCGGGT | GGCTTTGGCT  |
| 26851 | ATTCTCCTGG | AACAGGCGGC | TATTACCACC | ACACCTCGTA | ATAACCTTAA  |
|       | TAAGAGGACC | TTGTCCGCCG | ATAATGGTGG | TGTGGAGCAT | TATTGGAATT  |
| 26901 | TCCCCGTAGT | TGGCCCGCTG | CCCTGGTGTA | CCAGGAAAGT | CCCGCTCCCA  |
|       | AGGGGCATCA | ACCGGGCGAC | GGGACCACAT | GGTCCTTTCA | GGGCGAGGGT  |
| 26951 | CCACTGTGGT | ACTTCCAGAG | GACGCCCAGG | CCGAAGTTCA | GATGACTAAC  |
|       | GGTGACACCA | TGAAGGGTCT | CTGCGGGTCC | GGCTTCAAGT | CTACTGATTG  |
| 27001 | TCAGGGGCGC | AGCTTGCGGG | CGGCTTTCGT | CACAGGGTGC | GGTCGCCCGG  |
|       | AGTCCCCGCG | TCGAACGCC  | GCCGAAAGCA | GTGTCCCACG | CCAGCGGGCC  |
| 27051 | GCAGGGTATA | ACTCACCTGA | CAATCAGAGG | GCGAGGTATT | CAGCTCAACG  |
|       | CGTCCCATAT | TGAGTGGACT | GTTAGTCTCC | CGCTCCATAA | GTCGAGTTGC  |
| 27101 | ACGAGTCGGT | GAGCTCCTCG | CTTGGTCTCC | GTCCGGACGG | GACATTTTCAG |
|       | TGCTCAGCCA | CTCGAGGAGC | GAACCAGAGG | CAGGCCTGCC | CTGTAAAGTC  |
| 27151 | ATCGGCGGCG | CCGGCCGCTC | TTCATTCACG | CCTCGTCAGG | CAATCCTAAC  |
|       | TAGCCGCCGC | GGCCGGCGAG | AAGTAAGTGC | GGAGCAGTCC | GTTAGGATTG  |

FIG. 10A-34



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27201 TCTGCAGACC TCGTCCTCTG AGCCGCGCTC TGGAGGCATT GGA ACTCTGC  
 AGACGTCTGG AGCAGGAGAC TCGGCGCGAG ACCTCCGTAA CCTTGAGACG

27251 AATTTATTGA GGAGTTTGTG CCATCGGTCT ACTTTAACCC CTTCTCGGGA  
 TTAAATAACT CCTCAAACAC GGTAGCCAGA TGA AATTGGG GAAGAGCCCT

27301 CCTCCCGGCC ACTATCCGGA TCAATTTATT CCTAACTTTG ACGCGGTAAA  
 GGAGGGCCGG TGATAGGCCT AGTTAAATAA GGATTGAAAC TGCGCCATTT

27351 GGACTCGGCG GACGGCTACG ACTGAATGTT AAGTGGAGAG GCAGAGCAAC  
 CCTGAGCCGC CTGCCGATGC TGACTTACAA TTCACCTCTC CGTCTCGTTG

27401 TGC GCCTGAA ACACCTGGTC CACTGTCGCC GCCACAAGTG CTTTGCCCGC  
 ACGCGGACTT TGTGGACCAG GTGACAGCGG CGGTGTTTAC GAAACGGGCG

27451 GACTCCGGTG AGTTTTGCTA CTTTGAATTG CCCGAGGATC ATATCGAGGG  
 CTGAGGCCAC TCAAAACGAT GAAACTTAAC GGGCTCCTAG TATAGCTCCC

27501 CCCGGCGCAC GCGGTCCGGC TTACCGCCCA GGGAGAGCTT GCCCGTAGCC  
 GGGCCGCGTG CCGCAGGCCG AATGGCGGGT CCCTCTCGAA CGGGCATCGG

27551 TGATTCGGGA GTTTACCCAG CGCCCCCTGC TAGTTGAGCG GGACAGGGGA  
 ACTAAGCCCT CAAATGGGTC GCGGGGGACG ATCAACTCGC CCTGTCCCCT

27601 CCCTGTGTTT TCACTGTGAT TTGCAACTGT CCTAACCCCTG GATTACATCA  
 GGGACACAAG AGTGACACTA AACGTTGACA GGATTGGGAC CTAATGTAGT

27651 AGATCTTTGT TGCCATCTCT GTGCTGAGTA TAATAAATAC AGAAATTA  
 TCTAGAAACA ACGGTAGAGA CACGACTCAT ATTATTTATG TCTTTAATTT

27701 ATATACTGGG GCTCCTATCG CCATCCTGTA AACGCCACCG TCTTCACCCG  
 TATATGACCC CGAGGATAGC GGTAGGACAT TTGCGGTGGC AGAAGTGGGC

27751 CCAAGCAAA CCAAGGCGAA CCTTACCTGG TACTTTTAAAC ATCTCTCCCT  
 GGGTTCGTTT GGTTCGCTT GGAATGGACC ATGAAAATTG TAGAGAGGGA

27801 CTGTGATTTA CAACAGTTTC AACCCAGACG GAGTGAGTCT ACGAGAGAAC  
 GACACTAAAT GTTGTCAAAG TTGGGTCTGC CTCACTCAGA TGCTCTCTTG

27851 CTCTCCGAGC TCAGCTACTC CATCAGAAA AACACCACC TCCTTACCTG  
 GAGAGGCTCG AGTCGATGAG GTAGTCTTTT TTGTGGTGGG AGGAATGGAC

27901 CCGGGAACGT ACGAGTGCGT CACCGGCCGC TGCACCACAC CTACCGCCTG  
 GGCCCTTGCA TGCTCACGCA GTGGCCGGCG ACGTGGTGTG GATGGCGGAC

27951 ACCGTAAACC AGACTTTTTTC CGGACAGACC TCAATAACTC TGTTTACCAG  
 TGGCATTG TCTGAAAAAG GCCTGTCTGG AGTTATTGAG ACAAATGGTC

FIG. 10A-35



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|       |                          |                          |                           |                          |                           |
|-------|--------------------------|--------------------------|---------------------------|--------------------------|---------------------------|
| 28001 | AACAGGAGGT<br>TTGTCCTCCA | GAGCTTAGAA<br>CTCGAATCTT | AACCCTTAGG<br>TTGGGAATCC  | GTATTAGGCC<br>CATAATCCGG | AAAGGCGCAG<br>TTTCCGCGTC  |
| 28051 | CTACTGTGGG<br>GATGACACCC | GTTTATGAAC<br>CAAATACTTG | AATTC AAGCA<br>TTAAGTTCGT | ACTCTACGGG<br>TGAGATGCC  | CTATTCTAAT<br>GATAAGATTA  |
| 28101 | TCAGGTTTCT<br>AGTCCAAAGA | CTAGAATCGG<br>GATCTTAGCC | GGTTGGGGTT<br>CCAACCCCAA  | ATTCTCTGTC<br>TAAGAGACAG | TTGTGATTCT<br>AACACTAAGA  |
| 28151 | CTTTATTCTT<br>GAAATAAGAA | ATACTAACGC<br>TATGATTGCG | TTCTCTGCCT<br>AAGAGACGGA  | AAGGCTCGCC<br>TTCCGAGCGG | GCCTGCTGTG<br>CGGACGACAC  |
| 28201 | TGCACATTTG<br>ACGTGTAAAC | CATTTATTGT<br>GTAAATAACA | CAGCTTTTTA<br>GTCGAAAAAT  | AACGCTGGGG<br>TTGCGACCCC | TCGCCACCCA<br>AGCGGTGGGT  |
| 28251 | AGATGATTAG<br>TCTACTAATC | GTACATAATC<br>CATGTATTAG | CTAGGTTTAC<br>GATCCAAATG  | TCACCCTTGC<br>AGTGGGAACG | GTCAGCCCAC<br>CAGTCGGGTG  |
| 28301 | GGTACCACCC<br>CCATGGTGGG | AAAAGGTGGA<br>TTTTCCACCT | TTTTAAGGAG<br>AAAATTCCTC  | CCAGCCTGTA<br>GGTCGGACAT | ATGTTACATT<br>TACAATGTAA  |
| 28351 | CGCAGCTGAA<br>GCGTCGACTT | GCTAATGAGT<br>CGATTACTCA | GCACCACTCT<br>CGTGGTGAGA  | TATAAAATGC<br>ATATTTTACG | ACCACAGAAC<br>TGGTGTCTTG  |
| 28401 | ATGAAAAGCT<br>TACTTTTCGA | GCTTATTCGC<br>CGAATAAGCG | CACAAAAACA<br>GTGTTTTTGT  | AAATTGGCAA<br>TTTAACCGTT | GTATGCTGTT<br>CATACGACAA  |
| 28451 | TATGCTATTT<br>ATACGATAAA | GGCAGCCAGG<br>CCGTCGGTCC | TGACACTACA<br>ACTGTGATGT  | GAGTATAATG<br>CTCATATTAC | TTACAGTTTT<br>AATGTCAAAA  |
| 28501 | CCAGGGTAAA<br>GGTCCCATTT | AGTCATAAAA<br>TCAGTATTTT | CTTTTATGTA<br>GAAAATACAT  | TACTTTTCCA<br>ATGAAAAGGT | TTTTATGAAA<br>AAAATACTTT  |
| 28551 | TGTGCGACAT<br>ACACGCTGTA | TACCATGTAC<br>ATGGTACATG | ATGAGCAAAC<br>TACTCGTTTG  | AGTATAAGTT<br>TCATATTCAA | GTGGCCCCCA<br>CACCGGGGGT  |
| 28601 | CAAAATTGTG<br>GTTTTAACAC | TGGAAAACAC<br>ACCTTTTGTG | TGGCACTTTC<br>ACCGTGAAAG  | TGCTGCACTG<br>ACGACGTGAC | CTATGCTAAT<br>GATACGATTA  |
| 28651 | TACAGTGCTC<br>ATGTCACGAG | GCTTTGGTCT<br>CGAAACCAGA | GTACCCTACT<br>CATGGGATGA  | CTATATTAAA<br>GATATAATTT | TACAAAAGCA<br>ATGTTTTTCGT |
| 28701 | GACGCAGCTT<br>CTGCGTCGAA | TATTGAGGAA<br>ATAACTCCTT | AAGAAAATGC<br>TTCTTTTACG  | CTTAATTTAC<br>GAATTAAATG | TAAGTTACAA<br>ATCAATGTT   |
| 28751 | AGCTAATGTC<br>TCGATTACAG | ACCACTAACT<br>TGGTGATTGA | GCTTTACTCG<br>CGAAATGAGC  | CTGCTTGCAA<br>GACGAACGTT | AACAAATTC A<br>TTGTTTAAGT |

FIG. 10A-36



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28801 AAAAGTTAGC ATTATAATTA GAATAGGATT TAAACCCCCC GGTCATTTCC  
 TTTTCAATCG TAATATTAAT CTTATCCTAA ATTTGGGGGG CCAGTAAAGG

28851 TGCTCAATAC CATTCCCCTG AACCAATTGAC TCTATGTGGG ATATGCTCCA  
 ACGAGTTATG GTAAGGGGAC TTGTTAACTG AGATACACCC TATACGAGGT

28901 GCGCTACAAC CTTGAAGTCA GGCTTCCTGG ATGTCAGCAT CTGACTTTGG  
 CGCGATGTTG GAACTTCAGT CCGAAGGACC TACAGTCGTA GACTGAAACC

28951 CCAGCACCTG TCCCGCGGAT TTGTTCCAGT CCAACTACAG CGACCCACCC  
 GGTCGTGGAC AGGGCGCCTA AACCAAGGTCA GGTTCATGTC GCTGGGTGGG

29001 TAACAGAGAT GACCAACACA ACCAACGCGG CCGCCGCTAC CGGACTTACA  
 ATTGTCTCTA CTGGTTGTGT TGGTTGCGCC GGCGGCGATG GCCTGAATGT

29051 TCTACCACAA ATACACCCA AGTTTCTGCC TTTGTCAATA ACTGGGATAA  
 AGATGGTGTT TATGTGGGGT TCAAAGACGG AAACAGTTAT TGACCCTATT

29101 CTTGGGCATG TGGTGGTTCT CCATAGCGCT TATGTTTGTA TGCCTTATTA  
 GAACCCGTAC ACCACCAAGA GGTATCGCGA ATACAAACAT ACGGAATAAT

29151 TTATGTGGCT CATCTGCTGC CTAAAGCGCA AACGCGCCCG ACCACCCATC  
 AATACACCGA GTAGACGACG GATTTGCGGT TTGCGCGGGC TGGTGGGTAG

29201 TATAGTCCCA TCATTGTGCT ACACCCAAAC AATGATGGAA TCCATAGATT  
 ATATCAGGGT AGTAACACGA TGTGGGTTTG TTACTIONTTT AGGTATCTAA

29251 GGACGGACTG AAACACATGT TCTTTTCTCT TACAGTATGA TTAAATGAGA  
 CCTGCCTGAC TTTGTGTACA AGAAAAGAGA ATGTCATACT AATTTACTCT

29301 CATGATTCCCT CGAGTTTTTA TATTACTGAC CCTTGTTGCG CTTTTTTGTG  
 GTACTAAGGA GCTCAAAAAT ATAATGACTG GGAACAACGC GAAAAAACAC

29351 CGTGCTCCAC ATTGGCTGCG GTTTCTCACA TCGAAGTAGA CTGCATTCCA  
 GCACGAGGTG TAACCGACGC CAAAGAGTGT AGCTTCATCT GACGTAAGGT

29401 GCCTTCACAG TCTATTTGCT TTACGGATTT GTCACCCTCA CGCTCATCTG  
 CGGAAGTGTC AGATAAACGA AATGCCTAAA CAGTGGGAGT GCGAGTAGAC

29451 CAGCCTCATC ACTGTGGTCA TCGCCTTTAT CCAGTGCATT GACTGGGTCT  
 GTCGGAGTAG TGACACCAGT AGCGGAAATA GGTCACGTAA CTGACCCAGA

29501 GTGTGCGCTT TGCATATCTC AGACACCATC CCCAGTACAG GGACAGGACT  
 CACACGCGAA ACGTATAGAG TCTGTGGTAG GGGTCATGTC CCTGTCCTGA

29551 ATAGCTGAGC TTCTTAGAAT TCTTTAATTA TGAAATTTAC TGTGACTTTT  
 TATCGACTCG AAGAATCTTA AGAAATTAAT ACTTTAAATG AACTGAAAA

FIG. 10A-37



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|       |                           |                           |                           |                          |                           |
|-------|---------------------------|---------------------------|---------------------------|--------------------------|---------------------------|
| 29601 | CTGCTGATTA<br>GACGACTAAT  | TTTGCACCCT<br>AAACGTGGGA  | ATCTGCGTTT<br>TAGACGCAA   | TGTTCCCCGA<br>ACAAGGGGCT | CCTCCAAGCC<br>GGAGGTTCGG  |
| 29651 | TCAAAGACAT<br>AGTTTCTGTA  | ATATCATGCA<br>TATAGTACGT  | GATTCACCTCG<br>CTAAGTGAGC | TATATGGAAT<br>ATATACCTTA | ATTCCAAGTT<br>TAAGGTTCAA  |
| 29701 | GCTACAATGA<br>CGATGTTACT  | AAAAAGCGAT<br>TTTTTTCGCTA | CTTTCGGAAG<br>GAAAGGCTTC  | CCTGGTTATA<br>GGACCAATAT | TGCAATCATC<br>ACGTTAGTAG  |
| 29751 | TCTGTTATGG<br>AGACAATACC  | TGTTCTGCAG<br>ACAAGACGTC  | TACCATCTTA<br>ATGGTAGAAT  | GCCCTAGCTA<br>CGGGATCGAT | TATATCCCTA<br>ATATAGGGAT  |
| 29801 | CCTTGACATT<br>GGAACGTAA   | GGCTGGAACG<br>CCGACCTTGC  | CAATAGATGC<br>GTTATCTACG  | CATGAACCAC<br>GTACTTGGTG | CCAAC TTTCC<br>GGTTGAAAGG |
| 29851 | CCGCGCCCGC<br>GGCGCGGGCG  | TATGCTTCCA<br>ATACGAAGGT  | CTGCAACAAG<br>GACGTTGTTC  | TTGTTGCCGG<br>ACAACGGCC  | CGGCTTTGTC<br>GCCGAAACAG  |
| 29901 | CCAGCCAATC<br>GGTCGGTTAG  | AGCCTCGCCC<br>TCGGAGCGGG  | ACCTTCTCCC<br>TGGAAGAGGG  | ACCCCCTG<br>TGGGGGTGAC   | AAATCAGCTA<br>TTTAGTCGAT  |
| 29951 | CTTTAATCTA<br>GAAATTAGAT  | ACAGGAGGAG<br>TGTCCTCCTC  | ATGACTGACA<br>TACTGACTGT  | CCCTAGATCT<br>GGGATCTAGA | AGAAATGGAC<br>TCTTTACCTG  |
| 30001 | GGAATTATTA<br>CCTTAATAAT  | CAGAGCAGCG<br>GTCTCGTCGC  | CCTGCTAGAA<br>GGACGATCTT  | AGACGCAGGG<br>TCTGCGTCCC | CAGCGGCCGA<br>GTCGCCGGCT  |
| 30051 | GCAACAGCGC<br>CGTTGTCGCG  | ATGAATCAAG<br>TACTTAGTTC  | AGCTCCAAGA<br>TCGAGGTTCT  | CATGGTTAAC<br>GTACCAATTG | TTGCACCAGT<br>AACGTGGTCA  |
| 30101 | GCAAAAGGGG<br>CGTTTTCCCC  | TATCTTTTGT<br>ATAGAAAACA  | CTCGTAAAGC<br>GAGCATTTTCG | AGGCCAAAGT<br>TCCGGTTTCA | CACCTACGAC<br>GTGGATGCTG  |
| 30151 | AGTAATACCA<br>TCATTATGGT  | CCGGACACCG<br>GGCCTGTGGC  | CCTTAGCTAC<br>GGAATCGATG  | AAGTTGCCAA<br>TTCAACGGTT | CCAAGCGTCA<br>GGTTCGCAGT  |
| 30201 | GAAATTGGTG<br>CTTTAACCAC  | GTCATGGTGG<br>CAGTACCACC  | GAGAAAAGCC<br>CTCTTTTCGG  | CATTACCATA<br>GTAATGGTAT | ACTCAGCACT<br>TGAGTCGTGA  |
| 30251 | CGGTAGAAAC<br>GCCATCTTTG  | CGAAGGCTGC<br>GCTTCCGACG  | ATTCACCTCAC<br>TAAGTGAGTG | CTTGTCAAGG<br>GAACAGTTCC | ACCTGAGGAT<br>TGGACTCCTA  |
| 30301 | CTCTGCACCC<br>GAGACGTGGG  | TTATTAAGAC<br>AATAATTCTG  | CCTGTGCGGT<br>GGACACGCCA  | CTCAAAGATC<br>GAGTTTCTAG | TTATTCCCTT<br>AATAAGGGAA  |
| 30351 | TAAC TAATAA<br>ATTGATTATT | AAAAAAATAA<br>TTTTTTTATT  | TAAAGCATCA<br>ATTCGTAGT   | CTTACTTAAA<br>GAATGAATTT | ATCAGTTAGC<br>TAGTCAATCG  |

FIG. 10A-38



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30401 AAATTTCTGT CCAGTTTATT CAGCAGCACC TCCTTGCCCT CCTCCCAGCT  
 TTTAAAGACA GGTCAAATAA GTCGTCTGG AGGAACGGGA GGAGGGTCGA

30451 CTGGTATTGC AGCTTCCTCC TGGCTGCAAA CTTTCTCCAC AATCTAAATG  
 GACCATAACG TCGAAGGAGG ACCGACGTTT GAAAGAGGTG TTAGATTTAC

30501 GAATGTCAGT TTCCTCCTGT TCCTGTCCAT CCGCACCCAC TATCTTCATG  
 CTTACAGTCA AAGGAGGACA AGGACAGGTA GGCCTGGGTG ATAGAAGTAC

30551 TTGTTGCAGA TGAAGCGCGC AAGACCGTCT GAAGATACCT TCAACCCCGT  
 AACAACTGTCT ACTTCGCGCG TTCTGGCAGA CTTCTATGGA AGTTGGGGCA

30601 GTATCCATAT GACACGGAAA CCGGTCTTCC AACTGTGCCT TTTCTTACTC  
 CATAGGTATA CTGTGCCTTT GGCCAGGAGG TTGACACGGA AAAGAATGAG

30651 CTCCCTTTGT ATCCCCAAT GGGTTTCAAG AGAGTCCCCC TGGGGTACTC  
 GAGGGAAACA TAGGGGGTTA CCCAAAGTTC TCTCAGGGGG ACCCCATGAG

30701 TCTTTGCGCC TATCCGAACC TCTAGTTACC TCCAATGGCA TGCTTGCGCT  
 AGAAACGCGG ATAGGCTTGG AGATCAATGG AGGTTACCGT ACGAACGCGA

30751 CAAAATGGGC AACGGCCTCT CTCTGGACGA GGCCGGCAAC CTTACCTCCC  
 GTTTTACCCG TTGCCGGAGA GAGACCTGCT CCGGCCGTTG GAATGGAGGG

30801 AAAATGTAAC CACTGTGAGC CCACCTCTCA AAAAAACCAA GTCAAACATA  
 TTTTACATTG GTGACACTCG GGTGGAGAGT TTTTTTGGTT CAGTTTGTAT

30851 AACCTGGAAA TATCTGCACC CCTCACAGTT ACCTCAGAAG CCCTAACTGT  
 TTGGACCTTT ATAGACGTGG GGAGTGTCAA TGGAGTCTTC GGGATTGACA

30901 GGCTGCCGCC GCACCTCTAA TGGTCGCGGG CAACACACTC ACCATGCAAT  
 CCGACGGCGG CGTGGAGATT ACCAGCGCCC GTTGTGTGAG TGGTACGTTA

30951 CACAGGCCCC GCTAACCGTG CACGACTCCA AACTTAGCAT TGCCACCCAA  
 GTGTCCGGGG CGATTGGCAC GTGCTGAGGT TTGAATCGTA ACGGTGGGTT

31001 GGACCCCTCA CAGTGTCAGA AGGAAAGCTA GCCCTGCAAA CATCAGGCC  
 CCTGGGGAGT GTCACAGTCT TCCTTTCGAT CGGGACGTTT GTAGTCCGGG

31051 CCTCACCACC ACCGATAGCA GTACCCTTAC TATCACTGCC TCACCCCTT  
 GGAGTGGTGG TGGCTATCGT CATGGGAATG ATAGTGACGG AGTGGGGGAA

31101 TAACTACTGC CACTGGTAGC TTGGGCATTG ACTTGAAAGA GCCCATTTAT  
 ATTGATGACG GTGACCATCG AACCCTGTAAC TGAACCTTCT CGGGTAAATA

31151 ACACAAAATG GAAAACCTAGG ACTAAAGTAC GGGGCTCCTT TGCATGTAAC  
 TGTGTTTTAC CTTTTGATCC TGATTTTCATG CCCCAGGAA ACGTACATTG

FIG. 10A-39



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|       |                          |                           |                          |                           |                            |
|-------|--------------------------|---------------------------|--------------------------|---------------------------|----------------------------|
| 31201 | AGACGACCTA<br>TCTGCTGGAT | AACACTTTGA<br>TTGTGAAACT  | CCGTAGCAAC<br>GGCATCGTTG | TGGTCCAGGT<br>ACCAGGTCCA  | GTGACTATTA<br>CACTGATAAT   |
| 31251 | ATAATACTTC<br>TATTATGAAG | CTTGCAAACCT<br>GAACGTTTGA | AAAGTTACTG<br>TTTCAATGAC | GAGCCTTGGG<br>CTCGGAACCC  | TTTTGATTCA<br>AAAACCTAAGT  |
| 31301 | CAAGGCAATA<br>GTTCCGTTAT | TGCAACTTAA<br>ACGTTGAATT  | TGTAGCAGGA<br>ACATCGTCCT | GGACTAAGGA<br>CCTGATTCCT  | TTGATTCTCA<br>AACTAAGAGT   |
| 31351 | AAACAGACGC<br>TTTGTCTGCG | CTTATACTTG<br>GAATATGAAC  | ATGTTAGTTA<br>TACAATCAAT | TCCGTTTGAT<br>AGGCAAACCTA | GCTCAAAACC<br>CGAGTTTTGG   |
| 31401 | AACTAAATCT<br>TTGATTTAGA | AAGACTAGGA<br>TTCTGATCCT  | CAGGGCCCTC<br>GTCGCCGGAG | TTTTTATAAA<br>AAAATATTT   | CTCAGCCCAC<br>GAGTCGGGTG   |
| 31451 | AACTTGGATA<br>TTGAACCTAT | TTAACTACAA<br>AATTGATGTT  | CAAAGGCCTT<br>GTTTCCGGAA | TACTTGTTTA<br>ATGAACAAAT  | CAGCTTCAAA<br>GTCGAAGTTT   |
| 31501 | CAATTCCAAA<br>GTTAAGGTTT | AAGCTTGAGG<br>TTCGAACTCC  | TTAACCTAAG<br>AATTGGATTC | CACTGCCAAG<br>GTGACGGTTC  | GGGTGATGT<br>CCCAACTACA    |
| 31551 | TTGACGCTAC<br>AACTGCGATG | AGCCATAGCC<br>TCGGTATCGG  | ATTAATGCAG<br>TAATTACGTC | GAGATGGGCT<br>CTCTACCCGA  | TGAATTTGGT<br>ACTTAAACCA   |
| 31601 | TCACCTAATG<br>AGTGGATTAC | CACCAAACAC<br>GTGGTTTGTG  | AAATCCCCTC<br>TTTAGGGGAG | AAAACAAAAA<br>TTTTGTTTTT  | TTGGCCATGG<br>AACC GG TACC |
| 31651 | CCTAGAATTT<br>GGATCTTAAA | GATTCAAACA<br>CTAAGTTTGT  | AGGCTATGGT<br>TCCGATACCA | TCCTAAACTA<br>AGGATTTGAT  | GGAAGTGGCC<br>CCTTGACCGG   |
| 31701 | TTAGTTTTGA<br>AATCAAAACT | CAGCACAGGT<br>GTCGTGTCCA  | GCCATTACAG<br>CGGTAATGTC | TAGGAAACAA<br>ATCCTTTGTT  | AAATAATGAT<br>TTTATTACTA   |
| 31751 | AAGCTAACTT<br>TTCGATTGAA | TGTGGACCAC<br>ACACCTGGTG  | ACCAGCTCCA<br>TGGTCGAGGT | TCTCCTAACT<br>AGAGGATTGA  | GTAGACTAAA<br>CATCTGATTT   |
| 31801 | TGCAGAGAAA<br>ACGTCTCTTT | GATGCTAAAC<br>CTACGATTTG  | TCACTTTGGT<br>AGTGAAACCA | CTTAACAAAA<br>GAATTGTTTT  | TGTGGCAGTC<br>ACACCGTCAG   |
| 31851 | AAATACTTGC<br>TTTATGAACG | TACAGTTTCA<br>ATGTCAAAGT  | GTTTTGGCTG<br>CAAACCGAC  | TTAAAGGCAG<br>AATTTCCGTC  | TTTGGCTCCA<br>AAACCGAGGT   |
| 31901 | ATATCTGGAA<br>TATAGACCTT | CAGTTCAAAG<br>GTCAAGTTTC  | TGCTCATCTT<br>ACGAGTAGAA | ATTATAAGAT<br>TAATATTCTA  | TTGACGAAAA<br>AACTGCTTTT   |
| 31951 | TGGAGTGCTA<br>ACCTCACGAT | CTAAACAATT<br>GATTTGTAA   | CCTTCCTGGA<br>GGAAGGACCT | CCCAGAATAT<br>GGGTCTTATA  | TGGAACTTTA<br>ACCTTGAAAT   |

FIG. 10A-40



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|       |                          |                           |                          |                           |                           |
|-------|--------------------------|---------------------------|--------------------------|---------------------------|---------------------------|
| 32001 | GAAATGGAGA<br>CTTTACCTCT | TCTTACTGAA<br>AGAATGACTT  | GGCACAGCCT<br>CCGTGTCGGA | ATACAAACGC<br>TATGTTTGCG  | TGTTGGATTT<br>ACAACCTAAA  |
| 32051 | ATGCCTAACC<br>TACGGATTGG | TATCAGCTTA<br>ATAGTCGAAT  | TCCAAAATCT<br>AGGTTTTAGA | CACGGTAAAA<br>GTGCCATTTT  | CTGCCAAAAG<br>GACGGTTTTC  |
| 32101 | TAACATTGTC<br>ATTGTAACAG | AGTCAAGTTT<br>TCAGTTCAA   | ACTTAAACGG<br>TGAATTTGCC | AGACAAAACCT<br>TCTGTTTTGA | AAACCTGTAA<br>TTTGGACATT  |
| 32151 | CACTAACCAT<br>GTGATTGGTA | TACACTAAAC<br>ATGTGATTTG  | GGTACACAGG<br>CCATGTGTCC | AAACAGGAGA<br>TTTGTCCTCT  | CACAACTCCA<br>GTGTTGAGGT  |
| 32201 | AGTGCATACT<br>TCACGTATGA | CTATGTCATT<br>GATACAGTAA  | TTCATGGGAC<br>AAGTACCCTG | TGGTCTGGCC<br>ACCAGACCGG  | ACAACCTACAT<br>TGTTGATGTA |
| 32251 | TAATGAAATA<br>ATTACTTTAT | TTTGCCACAT<br>AAACGGTGTA  | CCTCTTACAC<br>GGAGAATGTG | TTTTTCATAC<br>AAAAGTATG   | ATTGCCCAAG<br>TAACGGGTTC  |
| 32301 | AATAAAGAAT<br>TTATTTCTTA | CGTTTGTGTT<br>GCAAACACAA  | ATGTTTCAAC<br>TACAAAGTTG | GTGTTTATTT<br>CACAAATAAA  | TTCAATTGCA<br>AAGTTAACGT  |
| 32351 | GAAAATTTCA<br>CTTTTAAAGT | AGTCATTTTT<br>TCAGTAAAAA  | CATTCAGTAG<br>GTAAGTCATC | TATAGCCCA<br>ATATCGGGGT   | CCACCACATA<br>GGTGGTGTAT  |
| 32401 | GCTTATACAG<br>CGAATATGTC | ATCACCGTAC<br>TAGTGGCATG  | CTTAATCAA<br>GAATTAGTTT  | CTCACAGAAC<br>GAGTGTCTTG  | CCTAGTATTC<br>GGATCATAAG  |
| 32451 | AACCTGCCAC<br>TTGGACGGTG | CTCCCTCCCA<br>GAGGGAGGGT  | ACACACAGAG<br>TGTGTGTCTC | TACACAGTCC<br>ATGTGTCAGG  | TTTCTCCCCG<br>AAAGAGGGGC  |
| 32501 | GCTGGCCTTA<br>CGACCGGAAT | AAAAGCATCA<br>TTTTTCGTAGT | TATCATGGGT<br>ATAGTACCCA | AACAGACATA<br>TTGTCTGTAT  | TTCTTAGGTG<br>AAGAATCCAC  |
| 32551 | TTATATTCCA<br>AATATAAGGT | CACGGTTTTCC<br>GTGCCAAAGG | TGTCGAGCCA<br>ACAGCTCGGT | AACGCTCATC<br>TTGCGAGTAG  | AGTGATATTA<br>TCACTATAAT  |
| 32601 | ATAAACTCCC<br>TATTTGAGGG | CGGGCAGCTC<br>GCCCGTCGAG  | ACTTAAGTTC<br>TGAATTCAAG | ATGTCGCTGT<br>TACAGCGACA  | CCAGCTGCTG<br>GGTCGACGAC  |
| 32651 | AGCCACAGGC<br>TCGGTGTCCG | TGCTGTCCAA<br>ACGACAGGTT  | CTTGCGGTTG<br>GAACGCCAAC | CTTAACGGGC<br>GAATTGCCCG  | GGCGAAGGAG<br>CCGCTTCCTC  |
| 32701 | AAGTCCACGC<br>TTCAGGTGCG | CTACATGGGG<br>GATGTACCCC  | GTAGAGTCAT<br>CATCTCAGTA | AATCGTGCA<br>TTAGCACGTA   | CAGGATAGGG<br>GTCCTATCCC  |
| 32751 | CGGTGGTGCT<br>GCCACCACGA | GCAGCAGCGC<br>CGTCGTCGCG  | GCGAATAAAC<br>CGCTTATTTG | TGCTGCCGCC<br>ACGACGGCGG  | GCCGCTCCGT<br>CGGCGAGGCA  |

FIG. 10A-41



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|       |                           |                          |                           |                          |                          |
|-------|---------------------------|--------------------------|---------------------------|--------------------------|--------------------------|
| 32801 | CCTGCAGGAA<br>GGACGTCCTT  | TACAACATGG<br>ATGTTGTACC | CAGTGGTCTC<br>GTCACCAGAG  | CTCAGCGATG<br>GAGTCGCTAC | ATTCGCACCG<br>TAAGCGTGGC |
| 32851 | CCCGCAGCAT<br>GGCCGTCGTA  | AAGGCGCCTT<br>TTCCGCGGAA | GTCCTCCGGG<br>CAGGAGGCCC  | CACAGCAGCG<br>GTGTCGTCGC | CACCCTGATC<br>GTGGGACTAG |
| 32901 | TCACTTAAAT<br>AGTGAATTTA  | CAGCACAGTA<br>GTCGTGTCAT | ACTGCAGCAC<br>TGACGTCGTG  | AGCACCACAA<br>TCGTGGTGTT | TATTGTTCAA<br>ATAACAAGTT |
| 32951 | AATCCCACAG<br>TTAGGGTGTC  | TGCAAGGCGC<br>ACGTTCCGCG | TGTATCCAAA<br>ACATAGGTTT  | GCTCATGGCG<br>CGAGTACCGC | GGGACCACAG<br>CCCTGGTGTC |
| 33001 | AACCCACGTG<br>TTGGGTGCAC  | GCCATCATA<br>CGGTAGTATG  | CACAAGCGCA<br>GTGTTCCGCGT | GGTAGATTAA<br>CCATCTAATT | GTGGCGACCC<br>CACCGCTGGG |
| 33051 | CTCATAAACA<br>GAGTATTTGT  | CGCTGGACAT<br>GCGACCTGTA | AAACATTACC<br>TTTGTAATGG  | TCTTTTGGCA<br>AGAAAACCGT | TGTTGTAATT<br>ACAACATTAA |
| 33101 | CACCACCTCC<br>GTGGTGGAGG  | CGGTACCATA<br>GCCATGGTAT | TAAACCTCTG<br>ATTTGGAGAC  | ATTAAACATG<br>TAATTTGTAC | GCGCCATCCA<br>CGCGGTAGGT |
| 33151 | CCACCATCCT<br>GGTGGTAGGA  | AAACCAGCTG<br>TTTGGTCGAC | GCCAAAACCT<br>CGGTTTTGGA  | GCCCGCCGGC<br>CGGGCGGCCG | TATAACTGTC<br>ATATGTGACG |
| 33201 | AGGGAACCGG<br>TCCCTTGGCC  | GACTGGAACA<br>CTGACCTTGT | ATGACAGTGG<br>TACTGTCACC  | AGAGCCCAGG<br>TCTCGGGTCC | ACTCGTAACC<br>TGAGCATTGG |
| 33251 | ATGGATCATC<br>TACCTAGTAG  | ATGCTCGTCA<br>TACGAGCAGT | TGATATCAAT<br>ACTATAGTTA  | GTTGGCACAA<br>CAACCGTGTT | CACAGGCACA<br>GTGTCCTGTG |
| 33301 | CGTGCATACA<br>GCACGTATGT  | CTTCCTCAGG<br>GAAGGAGTCC | ATTACAAGCT<br>TAATGTTCGA  | CCTCCCGCGT<br>GGAGGGCGCA | TAGAACCATA<br>ATCTTGGTAT |
| 33351 | TCCCAGGGAA<br>AGGGTCCCTT  | CAACCCATTC<br>GTTGGGTAAG | CTGAATCAGC<br>GACTTAGTCG  | GTAAATCCCA<br>CATTTAGGGT | CACTGCAGGG<br>GTGACGTCCC |
| 33401 | AAGACCTCGC<br>TTCTGGAGCG  | ACGTAACTCA<br>TGCATTGAGT | CGTTGTGCAT<br>GCAACACGTA  | TGTCAAAGTG<br>ACAGTTTCAC | TTACATTCGG<br>AATGTAAGCC |
| 33451 | GCAGCAGCGG<br>CGTCGTCGCC  | ATGATCCTCC<br>TACTAGGAGG | AGTATGGTAG<br>TCATACCATC  | CGCGGGTTTC<br>GCGCCCAAAG | TGTCTCAAAA<br>ACAGAGTTTT |
| 33501 | GGAGGTAGAC<br>CCTCCATCTG  | GATCCCTACT<br>CTAGGGATGA | GTACGGAGTG<br>CATGCCTCAC  | CGCCGAGACA<br>GCGGCTCTGT | ACCGAGATCG<br>TGGCTCTAGC |
| 33551 | TGTTGGTTCGT<br>ACAACCAGCA | AGTGTCATGC<br>TCACAGTACG | CAAATGGAAC<br>GTTTACCTTG  | GCCGGACGTA<br>CGGCTGCAT  | GTCATATTTT<br>CAGTATAAAG |

FIG. 10A-42



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|       |                           |                           |                          |                          |                          |
|-------|---------------------------|---------------------------|--------------------------|--------------------------|--------------------------|
| 33601 | CTGAAGCAAA<br>GACTTCGTTT  | ACCAGGTGCG<br>TGGTCCACGC  | GGCGTGACAA<br>CCGCACTGTT | ACAGATCTGC<br>TGTCTAGACG | GTCTCCGGTC<br>CAGAGGCCAG |
| 33651 | TCGCCGCTTA<br>AGCGGCCGAAT | GATCGCTCTG<br>CTAGCCGAGAC | TGTAGTAGTT<br>ACATCATCAA | GTAGTATATC<br>CATCATATAG | CACTCTCTCA<br>GTGAGAGAGT |
| 33701 | AAGCATCCAG<br>TTCGTAGGTC  | GCGCCCCCTG<br>CGCGGGGGAC  | GCTTCGGGTT<br>CGAAGCCCAA | CTATGTAAAC<br>GATACATTTG | TCCTTCATGC<br>AGGAAGTACG |
| 33751 | GCCGCTGCCC<br>CGGCGACGGG  | TGATAACATC<br>ACTATTGTAG  | CACCACCGCA<br>GTGGTGGCGT | GAATAAGCCA<br>CTTATTCGGT | CACCCAGCCA<br>GTGGGTCCGT |
| 33801 | ACCTACACAT<br>TGGATGTGTA  | TCGTTCTGCG<br>AGCAAGACGC  | AGTCACACAC<br>TCAGTGTGTG | GGGAGGAGCG<br>CCCTCCTCGC | GGAAGAGCTG<br>CCTTCTCGAC |
| 33851 | GAAGAACCAT<br>CTTCTTGGTA  | GTTTTTTTTT<br>CAAAAAAAAA  | TTATTCCAAA<br>AATAAGGTTT | AGATTATCCA<br>TCTAATAGGT | AAACCTCAAA<br>TTTGGAGTTT |
| 33901 | ATGAAGATCT<br>TACTTCTAGA  | ATTAAGTGAA<br>TAATTCACTT  | GCGCTCCCC<br>GCGCGAGGGG  | TCCGGTGGCG<br>AGGCCACCGC | TGGTCAAAC<br>ACCAGTTTGA  |
| 33951 | CTACAGCCAA<br>GATGTCGGTT  | AGAACAGATA<br>TCTTGTCCTAT | ATGGCATTG<br>TACCGTAAAC  | TAAGATGTTG<br>ATTCTACAAC | CACAATGGCT<br>GTGTTACCGA |
| 34001 | TCCAAAAGGC<br>AGGTTTTCCG  | AAACGGCCCT<br>TTTGCCGGGA  | CACGTCCAAG<br>GTGCAGGTTT | TGGACGTAAA<br>ACCTGCATTT | GGCTAAACCC<br>CCGATTTGGG |
| 34051 | TTCAGGGTGA<br>AAGTCCCCT   | ATCTCCTCTA<br>TAGAGGAGAT  | TAAACATTCC<br>ATTTGTAAGG | AGCACCTTCA<br>TCGTGGAAGT | ACCATGCCCA<br>TGGTACGGGT |
| 34101 | AATAATTCTC<br>TTATTAAGAG  | ATCTCGCCAC<br>TAGAGCGGTG  | CTTCTCAATA<br>GAAGAGTTAT | TATCTCTAAG<br>ATAGAGATTC | CAAATCCCGA<br>GTTTAGGGCT |
| 34151 | ATATTAAGTC<br>TATAATTCAG  | CGGCCATTGT<br>GCCGGTAACA  | AAAAATCTGC<br>TTTTTAGACG | TCCAGAGCGC<br>AGGTCTCGCG | CCTCCACCTT<br>GGAGGTGGAA |
| 34201 | CAGCCTCAAG<br>GTCGGAGTTC  | CAGCGAATCA<br>GTCGCTTAGT  | TGATTGCAAA<br>ACTAACGTTT | AATTCAGGTT<br>TTAAGTCCAA | CCTCACAGAC<br>GGAGTGTCTG |
| 34251 | CTGTATAAGA<br>GACATATTCT  | TTCAAAAGCG<br>AAGTTTTTCGC | GAACATTAAC<br>CTTGTAATTG | AAAAATACCG<br>TTTTTATGGC | CGATCCCGTA<br>GCTAGGGCAT |
| 34301 | GGTCCCTTCG<br>CCAGGGAAGC  | CAGGGCCAGC<br>GTCCCGGTCG  | TGAACATAAT<br>ACTTGTATTA | CGTGCAGGTC<br>GCACGTCCAG | TGCACGGACC<br>ACGTGCCTGG |
| 34351 | AGCGCGGCCA<br>TCGCGCCGGT  | CTTCCCCGCC<br>GAAGGGGCGG  | AGGAACCATG<br>TCCTTGGTAC | ACAAAAGAAC<br>TGTTTTCTTG | CCACACTGAT<br>GGTGTGACTA |

FIG. 10A-43



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|       |             |             |             |            |             |
|-------|-------------|-------------|-------------|------------|-------------|
| 34401 | TATGACACGC  | ATACTCGGAG  | CTATGCTAAC  | CAGCGTAGCC | CCGATGTAAG  |
|       | ATACTGTGCG  | TATGAGCCTC  | GATACGATTG  | GTCGCATCGG | GGCTACATTC  |
| 34451 | CTTGTTGCAT  | GGGCGGCGAT  | ATAAAAATGCA | AGGTGCTGCT | CAAAAAATCA  |
|       | GAACAACGTA  | CCCGCCGCTA  | TATTTTACGT  | TCCACGACGA | GTTTTTTAGT  |
| 34501 | GGCAAAGCCT  | CGCGCAAAAA  | AGAAAGCACA  | TCGTAGTCAT | GCTCATGCAG  |
|       | CCGTTTCGGA  | GCGCGTTTTT  | TCTTTCGTGT  | AGCATCAGTA | CGAGTACGTC  |
| 34551 | ATAAAGGCAG  | GTAAGCTCCG  | GAACCACCAC  | AGAAAAAGAC | ACCATTTTTTC |
|       | TATTTCCGTC  | CATTCGAGGC  | CTTGGTGGTG  | TCTTTTTCTG | TGGTAAAAAG  |
| 34601 | TCTCAAACAT  | GTCTGCGGGT  | TTCTGCATAA  | ACACAAAATA | AAATAACAAA  |
|       | AGAGTTTGTA  | CAGACGCCCA  | AAGACGTATT  | TGTGTTTTAT | TTTATTGTTT  |
| 34651 | AAAACATTTA  | AACATTAGAA  | GCCTGTCTTA  | CAACAGGAAA | AACAACCCTT  |
|       | TTTTGTAAAT  | TTGTAATCTT  | CGGACAGAAT  | GTTGTCCTTT | TTGTTGGGAA  |
| 34701 | ATAAGCATAA  | GACGGACTAC  | GGCCATGCCG  | GCGTGACCGT | AAAAAAACTG  |
|       | TATTCGTATT  | CTGCCTGATG  | CCGGTACGGC  | CGCACTGGCA | TTTTTTTTGAC |
| 34751 | GTCACCGTGA  | TTAAAAAGCA  | CCACCGACAG  | CTCCTCGGTC | ATGTCCGGAG  |
|       | CAGTGGCACT  | AATTTTTTCGT | GGTGGCTGTC  | GAGGAGCCAG | TACAGGCCTC  |
| 34801 | TCATAATGTA  | AGACTCGGTA  | AACACATCAG  | GTTGATTCAC | ATCGGTCAGT  |
|       | AGTATTACAT  | TCTGAGCCAT  | TTGTGTAGTC  | CAACTAAGTG | TAGCCAGTCA  |
| 34851 | GCTAAAAAGC  | GACCGAAATA  | GCCCGGGGGA  | ATACATACCC | GCAGGCGTAG  |
|       | CGATTTTTTCG | CTGGCTTTAT  | CGGGCCCCCT  | TATGTATGGG | CGTCCGCATC  |
| 34901 | AGACAACATT  | ACAGCCCCCA  | TAGGAGGTAT  | AACAAAATTA | ATAGGAGAGA  |
|       | TCTGTTGTAA  | TGTCGGGGGT  | ATCCTCCATA  | TTGTTTTAAT | TATCCTCTCT  |
| 34951 | AAAACACATA  | AACACCTGAA  | AAACCCTCCT  | GCCTAGGCAA | AATAGCACCC  |
|       | TTTTGTGTAT  | TTGTGGACTT  | TTTGGGAGGA  | CGGATCCGTT | TTATCGTGGG  |
| 35001 | TCCCGCTCCA  | GAACAACATA  | CAGCGCTTCC  | ACAGCGGCAG | CCATAACAGT  |
|       | AGGGCGAGGT  | CTTGTTGTAT  | GTCGCGAAGG  | TGTCGCCGTC | GGTATTGTCA  |
| 35051 | CAGCCTTACC  | AGTAAAAAAG  | AAAACCTATT  | AAAAAAACAC | CACTCGACAC  |
|       | GTCGGAATGG  | TCATTTTTTC  | TTTTGGATAA  | TTTTTTTGTG | GTGAGCTGTG  |
| 35101 | GGCACCAGCT  | CAATCAGTCA  | CAGTGTAATA  | AAGGGCCAAG | TGCAGAGCGA  |
|       | CCGTGGTCGA  | GTTAGTCAGT  | GTCACATTTT  | TTCCCGGTTC | ACGTCTCGCT  |
| 35151 | GTATATATAG  | GACTAAAAAA  | TGACGTAACG  | GTTAAAGTCC | ACAAAAAACA  |
|       | CATATATATC  | CTGATTTTTT  | ACTGCATTGC  | CAATTCAGG  | TGTTTTTTGT  |

FIG. 10A-44



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35201 CCCAGAAAAC CGCACGCGAA CCTACGCCCA GAAACGAAAG CCAAAAAACC  
 GGGTCTTTTG GCGTGCGCTT GGATGCGGGT CTTTGCTTTC GGTTTTTTGG

35251 CACAACCTCC TCAAATCGTC ACTTCCGTTT TCCCACGTTA CGTCACTTCC  
 GTGTTGAAGG AGTTTAGCAG TGAAGGCAAA AGGGTGCAAT GCAGTGAAGG

35301 CATTTTAAGA AACTACAAT TCCCAACACA TACAAGTTAC TCCGCCCTAA  
 GTAAAATTCT TTTGATGTTA AGGGTTGTGT ATGTTCAATG AGGCGGGATT

35351 AACCTACGTC ACCCGCCCCG TTCCCACGCC CCGCGCCACG TCACAAACTC  
 TTGGATGCAG TGGGCGGGGC AAGGGTGCCG GCGCGGGTGC AGTGTTTGAG

35401 CACCCCCTCA TTATCATATT GGCTTCAATC CAAAATAAGG TATATTATTG  
 GTGGGGGAGT AATAGTATAA CCGAAGTTAG GTTTTATTCC ATATAATAAC

PacI  
~~~~~

35451 ATGATGTTAA TTAAGAATTC GGATCTGCGA CGCGAGGCTG GATGGCCTTC  
 TACTACAATT AATTCTTAAG CCTAGACGCT GCGCTCCGAC CTACCGGAAG

35501 CCCATTATGA TTCTTCTCGC TTCCGGCGGC ATCGGGATGC CCGCGTTGCA  
 GGGTAATACT AAGAAGAGCG AAGGCCGCCG TAGCCCTACG GCGCAACGT

35551 GGCCATGCTG TCCAGGCAGG TAGATGACGA CCATCAGGGA CAGCTTCAAG  
 CCGGTACGAC AGGTCCGTCC ATCTACTGCT GGTAGTCCCT GTCGAAGTTC

35601 GCCAGCAAAA GGCCAGGAAC CGTAAAAAGG CCGCGTTGCT GCGGTTTTTC  
 CGGTCGTTTT CCGGTCCTTG GCATTTTTTC GCGCAACGA CCGCAAAAAG

35651 CATAGGCTCC GCCCCCTGA CGAGCATCAC AAAAATCGAC GCTCAAGTCA  
 GTATCCGAGG CGGGGGGACT GCTCGTAGTG TTTTLAGCTG CGAGTTCAGT

35701 GAGGTGGCGA AACCCGACAG GACTATAAAG ATACCAGGCG TTTCCCCCTG  
 CTCCACCGCT TTGGGCTGTC CTGATATTTT TATGGTCCGC AAAGGGGGAC

35751 GAAGCTCCCT CGTGCGCTCT CCTGTTCCGA CCCTGCCGCT TACCGGATAC  
 CTTGAGGGA GCACGCGAGA GGACAAGGCT GGGACGGCGA ATGGCCTATG

35801 CTGTCCGCCT TTCTCCCTTC GGAAGCGTG GCGCTTTCCT ATAGCTCACG  
 GACAGGCGGA AAGAGGGAAG CCCTTCGCAC CCGGAAAGAG TATCGAGTGC

35851 CTGTAGGTAT CTCAGTTCGG TGTAGGTCGT TCGCTCCAAG CTGGGCTGTG  
 GACATCCATA GAGTCAAGCC ACATCCAGCA AGCGAGGTTC GACCCGACAC

35901 TGCACGAACC CCCCCTTCAG CCCGACCGCT GCGCCTTATC CGGTAACAT  
 ACGTGCTTGG GGGGCAAGTC GGGCTGGCGA CCGGGAATAG GCCATTGATA

FIG. 10A-45



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35951	CGTCTTGAGT GCAGAACTCA	CCAACCCGGT GGTTGGGCCA	AAGACACGAC TTCTGTGCTG	TTATCGCCAC AATAGCGGTG	TGGCAGCAGC ACCGTCGTCG
36001	CACTGGTAAC GTGACCATTC	AGGATTAGCA TCCTAATCGT	GAGCGAGGTA CTCGCTCCAT	TGTAGGCGGT ACATCCGCCA	GCTACAGAGT CGATGTCTCA
36051	TCTTGAAGTG AGAACTTCAC	GTGGCCTAAC CACCGGATTG	TACGGCTACA ATGCCGATGT	CTAGAAGGAC GATCTTCCTG	AGTATTTGGT TCATAAACCA
36101	ATCTGCGCTC TAGACGCGAG	TGCTGAAGCC ACGACTTCGG	AGTTACCTTC TCAATGGAAG	GGAAAAGAG CCTTTTTCTC	TTGGTAGCTC AACCATCGAG
36151	TTGATCCGGC AACTAGGCCG	AAACAAACCA TTTGTTTGGT	CCGCTGGTAG GGCGACCATC	CGGTGGTTTT GCCACCAAAA	TTTGTTTGCA AAACAAACGT
36201	AGCAGCAGAT TCGTCGTCTA	TACGCGCAGA ATGCGCGTCT	AAAAAAGGAT TTTTTTCCTA	CTCAAGAAGA GAGTTCTTCT	TCCTTTGATC AGGAAACTAG
36251	TTTTCTACGG AAAAGATGCC	GGTCTGACGC CCAGACTGCG	TCAGTGGAAC AGTCACCTTG	GAAACTCAC CTTTTGAGTG	GTTAAGGGAT CAATTCCCTA
36301	TTTGGTCATG AAACCAGTAC	AGATTATCAA TCTAATAGTT	AAAGGATCTT TTTCCTAGAA	CACCTAGATC GTGGATCTAG	CTTTTAAATC GAAAATTTAG
36351	AATCTAAAGT TTAGATTTCA	ATATATGAGT TATATACTCA	AACTTGGTC TTTGAACCAG	TGACAGTTAC ACTGTCAATG	CAATGCTTAA GTTACGAATT
36401	TCAGTGAGGC AGTCACTCCG	ACCTATCTCA TGGATAGAGT	GCGATCTGTC CGCTAGACAG	TATTTTCGTT ATAAAGCAAG	ATCCATAGTT TAGGTATCAA
36451	GCCTGACTCC CGGACTGAGG	CCGTCGTGTA GGCAGCACAT	GATAACTACG CTATTGATGC	ATACGGGAGG TATGCCCTCC	GCTTACCATC CGAATGGTAG
36501	TGGCCCCAGT ACCGGGGTCA	GCTGCAATGA CGACGTTACT	TACCGCGAGA ATGGCGCTCT	CCCACGCTCA GGGTGCGAGT	CCGGCTCCAG GGCCGAGGTC
36551	ATTTATCAGC TAAATAGTCG	AATAAACCAG TTATTTGGTC	CCAGCCGGAA GGTCGGCCTT	GGCCGAGCG CCCGGCTCGC	CAGAAGTGGT GTCTTCACCA
36601	CCTGCAACTT GGACGTTGAA	TATCCGCCTC ATAGGCGGAG	CATCCAGTCT GTAGGTCAGA	ATTAATTGTT TAATTAACAA	GCCGGGAAGC CGGCCCTTCG
36651	TAGAGTAAGT ATCTCATTCA	AGTTCGCCAG TCAAGCGGTC	TTAATAGTTT AATTATCAAA	GCGCAACGTT CGCGTTGCAA	GTTGCCATTG CAACGGTAAC
36701	CTACAGGCAT GATGTCCGTA	CGTGGTGTC GCACCACAGT	CGCTCGTCGT GCGAGCAGCA	TTGGTATGGC AACCATACCG	TTCATTCAGC AAGTAAGTCG

FIG. 10A-46



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36751 TCCGGTCCC AACGATCAAG GCGAGTTACA TGATCCCCCA TGTTGTGCAA  
 AGGCCAAGGG TTGCTAGTTC CGCTCAATGT ACTAGGGGGT ACAACACGTT

36801 AAAAGCGGTT AGCTCCTTCG GTCCTCCGAT CGTTGTCAGA AGTAAGTTGG  
 TTTTCGCCAA TCGAGGAAGC CAGGAGGCTA GCAACAGTCT TCATTCAACC

36851 CCGCAGTGTT ATCACTCATG GTTATGGCAG CACTGCATAA TTCTCTTACT  
 GGCGTCACAA TAGTGAGTAC CAATACCGTC GTGACGTATT AAGAGAATGA

36901 GTCATGCCAT CCGTAAGATG CTTTTCTGTG ACTGGTGAGT ACTCAACCAA  
 CAGTACGGTA GGCATTCTAC GAAAAGACAC TGACCACTCA TGAGTTGGTT

36951 GTCATTCTGA GAATAGTGTA TGCGGCGACC GAGTTGCTCT TGCCCGGCGT  
 CAGTAAGACT CTTATCACAT ACGCCGCTGG CTCAACGAGA ACGGGCCGCA

37001 CAACACGGGA TAATACCGCG CCACATAGCA GAACTTTAAA AGTGCTCATC  
 GTTGTGCCCT ATTATGGCGC GGTGTATCGT CTTGAAATTT TCACGAGTAG

37051 ATTGGAAAAC GTTCTTCGGG GCGAAAAC TC AAGGATCT TACCGCTGTT  
 TAACCTTTTG CAAGAAGCCC CGCTTTTGAG AGTTCCTAGA ATGGCGACAA

37101 GAGATCCAGT TCGATGTAAC CCACTCGTGC ACCCAACTGA TCTTCAGCAT  
 CTCTAGGTCA AGCTACATTG GGTGAGCACG TGGGTTGACT AGAAGTCGTA

37151 CTTTTACTTT CACCAGCGTT TCTGGGTGAG CAAAACAGG AAGGCAAAT  
 GAAAATGAAA GTGGTCGCAA AGACCCACTC GTTTTTGTC TCCGTTTTA

37201 GCCGCAAAA AGGGAATAAG GCGGACACGG AAATGTTGAA TACTCATACT  
 CGGCGTTTTT TCCCTTATTC CCGCTGTGCC TTTACAACCTT ATGAGTATGA

37251 CTTCTTTTT CAATATTATT GAAGCATTTA TCAGGGTTAT TGTCTCATGA  
 GAAGGAAAA GTTATAATAA CTTCGTAAAT AGTCCCAATA ACAGAGTACT

37301 GCGGATACAT ATTTGAATGT ATTTAGAAAA ATAAACAAAT AGGGGTCCG  
 CGCCTATGTA TAAACTTACA TAAATCTTTT TATTTGTTTA TCCCAAGGC

37351 CGCACATTTT CCCGAAAAGT GCCACCTGAC GTCTAAGAAA CCATTATTAT  
 GCGTGTAAG GGGCTTTTCA CGGTGGACTG CAGATTCTTT GGTAATAATA

37401 CATGACATTA ACCTATAAAA ATAGGCGTAT CACGAGGCC TTCGTCTTC  
 GTACTGTAAT TGGATATTTT TATCCGCATA GTGCTCCGGG AAAGCAGAAG

37451 AAGAATTGGA TCCGAATTCT TAAT  
 TTCTTAACCT AGGCTTAAGA ATTA

FIG. 10A-47



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1 catcatcaat aatatacctt atagatggaa tgggtgccaat atgtaaataga ggtgatttta  
 61 aaaagtgtgg gccgtgtggt gattggctgt ggggttaacg gttaaaaggg gcggcgccgc  
 121 cgtgggaaaa tgacgtttta tgggggtgga gtttttttgc aagtgtgcgc gggaaatggt  
 181 acgcataaaa aggcttcttt tctcacggaa ctacttagtt tccccacggg atttaacagg  
 241 aatgaggta gttttgaccg gatgcaagtg aaaattgctg attttcgcgc gaaaactgaa  
 301 tgaggaagtg tttttctgaa taatgtggta tttatggcag ggtggagtat ttgttcaggg  
 361 ccaggtagac tttgacccat tacgtggagg tttcgattac cgtgtttttt acctgaattt  
 421 ccgcgtaccg tgtcaaagtc ttctgttttt acgtagggtg cagctgatcg ctagggtatt  
 481 tatacctcag ggtttgtgtc aagaggccac tcttgagtgc cagcgagaag agttttctcc  
 541 tctgcgccgg cagtttaata ataaaaaaat gagagatttg cgatttctgc ctcaggaaat  
 601 aatctctgct gagactggaa atgaaatatt ggagcttgtg gtgcacgcc tgatgggaga  
 661 cgatccggag ccacctgtgc agctttttga gcctcctacg cttcaggaac tgtatgattt  
 721 agaggtagag ggatcggagg attctaataga ggaagctgtg aatggctttt ttaccgattc  
 781 tatgctttta gctgctaata aaggattaga attagatccg cttttggaca ctttcaatac  
 841 tccaggggtg attgtggaaa gcggtacagg tgtaagaaaa ttacctgatt tgagttccgt  
 901 ggactgtgat ttgcaactgct atgaagacgg gtttcctccg agtgatgagg aggaccatga  
 961 aaaggagcag tccatgcaga ctgcagcggg tgagggagtg aaggctgcca atgtttggtt  
 1021 tcagttggat tgcccggagc ttctggacat ggctgtaagt cttgtgaatt tcacaggaaa  
 1081 aatactggag taaaggaact gttatgttcg cttttgttat atgaaaacc actgccactt  
 1141 tatttacagt aaagtgtggt taagttaaaa tttaaaggaa tatgctggtt ttcacatgta  
 1201 tattgagtgt gagttttgtg cttcttatta taagtcctgt gtctgatgct gatgaatcac  
 1261 catctcctga ttctactacc tcacctcctg atattcaagc acctgttctt gttggacgtgc  
 1321 gcaagcccat tcctgtgaag cttaagcctg ggaaacgtcc agcagtggag aaacttgagg  
 1381 acttgttaca ggggtggggac ggacctttgg acttgagtac acggaaacgt ccaagacaat  
 1441 aagtgttcca tatccgtggt tacttaaggt gacgtcaata tttgtgtgag agtgcaatgt  
 1501 aataaaaata tgtaactgt tcactggttt ttattgcttt ttgggcgggg actcaggtat  
 1561 ataagtagaa gcagacctgt gtggttagct cataggagct ggctttcatc catggagggt  
 1621 tgggccaattt tggaagacct taggaagact aggcaactgt tagagagcgc ttcggacgga  
 1681 gtctccggtt tttggagatt ctggttcgtc agtgaattag ctagggtagt ttttaggata  
 1741 aacaggact ataaacaaga atttgaaaag ttgttggtag attgccaggg actttttgaa  
 1801 gctcttaatt tgggccaatca ggttcacttt aaagaaaaag ttttatcagt tttagacttt  
 1861 tcaaccccag gtagaactgc tgctgctgtg gcttttctta cttttatatt agataaatgg  
 1921 atcccgcaga ctcatctcag caggggatac gttttggatt tcatagccac agcattgtgg  
 1981 agaacatgga aggttcgcaa gatgaggaca atcttaggtt actggccagt gcagcctttg  
 2041 ggtgtagcgg gaatcctgag gcatccaccg gtcatgccag cggttctgga ggaggaacag  
 2101 caagaggaca acccgagagc cggcctggac cctccagtgg aggaggcggg gtagctgact  
 2161 tgtctcctga actgcaacgg gtgcttactg gatctacgtc cactggacgg gatagggcg  
 2221 ttaagagggg gagggcatcc agtggtagtg atgctagatc tgagttggct ttaagtttaa  
 2281 tgagtcgcag acgtcctgaa accatttggt ggcatgaggt tcagaaagag ggaagggatg  
 2341 aagtttctgt attgcaggag aaatattcac tggaaacagg gaaaacatgt tggttggagc  
 2401 cagaggatga ttgggagggt gccattaaaa attatgccaa gatagctttg aggctgata  
 2461 aacagtataa gatcagtaga cggattaata tccggaatgc ttgttacata tctggaaatg  
 2521 gggctgaggt ggtaatagat actcaagaca agacagttat tagatgctgc atgatggata  
 2581 tgtggcctgg agtagtcggt atggaagcag tcacttttgt aatgtttaag tttaggggag  
 2641 atggttataa tggaatagtg tttatggcca ataccaaact tatattgcat ggtttagact  
 2701 tttttgggtt caacaatacc tgtgtagatg cctggggaca ggttagtgta cgggggtgta  
 2761 gtttctatgc gtgttggatt gccacagctg gcagaaccaa gagtcaattg tctctgaaga  
 2821 aatgcatatt ccaaagatgt aacctgggca ttctgaatga aggcgaagca agggctccgtc  
 2881 actgcgcttc tacagatact ggatgtttta ttttaattaa gggaaatgcc agcgtaaagc  
 2941 ataacatgat ttgtggtgct tccgatgaga ggcttatca aatgctcact tgtgctgggtg  
 3001 ggcatgtgaa tatgctggct actgtgcata ttgtttccca tcaacgcaa aaatggcctg  
 3061 tttttgatca caatgtgttg accaagtgca ccatgcatgc aggtgggctg agaggaatgt  
 3121 ttatgcctta ccagtgtaac atgaatcatg tgaaagtgtt gttggaacca gatgcctttt  
 3181 ccagaatgag cctaacagga atctttgaca tgaacacgca aatctggaag atcctgaggt  
 3241 atgatgatac gagatcgagg gtgcgcgcat gcgaatgcgg aggcaagcat gccaggttcc  
 3301 agccggtgtg tgtagatgtg accgaagatc tcagaccgga tcatttgggt attgcccgca  
 3361 ctggagcaga gttcggatcc agtggagaag aaactgacta aggtgagtat tgggaaaact  
 3421 ttgggggtggg attttcagat ggacagattg agtaaaaatt tgttttttct gtcttgcagc  
 3481 tgacatgagt ggaaatgctt cttttaaggg gggagtcttc agcccttate tgacagggcg  
 3541 tctcccatcc tgggcaggag ttctgcagaa tgttatggga tctactgtgg atggaagacc  
 3601 cgttcaacce gccaatctt caacgctgac ctatgctact ttaagttctt cacctttgga  
 3661 cgcagctgca gccgctgccg ccgctctgt cgcgctaac actgtgcttg gaatgggtta

FIG. 11A-1



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3721 ctatggaagc atcgtggcta attccacttc ctctaataac ctttctacac tgactcagga  
 3781 caagttactt gtccttttgg cccagctgga ggctttgacc caacgtctgg gtgaactttc  
 3841 tcagcagggt gccgagttgc gagtacaac tgagtctgct gtcggcacgg caaagtctaa  
 3901 ataaaaaaaa ttccagaatc aatgaataaa taaacgagct tgttgttgat ttaaaatcaa  
 3961 gtgtttttat ttcatttttc gcgcacggta tgccttgac caccgatctc gatcattgag  
 4021 aactcgggtg attttttcca gaatcctata gaggtgggat tgaatgttta gatacatggg  
 4081 cattaggccg tctttggggg ggagatagct ccattgaagg gattcatgct ccggggtagt  
 4141 gttgtaaate acccagtcac aacaaggctc cagtgcattg tgttgcacaa tatcttttag  
 4201 aagtaggctg attgccacag ataagccctt ggtgtaggtg ttacaaacc ggttgagctg  
 4261 ggaggggtgc attcgagggt aaattatgtg cattttggat tggattttta agttggcaat  
 4321 attgccgcca agatcccgtc ttgggttcat gttatgaagg actaccaaga cgggtgatcc  
 4381 ggtacattta ggaaatttat cgtgcagctt ggatggaaaa gcgtggaaaa atttgagac  
 4441 acccttgtgt cctccgagat tttccatgca ctcatccatg ataatagcaa tggggccgtg  
 4501 ggcagcggcg cgggcaaaca cgttccgtgg gtctgacaca tcatagttat gttcctgagt  
 4561 taaatcatca taagccattt taatgaattt ggggaggagc gtaccagatt ggggtatgaa  
 4621 tgttccttcg ggccccggag catagttccc ctacacagatt tgcatttccc aagctttcag  
 4681 ttctgagggt ggaatcatgt ccacctgggg ggctatgaag aacaccgttt cgggggaggg  
 4741 ggtgattagt tgggatgata gcaagtttct gagcaattga gatttgccac atccggtggg  
 4801 gccataaata attccgatta caggttgcag gtggtagttt agggaaacggc aactgccgtc  
 4861 ttctcgaagc aagggggcca cctcgttcat catttccctt acatgcatat tttcccgcac  
 4921 caaatccatt aggaggcgtc ctccctcctag tgatagaagt tcttgtagtg aggaaaagt  
 4981 tttcagcggg tttagaccgt cagccatggg cattttggaa agagtttgct gcaaaagttc  
 5041 tagtctgttc cacagttcag tgatgtgttc tatggcatct cgatccagca gacctcctcg  
 5101 tttcgcgggt ttggacggct cctggagtag ggtatgagac gatgggctc cagcgtgccc  
 5161 agggttcggg cctccagggg tctcagtgtt cgagtcaggg ttgtttccgt cacagtgaag  
 5221 ggggtgtgcg ctgcttgggc gcttgccagg gtgcttca gactcattct gctggtggag  
 5281 aacttctgtc gcttggcggc ctgtatgtcg gccaaagtagc agtttaccat gagttcgtag  
 5341 ttgagcgcct cggctgcgtg gcctttggcg cggagcttac ctttggaaagt tttcttgcac  
 5401 accgggcagt ataggcattt cagcgcatac agcttggggc caaggaaaat ggattctggg  
 5461 gagtatgcat ccgcgccgca ggaggcgcaa acagtttcac attccaccag ccaggttaaa  
 5521 tccggttcat tggggtcaaa aacaagtttt ccgcatatt ttttgatgcg tttcttacct  
 5581 ttggtctcca taagttcgtg tcctcgttga gtgacaaaca ggctgtccgt atctccgtag  
 5641 actgatttta caggcctctt ctccagtggg gtgcctcggg cttcttcgta caggaactct  
 5701 gaccactctg atacaaaggc gcgcgtccag gccagcacia aggaggctat gtgggagggg  
 5761 tagcgatcgt tgtcaaccag ggggtccacc ttttccaaag tatgcaaaca catgtcacc  
 5821 tcttcaacat ccaggaatgt gattggcttg taggtgtatt tcacgtgacc tgggggtccc  
 5881 gctggggggg tataaaaggg ggcggttctt tgctcttctt cactgtcttc cggatcgctg  
 5941 tccaggaacg tcagctgttg gggtaggtat tcctctcga aggcgggcat gacctctgca  
 6001 ctccaggttg cagtttctaa gaacgaggag gatttgatat tgacagtgcc ggttgagatg  
 6061 cttttcatga ggttttctc attttgggca gaaaacacia tttttttatt gtcaagtttg  
 6121 gtggcaaatg atccatacag ggcgttggat aaaagtttgg caatggatcg catggtttgg  
 6181 ttcttttctt tgtccgcgcg ctctttggcg gcgatgttga gttggacata ctccgctgccc  
 6241 aggcacttcc attcggggaa gatagttggt aattcatctg gcacgattct cacttgccac  
 6301 cctcgattat gcaaggtaat taaatccaca ctgggtggcca cctcgcctcg aaggggttca  
 6361 ttggtccaac agagcctacc tcctttccta gaacagaaag ggggaagtgg gtctagcata  
 6421 agttcatcgg gaggtctgac atccatggta aagattcccg gaagtaaate cttatcaaaa  
 6481 tagctgatgg gagtggggtc atctaaggcc atttgccatt ctccagctgc cagtgccgcg  
 6541 tcatatgggt taaggggact gcccaggggc atgggatggg tgagagcaga ggcatacatg  
 6601 ccacagatgt catagacgta gatgggatcc tcaaagatgc ctatgtaggt tggatagcat  
 6661 cgccccctc tgatacttgc tcgcacatag tcatatagtt catgtgatgg cgctagcagc  
 6721 cccggaccca agttgggtgag attgggtttt tctgttctgt agacgatctg gcgaaagatg  
 6781 gcgtgagaat tgggaagagat ggtgggtctt tgaaaaatgt tgaaatgggc atgaggtaga  
 6841 cctacagagt ctctgacaaa gtgggcataa gattcttgaa gcttgggttac cagttcggcg  
 6901 gtgacaagta cgtctagggc gcagtagtca agtgtttctt gaatgatgct ataacctggt  
 6961 tggtttttct tttcccacag ttccggttg agaaggtatt cttcgcgatc cttccagtac  
 7021 tcttctagcg gaaaccctc tttgtctgca cggtaagatc ctagcatgta gaactgatta  
 7081 actgccttgt aagggcagca gcccttctct acgggtagag agtatgcttg agcagctttt  
 7141 cgtagcgaag cgtgagtaag ggcaaagggt tctctgacca tgactttgag aaattgggat  
 7201 ttgaagtcca tgtcgtcaca ggctccctgt tcccagagtt ggaagtctac ccgtttcttg  
 7261 taggcggggg tgggcaaagc gaaagtaaca tcattgaaga gaatcttacc ggctctgggc  
 7321 ataaaattgc gagtgatgag gaaaggctgt ggtacttccg ctcgattggt gatcacctgg  
 7381 gcagctagga cgatttctgc gaaaccgttg atgttgtgct ctacgatgta taattctatg

FIG. 11A-2



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7441 aaacgcggcg tgcctctgac gtgaggtagc ttactgagct catcaaaggt taggtctgtg  
7501 gggtcagata aggcgtagtg ttcgagagcc cttcgtgca ggtgaggatt tgcattgtagg  
7561 aatgatgacc aaagatctac cgccagtgtc gtttgtaact ggtcccata ctgacgaaaa  
7621 tgccggccaa ttgccatttt ttctggagtg acacagtaga aggttctggg gtctttgttg  
7681 catcgatccc acttgagttt aatggctaga tctgaggcca tgttgacgag acgctcttct  
7741 cctgagagtt tcatgaccag catgaaagga actagttgtt tgccaaagga tccatccag  
7801 gtgtaagttt ccacatcgta ggtcaggaag agtctttctg tgcgaggatg agagccgatc  
7861 gggagaact ggatttcctg ccaccagtgt gaggattggc tgttgatgtg atggaagtag  
7921 aagtttctgc ggcgcgccga gcattcgtgt ttgtgcttgt acagacggcc gcagtagtcg  
7981 cagcgttgca cgggttgat ctctgtaatg agctgtacct ggcttccctt gacgagaaat  
8041 ttcagtggga agccgaggcc tggcgattgt atctcgtgct ctctatatt cgctgtatcg  
8101 gcctgttcat cttctgtttc gatgggtggc atgctgacga gccccgcgg gaggcaagtc  
8161 cagacctcgg cgcgggaggg gcggagctga aggacgagag cgcgcaggct ggagctgtcc  
8221 agagtcctga gacgctgcgg actcaggtta gtaggtaggg acagaagatt aacttgcattg  
8281 atcttttcca gggcgtgcgg gaggttcaga tggtagttga ttccacagg ttcgtttgta  
8341 gagacgtcaa tggcttgacg ggttccgtgt ctttgggcg ccactaccgt acctttgttt  
8401 tttcttttga tgggtgggtg ctctcttctg tcttgcatgc tcagaagcgg tgacggggac  
8461 gcgcgcgggg cggcagcggg tgttccggac cggggggcat ggctggtagt ggcacgtcgg  
8521 cgcgcgcac gggcaggttc tggattgctg ctctgagaag acttgctgct gccaccacgc  
8581 gtcgattgac gtcttgtatc tgacgtctct gggtgaaagc taccggcccc gtgagcttga  
8641 acctgaaaga gagttcaaca gaatcaattt cggtagctgt aacggcagct tgtctcagta  
8701 tttcttgtac gtcaccagag ttgtcctggg aggcgatctc cgccatgaac tgctcgattt  
8761 ctctctctg aagatctccg cgaccgctc tttcagcggg ggccgcgagg tcattggaga  
8821 tacggcccat gagttgggag aatgcattca tgccgcctc gttccagacg cggctgtaa  
8881 ccacggcccc ctccgagctt cttgcgcgca tcaccacctg agcaggtta agctccactg  
8941 gtctgggtgaa gaccgcatag ttgcataggc gctgaaaaag gtagttgagt gtgggtggca  
9001 tgtgttcggc gacgaagaaa tacatgatcc atcgtctcag cggcatttcg ctaacatcgc  
9061 ccagagcttc caagcgtcc atggcctcgt agaagtccac ggcaaaatta aaaaactggg  
9121 agtttcgcgc ggacacggtc aattcctcct cgagaagacg gatgagttcg gctatgggtg  
9181 cccgtacttc gcgttcgaag gctcccgga tctcttctc ctcttctatc tcttctcca  
9241 ctaacatctc ttcttctctc tcaggcgggg gcggaggggg cacgcggcga cgtcgacggc  
9301 gcacgggcaa acggtcgatg aatcgttcaa tgacctctc gcggcggcgg cgcattgttt  
9361 cagtgcggc gcggccgttc tcgcgcggtc gcagagtaaa aacaccgccc cgcattctct  
9421 taaagtgggt actgggaggt tctccgtttg ggagggagag ggcgctgatt atacatttta  
9481 ttaattggcc cgtagggact gcgcgcagag atctgatcgt gtcaagatcc acgggatctg  
9541 aaaacctttc gacgaaagcg tctaaccagt cacagtcaca aggtaggctg agtacggctt  
9601 cttgtgggcg ggggtgggta tgtgttcggg ctgggtcttc tgtttcttct tcactctcggg  
9661 aaggtgagac gatgctgctg gtgatgaaat taaagtaggc agttctaaga cggcggatgg  
9721 tggcgaggag caccaggtct ttgggtccgg cttgctggat acgcaggcga ttggccattc  
9781 cccaagcatt atcctgacat ctagcaagat cttttagta gtcttgcatt agccgttcta  
9841 cgggcacttc ttcctcacc gttctgcat gcatacgtgt gaggccaaat ccgcgcatg  
9901 gttgtaccag tgccaagtca gctacgactc tttcggcgag gatggcttgc tgtacttggg  
9961 taagggtggc ttgaaagtca tcaaaatcca caaagcgggt gtaagccctt gtattaatgg  
10021 tgtaagcaca gttggccatg actgaccagt taactgtctg gtgaccaggg cgcacgagct  
10081 cgggtgtatt aaggcgcgaa taggcgcggg tgtcaaagat gtaatcgttg cagggtgcga  
10141 ccagatactg gtaccctata agaaaatgag gcggtgggtg gcggtagaga ggccatcgtt  
10201 ctgtagctgg agcgcaggg gcgaggtctt ccaacataag gcggtgatag ccgtagatgt  
10261 acctggacat ccaggtgatt cctgcggcgg tagtagaagc ccgaggaaac tcgcgtacgc  
10321 ggttccaaat gttgcgtagc ggcattgaagt agttcattgt aggcacgggt tgaccagtga  
10381 ggcgcgcgca gtcattgatg ctctatagac acggagaaaa tgaaagcgtt cagcgactcg  
10441 actccgtagc ctggaggaaac gtgaacgggt tgggtcgcgg tgtaccgccg ttcgagactt  
10501 gtactcgagc cggccggagc cgcggctaac gtggatttgg cactcccgtc tcgaccagc  
10561 ctacaaaaat ccaggatagc gaatcgagtc gttttgctgg tttccgaatg gcaggaagt  
10621 gagtcctatt tttttttttt ttttgcgct cagatgcatc ccgtgctgcg acagatgccc  
10681 cccaacaac agccccctc gcagcagcag cagcagcagc aaccacaaa ggctgtccct  
10741 gcaactactg caactgccgc cgtgagcggg gcgggacagc ccgctatga tctggacttg  
10801 gaagagggcg aaggactggc acgtctaggt gcgcttctgc ccgagcggca tccgcgagtt  
10861 caactgaaaa aagattctcg cgaggcgtat gtgccccaac agaacctatt tagagacaga  
10921 agcggcgagg agccggagga gatgcgagct tcccgttcta acgcgggtcg tgagctcgt  
10981 cacggtttgg accgaagacg agtgttgcca gacgaggatt tcgaagtga tgaagtgaca  
11041 gggatcagtc ctgccagggc acacgtggct gcagccaacc ttgtatcggc ttacgagcag  
11101 acagtaaagg aagagcgtaa cttccaaaag tcttttaata atcatgtgcg aaccctgatt

FIG. 11A-3



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11161 gcccgcaag aagttaccct tggtttgatg ctttgtggg atttgatgga agctatcatt  
11221 cagaacccta ctagcaaacc tctgaccgcc cagctgtttc tggtggtgca acacagcaga  
11281 gacaatgagg ctttcagaga ggcgctgctg aacatcaccg aacccgaggg gagatggttg  
11341 tatgatctta tcaacattct acagagtatc atagtgcagg agcggagcct gggcctggcc  
11401 gagaaggtag ctgccatcaa ttactcgggt ttgagcttgg gaaaatatta cgctcgcaaa  
11461 atctacaaga ctccatacgt tcccatagac aaggaggtga agatagatgg gttctacatg  
11521 cgcacgacgc tcaaggtctt gaccctgagc gatgatcttg ggggtgatcg caatgacaga  
11581 atgcatcgcg cggtagcgc cagcaggagg cgcgagtaa gcgacagggg actgatgcac  
11641 agtttgcaaa gagctctgac tggagctgga accgaggggt agaattactt cgacatggga  
11701 gctgacttgc agtggcagcc tagtcgcagg gctctgagcg ccgacgagcc aggatgtgag  
11761 cttccttaca tagaagaggc ggatgaaggc gaggaggaag agggcgagta cttggaagac  
11821 tgatggcaca acccgtgttt tttgctagat ggaacagcaa gcaccggatc ccgcaatgcg  
11881 ggcggcgctg cagagccagc cgtccggcat taactcctcg gacgattgga cccaggccat  
11941 gcaacgtatc atggcgttga cgactcgcaa cccgaagcc tttagacagc aaccaccaggc  
12001 caaccgtcta tcggccatca tggagctgt agtgccttcc cgatctaate cactcatga  
12061 gaaggctctg gccatcgtga acgcgttggg ggagaacaaa gctattcgtc cagatgaggc  
12121 cggactggta tacaacgctc tcttagaacg cgtggctcgc tacaacagta gcaatgtgca  
12181 aaccaatttg gaccgtatga taacagatgt acgcgaagcc gtgtctcagc gcgaaaggtt  
12241 ccagcgtgat gccaacctgg gttcgtggt ggcgttaaat gctttcttga gtactcagcc  
12301 tgctaatgtg ccgcgtggtc aacaggatta tactaacttt ttaagtgctt tgagactgat  
12361 ggtatcagaa gtacctcaga gcgaagtgt tcagtccggg cctgattact tctttcagac  
12421 tagcagacag ggcttgcaga cggtaaactt gagccaagct tttaaaaacc ttaaaggttt  
12481 gtggggagtg catgccccgg taggagaaag agcaaccgtg tctagcttgt taactccgaa  
12541 ctccgcctg ttattactgt tggtagctcc tttcaccgac agcggtagca tcgaccgtaa  
12601 ttcctatttg ggttacctac taaacctgta tcgcgaagcc atagggcaaa gtcagggtgga  
12661 cgagcagacc tatcaagaaa ttacccaagt cagtcgcgct ttgggacagg aagacactgg  
12721 cagtttgga gccactctga acttcttctg taccaatcgg tctcaaaaga tcctcctca  
12781 atatgctctt actgcggagg aggagaggat ccttagatat gtgcagcaga gcgtgggatt  
12841 gtttctgatg caagaggggg caactccgac tgcagcactg gacatgacag cgcgaaatat  
12901 ggagcccagc atgtatgcca gtaaccgacc tttcattaac aaactgctgg actacttgca  
12961 cagagctgcc gctatgaact ctgattattt caccaatgcc atcttaaacc cgcactggct  
13021 gccccacct ggtttctaca cgggcgaata tgacatgccc gaccctaag acggatttct  
13081 gtgggacgac gtggacagcg atgttttttc acctctttct gatcatcgca cgtggaaaaa  
13141 ggaaggcggg gatagaatgc attcttctgc atcgctgtcc ggggtcatgg gtgctaccgc  
13201 ggctgagccc gagtctgcaa gtctttttcc tagtctacce ttttctctac acagtgtacg  
13261 tagcagcgaa gtgggtagaa taagtcgccc gagtttaatg ggcgaagagg agtacctaaa  
13321 cgattccttg ctcagaccgg caagagaaaa aaatttccca aacaatggaa tagaaagttt  
13381 ggtggataaa atgagtagat ggaagactta tgctcaggat cacagagacg agcctgggat  
13441 catggggact acaagtagag cgagccgtag acgcccagcg catgacagac agaggggtct  
13501 tgtgtgggac gatgaggatt cggccgatga tagcagcgtg ttggacttgg gtgggagagg  
13561 aaggggcaac ccgtttgctc atttgcgcc tcgcttgggt ggtatgttgt gaaaaaaat  
13621 aaaaaagaaa aactcaccaa ggccatggcg acgagcgtac gttcgttctt ctttattatc  
13681 tgtgtctagt ataatgaggc gagtcgtgct aggcggagcg gtggtgtatc cggagggtcc  
13741 tcctccttcg tacgagagcg tgatgcagca gcagcaggcg acggcgggtg tgcaatcccc  
13801 actggaggct ccctttgtgc ctccgcgata cctggcacct acggagggca gaaacagcat  
13861 tcgttactcg gaactggcac ctacgtacga taccaccagg ttgtatctgg tggacaacaa  
13921 gtcggcggac attgcttctc tgaactatca gaatgaccac agcaacttct tgaccacggt  
13981 ggtgcagaac aatgacttta ccctacgga agccagcacc cagaccatta actttgatga  
14041 acgatcgcgg tggggcggtc agctaaagac catcatgcat actaacatgc caaacgtgaa  
14101 cgagtatatg tttagtaaca agttcaaagc gcgtgtgatg gtgtccagaa aacctccga  
14161 cgggtgctgca gttggggata cttatgatca caagcaggat attttggaa atgagtgggt  
14221 cgagtttact ttgccagaag gcaacttttc agttactatg actattgatt tgatgaacaa  
14281 tgccatcata gataattact tgaaagtggg tagacagaat ggagtgtctg aaagtacat  
14341 tgggtgtaag ttcgacacca ggaacttcaa gctgggatgg gatcccgaaa ccaagttgat  
14401 catgcctgga gtgtatacgt atgaagcctt ccctcctgac attgtcttac tgctggctg  
14461 cggagtggat tttaccgaga gtcgtttgag caaccttctt ggtatcagaa aaaaacagcc  
14521 atttcaagag ggttttaaga ttttgatga agatttagaa ggtggtaata ttccggccct  
14581 cttggatgta gatgcctatg agaacagtaa gaaagaacaa aaagccaaaa tagaagctgc  
14641 tacagctgct gcagaagcta aggcaaacat agttgccagc gactctacaa gggttgctaa  
14701 cgctggagag gtcagaggag acaattttgc gccaacacct gttccgactg cagaatcatt  
14761 attggccgat gtgtctgatg gaacggagct gaaactcact attcaacctg tagaaaaaga  
14821 tagtaagaat agaagctata atgtgttggg agacaaaatc aacacagcct atcgcagttg

FIG. 11A-4



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14881 gtatcttttcg tacaattatg gcgatcccga aaaaggagtg cgttcctgga cattgctcac  
 14941 cacctcagat gtcacctgcg gagcagagca ggtttactgg tcgcttccag acatgatgaa  
 15001 ggatcctgtc actttccgct ccactagaca agtcagtaac taccctgtgg tgggtgcaga  
 15061 gcttatgccc gtcttctcaa agagcttcta caacgaacaa gctgtgtact cccagcagct  
 15121 ccgccagtcc acctcgctta cgcacgtctt caaccgcttt cctgagaacc agattttaat  
 15181 ccgtccgccc ggcgccacca ttaccaccgt cagtgaaaac gttcctgctc tcacagatca  
 15241 cgggaccctg ccgttgcgca gcagtatccg gggagtccaa cgtgtgaccg ttactgacgc  
 15301 cagacgccgc acctgtccct acgtgtacaa ggcactgggc atagtcgcac cgcgcgtcct  
 15361 ttcaagccgc actttctaaa aaaaaaatgt ccattcttat ctgcgccagt aataacaccg  
 15421 gttgggggtct ggcgcgtcca agcaagatgt acggaggcgc acgcaaactg tctaccaaac  
 15481 atcccgtgcg tgttcgcgga cattttcgcg ctccatgggg tgccctcaag ggccgcactc  
 15541 gcgttcgaac caccgtcgat gatgtaatcg atcagggtgg tgccgacgcc cgtaattata  
 15601 ctctactgct gcctacatct actgtggatg cagttattga cagtgtagtg gctgacgctc  
 15661 gcaactatgc tcgacgtaag agccggcgaa ggcgcattgc cagacgccac cgagctacca  
 15721 ctgccatgcy agccgcaaga gctctgctac gaagagctag acgcgtgggg cgaagagcca  
 15781 tgcttagggc ggccagacgt gcagcttcgg gcgccagcgc cggcaggctc cgcaggcaag  
 15841 cagccgctgt cgcagcggcg actattgccc acatggcca atcgcgaaga ggcaatgtat  
 15901 actgggtgcy tgacgctgcc accggtcaac gtgtaccctg gcgcaccctg ccccctcgca  
 15961 cttagaagat actgagcagt ctccgatggt gtgtcccagc ggcgaggatg tccaagcgca  
 16021 aatacaagga agaaatgctg caggttatcg cacctgaagt ctacggcca ccgttgaagg  
 16081 atgaaaaaaaa accccgcaaa atcaagcggg ttaaaaagga caaaaagaa gaggaagatg  
 16141 gcgatgatgg gctggcgagg tttgtgcgcy agtttgcccc acggcgacgc gtgcaatggc  
 16201 gtgggcycaa agttcgacat gtgttgagac ctggaacttc ggtggtcttt acaccggcg  
 16261 agcgttcaag cgctactttt aagcgttctt atgatgaggt gtacggggat gatgatattc  
 16321 ttgagcaggc ggctgaccga ttaggcagat ttgcttatgg caagcgtagt agaataactt  
 16381 ccaaggatga gacagtgtca atacccttgg atcatggaaa tcccaccctt agtcttaaac  
 16441 cggtcacttt gcagcaagtg ttaccctgta ctccgcgaac aggtgttaaa cgcgaagggtg  
 16501 aagatttgta tcccactatg caactgatgg tacccaacag ccagaagttg gaggacgttt  
 16561 tggagaaagt aaaagtggat ccagatattc aacctgaggt taaagtgaga cccattaagc  
 16621 aggtagcgc tggtctgggg gtacaaactg tagacattaa gattcccact gaaagtatgg  
 16681 aagtgcaaac tgaaccgcga aagcctactg ccacctccac tgaagtgcaa acggatccat  
 16741 ggatgcccac gcctattaca actgacgcgc ccggtcccac tcgaagatcc cgacgaaagt  
 16801 acggtccagc aagtctgttg atgcccatt atgttgtaga cccatctatt attcctactc  
 16861 ctgggtaccg aggcactcgc tactatcgca gccgaaacag tacctcccgc cgtcgccgca  
 16921 agacacctgc aaatcgcagt cgtcgccgta gacgcacaag caaacgact cccggcgccc  
 16981 tgggtcggca agtgtaccgc aatggtagtg cggaaccttt gacactgccg cgtgcgcgtt  
 17041 accatccgag tatcactact taatcaatgt tgccgctgcc tccttgcaga tatggccctc  
 17101 acttgctgcc ttcgcgttcc catcactggt taccgaggaa gaaactcgcg ccgtagaaga  
 17161 gggatgttgg gacgcggaat gcgacgctac aggcgacggc gtgctatccg caagcaattg  
 17221 cggggtggtt ttttaccagc cttaattcca attatcgctg ctgcaattgg cgcgatacca  
 17281 ggcatagctt ccgtggcggt tcaggcctcg caacgacatt gacattggaa aaaaaacgta  
 17341 taaataaaaa aaaatacaat ggactctgac actcctggtc ctgtgactat gttttcttag  
 17401 agatggaaga catcaatttt tcatccttgg ctccgcgaca cggcacgaag ccgtacatgg  
 17461 gcacctggag cgacatcggc acgagccaac tgaacggggg cgccttcaat tggagcagta  
 17521 tctggagcgg gcttaaaaaat tttggctcaa ccataaaaac atacgggaac aaagcttggg  
 17581 acagcagtac aggacaggcg cttagaaata aacttaaaga ccagaacttc caacaaaaag  
 17641 tagtcgatgg gatagcttcc ggcataaatg gagtggtaga tttggctaac caggctgtgc  
 17701 agaaaaagat aaacagtcgt ttggaccgcg gccagcaac cccagggtgaa atgcaagtgg  
 17761 aggaagaaat tcctccgcca gaaaaacgag gcgacaagcg tccgcgtccc gatttggag  
 17821 agacgctggg gacgcgcgta gatgaaccgc cttcttatga ggaagcaacg aagcttggaa  
 17881 tgcccaccac tagaccgata gccccaatgg ccaccggggg gatgaaacct tctcagttgc  
 17941 atcgaccctg caccttggat ttgccccctc ccctgctgc tactgctgta cccgcttcta  
 18001 agcctgtcgc tgccccgaaa ccagtcgccc tagccaggtc acgtcccggg ggcgctcctc  
 18061 gtccaaatgc gcaactggcaa aatactctga acagcatcgt gggcttaggc gtgcaaagtg  
 18121 taaaacgccg tcgctgcttt taattaaata tggagtagcg ctttaacttgc ctatctgtgt  
 18181 atatgtgtca ttacacgccg tcacagcagc agaggaaaaa aggaagaggt cgtgcgtcga  
 18241 cgctgagtta ctttcaagat ggccacccca tcgatgctgc cccaatgggc ataatgcac  
 18301 atcgccggac aggatgcttc ggagtacctg agtccgggtc tgggtgcagtt cgcgccgccc  
 18361 acagacacct acttcaatct gggaaataag tttagaaatc ccaccgtagc gccgaccac  
 18421 gatgtgacca ccgaccgtag ccagcggctc atgttgcgct tcgtgcccgt tgaccgggag  
 18481 gacaatacat actcttacia agtgcggtac accctggccg tgggcygaaa cagagtgtg  
 18541 gatatggcca gcacgttctt tgacattagg ggcgtgttgg acagaggctc cagtttcaaa

FIG. 11A-5



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18601 ccctattctg gtacggctta caactctctg gctcctaaag gcgctccaaa tgcattctcaa  
 18661 tggattgcaa aaggcgtacc aactgcagca gccgcaggca atgggtgaaga agaacatgaa  
 18721 acagaggaga aaactgctac ttacactttt gccaatgctc ctgtaaaagc cgaggctcaa  
 18781 attacaaaag agggcttacc aataggtttg gagatttcag ctgaaaacga atctaaaccc  
 18841 atctatgcag ataaacttta tcagccagaa cctcaagtgg gagatgaaac ttggactgac  
 18901 ctagacggaa aaaccgaaga gtatggaggc agggctctaa agcctactac taacatgaaa  
 18961 ccctgttacg ggtcctatgc gaagcctact aatttaaaag gtggtcaggc aaaaccgaaa  
 19021 aactcggaac cgtcagagtga aaaaattgaa tatgatattg acatggaatt ttttgataac  
 19081 tcatcgcaaa gaacaaactt cagtcctaaa attgtcatgt atgcagaaaa tgtaggtttg  
 19141 gaaacgccag aactcctatg agtgtaaaaa cctggaacag aagacacaag ttccgaagct  
 19201 aatttgggac aacagtctat gcccaacaga cccaactaca ttggcttcag agataacttt  
 19261 attggactca tgtactataa cagtactggg aacatggggg tgctggctgg tcaagcgtct  
 19321 cagttaaatg cagtgggtga cttgcaggac agaaacacag aactttctta ccaactcttg  
 19381 cttgactctc tgggcgacag aaccagatac tttagcatgt ggaatcaggc tgtggacagt  
 19441 tatgatcctg atgtacgtgt tattgaaaat catgggtgtg aagatgaact tccaactat  
 19501 tgttttccac tggacggcat aggtgttcca acaaccagtt acaaatcaat agttccaaat  
 19561 ggagaagata ataataattg gaaagaacct gaagtaaag gaacaagtga gatcggacag  
 19621 ggtaatttgt ttgccatgga aattaacctt caagccaatc tatggcgaag tttcctttat  
 19681 tccaatgtgg ctctgtatct ccagactcg tacaataca ccccgctcaa tgtcactctt  
 19741 ccagaaaaca aaaacaccta cgactacatg aacgggcggg tggtgccgcc atctctagta  
 19801 gacacctatg tgaacattgg tgccagggtg tctctggatg ccatggaca tgfcaacca  
 19861 ttcaaccacc accgtaacgc tggcttgcgt taccgatcta tgcttctggg taacggacgt  
 19921 tatgtgcctt tccacataca agtgcctcaa aaattcttcg ctgttaaaaa cctgctgctt  
 19981 ctcccaggct cctacactta tgagtggaac tttaggaagg atgtgaacat ggttctacag  
 20041 agttccctcg gtaacgacct gcgggtagat ggcgccagca tcagtttcac gagcatcaac  
 20101 ctctatgcta cttttttccc catggctcac aacaccgctt ccacccttga agccatgctg  
 20161 cggaatgaca ccaatgatca gtcattcaac gactacctat ctgcagctaa catgctctac  
 20221 cccattcctg ccaatgcaac caatattccc atttccattc cttctcgcaa ctgggaggct  
 20281 ttcagaggct ggtcatttac cagactgaaa accaaagaaa ctccctcttt ggggtctgga  
 20341 tttgaccctt actttgtcta ttctggttct attccctacc tggatggtac cttctacctg  
 20401 aaccacactt ttaagaaggt ttccatcatg tttgactctt cagtgagctg gcctggaaat  
 20461 gacaggttac tatctcctaa cgaatttgaa ataaagcgca ctgtggatgg cgaaggctac  
 20521 aacgtagccc aatgcaacat gaccaaagac tggttcttgg tacagatgct cgccaactac  
 20581 aacatcggct atcagggtt ctacattcca gaaggataca aagatcgcat gtattcattt  
 20641 ttcagaaact tccagcccat gagcaggcag gtgggtgatg aggtcaatta caaagacttc  
 20701 aaggccgtcg ccatacccta ccaacacaac aactctggct ttgtgggta catggctccg  
 20761 accatgcgcc aagggtcaacc ctatcccgtt aactatccct atccactcat tggaaacaact  
 20821 gccgtaaata gtgttacgca gaaaaagttc ttgtgtgaca gaaccatgtg gcgcataccg  
 20881 ttctcgagca acttcatgtc tatggggggc cttacagact tgggacagaa tatgctctat  
 20941 gccaaactcag ctcatgctct ggacatgacc tttgagggtg atcccatgga tgagcccacc  
 21001 ctgctttatc ttctcttcca agtttttcgac gtgggtcagag tgcattcagc acaccgaggc  
 21061 atcatcgagg cagtctacct gcgtacaccg ttctcggccg gtaacgctac cacgtaagaa  
 21121 gcttcttgct tcttgcaaat agcagctgca accatggcct gcggatccca aaacggctcc  
 21181 agcgagcaag agctcagagc cattgtccaa gacctgggtt gcggacccta ttttttggga  
 21241 acctacgata agcgttccc ggggttcatg gccccgata agctcgctg tgccattgta  
 21301 aatacggccg gacgtgagac ggggggagag cactgggttg ctttcgggtg gaaccacagt  
 21361 tctaacacct gctacctttt tgatcctttt ggattctcgg atgatcgtct caaacagatt  
 21421 taccagtttg aatatgaggg tctcctgcgc cgcagegctc ttgctaccaa ggaccgctgt  
 21481 attacgtgg aaaaatctac ccagaccgtg cagggcccc gttctgccc ctgaggactt  
 21541 ttctgctgca tgttccttca cgcctttgtg cactggcctg accgtcccat ggacggaaac  
 21601 cccaccatga aattgctaac tggagtgcca aacaacatgc ttcattctcc taaagtccag  
 21661 cccaccctgt gtgacaatca aaaagcactc taccattttc ttaataccca ttcgccttat  
 21721 tttcgctctc atcgtacaca catcgaaagg gccactgcgt tcgaccgtat ggatgttcaa  
 21781 taatgactca tgtaaacaaac gtgttcaata aacatcactt tattttttta catgtatcaa  
 21841 ggctctggat tacttattta tttacaagtc gaatgggttc tgacgagaat cagaatgacc  
 21901 cgcaggcagt gatacgttgc ggaactgata cttgggttgc cacttgaatt cgggaatcac  
 21961 caacttggga accggtatat cgggcaggat gtcactccac agctttcttg tcagctgcaa  
 22021 agtccaagc aggtcaggag ccgaaatctt gaaatcaca ttaggaccag tgctctgagc  
 22081 gcgagagttg cggtaacacc gattgcagca ctgaaacacc atcagcgacg gatgtctcac  
 22141 gcttgccagc acgggtgggat ctgcaatcat gccacatcc agatcttcag cattggcaat  
 22201 gctgaacggg gtcattcttc aggtctgcct acccatggcg ggcaccat taggcttgtg  
 22261 gttgcaatcg cagtgcaggg ggatcagtat catcttggcc tgatcctgct tgattcctgg

FIG. 11A-6



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22321 atacacggct ctcatgaaag catcatattg cttgaaagcc tgctgggctt tactaccctc  
 22381 ggtataaaac atcccgcagg acctgctcga aaactgggta gctgcacagc cggcatcatt  
 22441 cacacagcag cggggcgtcat tgttggctat ttgcaccaca cttctgcccc agcgggtttg  
 22501 ggtgattttg gttcgcctcg gattctcctt taaggctcgt tgctcgttct cgtggccac  
 22561 atccatctcg ataactctgct ccttctgaat cataatattg ccatgcagge acttcagctt  
 22621 gccctcataa tcattgcagc catgaggcca caacgcacag cctgtacatt cccaattatg  
 22681 gtgggcgatc tgagaaaaag aatgtatcat tccctgcaga aatcttccca tcatcgtgct  
 22741 cagtgtcttg tgactagtga aagttaactg gatgcctcgg tgctcttctg ttacgtactg  
 22801 gtgacagatg cgcttgtatt gttcgtgttg ctcaggcatt agtttaaaac aggttctaag  
 22861 ttcgttatcc agcctgtact tctccatcag cagacacatc acttccatgc ctttctccca  
 22921 agcagacacc aggggcaagc taatcggatt cttaacagtg caggcagcag ctcccttagc  
 22981 cagaggggtca tctttagcga tcttctcaat gcttcttttg ccatccttct caacgatgcg  
 23041 cacgggcggg tagctgaaac ccactgctac aagttgcgcc tcttctcttt cttcttcgct  
 23101 gtcttgactg atgtcttgca tggggatag tttggctctt cttggcttct ttttgggggg  
 23161 tatcggagga ggaggactgt cgctccgttc cggagacagg gaggattgtg acgtttcgtc  
 23221 caccattacc aactgactgt cggtagaaga acctgacccc acacggcgac aggtgttttt  
 23281 cttcgggggc agaggtggag gcgattgcga agggctgcgg tccgacctgg aaggcggatg  
 23341 actggcagaa ccccttccgc gttcgggggt gtgctccctg tggcggtcgc ttaactgatt  
 23401 tccttcgagg ctggccattg tgttctccta ggcagagaaa caacagacat ggaaactcag  
 23461 ccattgctgt caacatcgcc acgagtcca tcacatctcg tcctcagcga cgaggaaaag  
 23521 gagcagagct taagcattcc accgcccagt cctgccacca cctctaccct agaagataag  
 23581 gaggtcagc catctcatga catgcagaat aaaaaagcga aagagtctga gacagacatc  
 23641 gagcaagacc cgggctatgt gacaccggtg gaacacgagg aagagttaa acgctttcta  
 23701 gagagagagg atgaaaactg cccaaaacag cgagcagata actatcacca agatgctgga  
 23761 aatagggatc agaacaccga ctacctcata gggcttgacg ggggaagacg gctccttaa  
 23821 catctagcaa gacagtcgct catagtcaag gatgcattat tggacagaa tgaagtgcc  
 23881 atcagtgtgg aagagctcag ctgcgctac gagcttaacc ttttttcacc tcgtactccc  
 23941 cccaaacgtc agccaaacgg cacctgcgag ccaaatcctc gcttaaactt ttatccagct  
 24001 tttgctgtgc cagaagtact ggctacctat cacatctttt ttaaaaatca aaaaattcca  
 24061 gtctcctgcc gcgctaactg caccgcgcc gatgccctac tcaatctggg acctggttca  
 24121 cgcttacctg atatagcttc cttggaagag gttccaaaga tcttcgaggg tctgggcaat  
 24181 aatgagactc gggccgcaa tgctctgcaa aaggagaaa atggcatgga tgagcatcac  
 24241 agcgttctgg tggaaattgga aggcgataat gccagactcg cagtactcaa gcgaagcgtc  
 24301 gaggtcacac acttcgcata tcccgtgtc aacctgcccc ctaaagtcat gacggcggtc  
 24361 atggaccagt tactcattaa gcgcgcaagt cccctttcag aagacatgca tgaccagat  
 24421 gcctgtgatg agggtaaacc agtggctcagt gatgagcagc taaccgatg gctgggcacc  
 24481 gactctcccc gggatttggga agagcgtcgc aagcttatga tggcctgggt gctgggtacc  
 24541 gtagaactag agtgtctccg acgtttcttt accgattcag aaacctgcg caaactcgaa  
 24601 gagaatctgc actacacttt tagacacggc tttgtgaggc aggcattgca gatattcaac  
 24661 gtggaactca ccaacctggg ttcctacatg ggtattctgc atgagaatcg cctaggacaa  
 24721 agcgtgctgc acagaccct taagggggaa gcccgccgtg attacatccg cgattgtgtc  
 24781 tatctctacc tgtgccacac gtggcaaac ggcattgggt tatggcagca atgtttagaa  
 24841 gaacagaact tgaaagagct tgacaagctc ttacagaaat ctcttaaggt tctgtggaca  
 24901 gggttcgacg agcgcaccgt cgcttccgac ctggcagacc tcactctccc agagcgtctc  
 24961 agggttactt tgcgaaacgg attgcctgac tttatgagcc agagcatgct taacaatttt  
 25021 cgctctttca tcctggaacg ctccggtatc ctgcccgcca cctgctgcgc actgccctcc  
 25081 gactttgtgc ctctcaccta ccgagagtgc cccccgccc tatggagtca ctgctacctg  
 25141 ttcgctctgg ccaactatct ctctaccac tcggatgtga tcgaggatgt gagcggagac  
 25201 ggcttgctgg agtgccactg ccgctgcaat ctgtgcacgc cccaccggtc cctagcttgc  
 25261 aacccccagt tgatgagcga aaccagata ataggcact ttgaattgca aggccccagc  
 25321 agccaaggcg atgggtcttc tcctgggcaa agtttaaaac tgacccccgg actgtggacc  
 25381 tccgcctact tgcgcaagt tgctccggaa gattaccacc cctatgaaat caagtctat  
 25441 gaggaccaat cacagcctcc aaaggccgaa ctttcggctt gcgtcatcac ccagggggca  
 25501 attctggccc aattgcaagc catccaaaaa tcccgcgaag aatttctact gaaaaaggt  
 25561 aagggggtct accttgacc ccagaccggc gaggaactca acacaagggt ccctcaggat  
 25621 gtccaacga cgagaaaaca agaagttgaa ggtgcagccg ccgccccag aagatatgga  
 25681 ggaagattgg gacagtcagg cagaggaggc ggaggaggac agtctggagg acagtctgga  
 25741 ggaagacagt ttggaggagg aaaacgagga ggcagaggag gtggaagaag taaccgcca  
 25801 caaacagtta tcctcggctg cggagacaag caacagcgt accatctccg ctccgagtcg  
 25861 aggaaccgg cggcgtccca gcagtagatg ggacgagacc ggacgcttcc cgaaccaac  
 25921 cagcgttcc aagaccggt aagaggatcg gcagggatac aagtcctggc gggggcataa  
 25981 gaatgccatc atctcctgct tgcatgagt cgggggcaac atatccttca cgcggcgcta

FIG. 11A-7



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26041 cttgctattc caccatgggg tgaactttcc gcgcaatggt ttgcattact accgtcacct  
 26101 ccacagcccc tactatagcc agcaaattccc gacagtctcg acagataaag acagcggcgg  
 26161 cgacctccaa cagaaaacca gcagcggcag ttagaaaata cacaacaagt gcagcaacag  
 26221 gaggattaaa gattacagcc aacgagccag cgcaaaccgg agagttaaga aatcggatct  
 26281 ttccaaccct gtatgccatc ttccagcaga gtcgggggtca agagcaggaa ctgaaaataa  
 26341 aaaaccgatc tctgcgttcg ctcaccagaa gttgtttgta tcacaagagc gaagatcaac  
 26401 ttcagcgcac tctcgaggac gccgaggctc tcttcaacaa gtactgcgcy ctgactctta  
 26461 aagagtaggc agcgaccgcy cttattcaaa aaaggcggga attacatcat cctcgacatg  
 26521 agtaaagaaa ttcccacgcc ttacatgtgg agttatcaac cccaaatggg attggcagca  
 26581 ggcgcctccc aggactactc caccgcgatg aattggctca ggcgcgggcy tctatgatt  
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 26701 accaccacgc cccgccaaac ccttaattccc agaaattggc ccgcccct agtgtaccag  
 26761 gaaagtcccg ctcccaccac tgtattactt cctcgagacy cccaggccga agtccaaatg  
 26821 actaatgcag gtgcgcagtt agctggcggc tccaccctat gtcgtcacag gcctcggcat  
 26881 aatataaaac gcctgatgat cagaggccga ggtatccage tcaacgacya gtcggtgacy  
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 27061 ggaatcggga ccggttcaatt tgtagaggag tttactccct ctgtctactt caacccttc  
 27121 tccggatctc ctgggcacta cccggacyg ttcataccga acttcgacyc gattagcyg  
 27181 tcagtggacy gctacgattg atgtctggtg acgcygctga gctatctcgg ctgcyacatc  
 27241 tagaccactg ccgcccctt cgctgctttg cccgggaact tattgagttc atctacttcg  
 27301 aactccccaa ggatcacctc caaggtccgg cccacggagt gcygattact atcgaaggca  
 27361 aatagactc tcgcctgcaa cgaattttct cccagcygcy cgtgctgatc gagcygacc  
 27421 agggaaacac cacggtttcc atctactgca tttgtaatca ccccgattg catgaaagcc  
 27481 tttgctgtct tatgtgtact gagtttaata aaaactgaa taagactctc ctacggactg  
 27541 ccgcttcttc aaccggtatt ttacaaccag aagaacaaaa cttttcctgt cgtccaggac  
 27601 tctgttaact tcaccttcc tactcaaaa ctagaagctc aacgactaca ccgcttttcc  
 27661 agaagcattt tccctactaa tactacttcc aaaaccggag gtgagctcca cggctctccct  
 27721 acagaaaacc cttgggtgga agcgggcctt gtagtactag gaattcttgc ggggtggctt  
 27781 gtgattattc tttgctacct atacacacct tgcttactt tcttagtggg gttgtggtat  
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 27901 gttctgcaa ttacgatcca tgtctagact ttgaccaga aaactgcaca cttacttttg  
 27961 caccgacac aagccgcatc tgtggagttc ttattaagtg cggatgggaa tgcaggctccg  
 28021 ttgaaattac acacaataac aaaacctgga acaatacctt atccaccaca tgggagccag  
 28081 gagttcccga gtggtacact gtctctgtcc gaggtcctga cggttccatc cgcattagta  
 28141 acaacacttt cattttttct gaaatgtgcy atctggccat gttcatgacy aaacagtatt  
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 28321 aaaacgcaa taacaaagaa aaaatgcctt aacctcttcc tgtttacaga catggcttct  
 28381 cttacatctc tcatatttgt cagcattgtc actgcccgtc acggacaaac agtctgtctc  
 28441 atcccactag gacataatta cactctcata ggaccccaa tcaactcaga ggtcatctgg  
 28501 accaaactgg gaagcgttga ttactttgat ataactgtga acaaaacaaa accaataata  
 28561 gtaacttgca acatacaaaa tcttacattg ataatgtta gcaaagtta cagcggttac  
 28621 tattatgggt atgacagata cagtagtcaa tatagaaatt acttggttcgy tgttaccag  
 28681 ttgaaaacca cgaaaatgcc aaatatggca aagattcgyt ccgatgacaa tctctagaa  
 28741 acttttacct ctcccaccac acccgacyaa aaaaacatcc cagattcaat gattgcaatt  
 28801 gttgcagcyg tggcagtggt gatggcacta ataataatat gcatgctttt atatgcttgt  
 28861 cgctacaaaa agtttcatcc taaaaacaa gatctcctac taaggcttaa catttaattt  
 28921 ctttttatac agccatgggt tccactacca cattccttat gcttactagt ctcgcaactc  
 28981 tgacttctgc tcgctcacac ctactgtaa ctataggctc aaactgcaca ctaaaaggac  
 29041 ctcaaggtgg tcatgtcttt tggtaggaa tatatgacaa tggatggttt acaaaacct  
 29101 gtgaccaacc tggtagattt ttctgcaacg gcagagacct aaccattatc aacgtgacy  
 29161 caaatgacaa aggttctat tatggaaccg actataaaag tagtttagat tataacatta  
 29221 ttgtactgcc atctaccact ccagcaccct gcacaactac tttctctacy agcagtgctc  
 29281 ctaacaatac aatttccaat ccaaccttgc ccgcyctttt aaaacgcact gtgaataatt  
 29341 ctacaacttc acatacaaca atttccactt caacaatcag catcatcgyt gcagtgacaa  
 29401 ttggaatata tattcttgtt ttaccataa cctactacyc ctgctgctat agaaaagaca  
 29461 aacataaagg tgatccatta cttagatttg atatttaatt tgttcttttt ttttatttac  
 29521 agtatgggtga acaccaatca tggtagctag aaatttcttc ttcaccatac tcatctgtgc  
 29581 ttttaaatgt tgcyctactt tcacagcagc agccacagca accccagact gtataggacy  
 29641 atttgcttcc tatgcacttt ttgcttttgt tacttgcacy tgcgtatgta gcatagctc  
 29701 cctggttatt aatttttcc aacttctaga ctggatcctt gtcgcaattg cctacctcgy

FIG. 11A-8



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29761 ccaccatccc gaataaccgca accaaaatat cgcggcactt cttagactca tctaaaacca  
 29821 tgcaggctat actaccaata tttttgcttc tattgcttcc ctacgctgtc tcaaccccag  
 29881 ctgcctatag tactccacca gaacacctta gaaaatgcaa attccaacaa ccgtgggtcat  
 29941 ttcttgcttg ctatcgagaa aatcagaaa tcccccaaa ttaataatg attgctggaa  
 30001 taattaatat aatctgttgc accataattt catttttgat atacccccta tttgattttg  
 30061 gctggaatgc tcccaatgca catgatcatc cacaagacce agaggaacac attccccac  
 30121 aaaacatgca acatccaata gcgctaatag attacgaaag tgaaccacaa cccccactac  
 30181 tccctgctat tagttacttc aacctaacgg gcggagatga ctgaaacact caccacctcc  
 30241 aattccgccc aggatctgct cgatatggac ggccgcgtct cagaacaacg acttgcccaa  
 30301 ctacgcatcc gccagcagca ggaacgcgtg gccaaagagc tcagagatgt catccaaatt  
 30361 caccaatgca aaaaaggcat attctgtttg gtaaaacaag ccaagatate ctacgagatc  
 30421 accgctactg accatcgctt ctcttacgaa cttggccccc aacgacaaaa atttacctgc  
 30481 atgggtgggaa tcaaccccat agttatcacc caacaaagtg gagatactaa ggggttgcatt  
 30541 cactgctcct gcgattccat cgagtgcacc tacaccctgc tgaagaccct atgcggccta  
 30601 agagacctgc taccaatgaa ttaaaaaaaa atgattaata aaaaatcact tacttgaat  
 30661 cagcaataag gtctctgttg aaattttctc ccagcagcac ctcacttccc tcttcccaac  
 30721 tctggtattc taaaccccg tccagcggcat actttctcca tactttaaag gggatgtcaa  
 30781 attttagctc ctctcctgta cccacaatct tcatgtcttt cttcccagat gaccaagaga  
 30841 gtccggctca gtgactcctt caaccctgtc taccctatg aagatgaaag cacctcccaa  
 30901 cacccttta taaaccagg gtttatttcc ccaatggct tcacacaaag cccagacgga  
 30961 gttcttactt taaaatgttt aaccccacta acaaccacag gcggatctct acagctaaaa  
 31021 gtgggagggg gacttacagt ggatgacact gatggtacct tacaagaaaa catacgtgct  
 31081 acagcaccca ttactaaaaa taatcactct gtagaactat ccattggaaa tggattagaa  
 31141 actcaaaaaca ataaactatg tgccaaattg ggaaatgggt taaaatttaa caacggtgac  
 31201 atttgtataa aggatagtat taacacctta tggactggaa taaaccctcc acctaactgt  
 31261 caaattgtgg aaaacactaa tacaatgat ggcaaactta ctttagtatt agtaaaaaat  
 31321 ggagggcttg ttaatggcta cgtgtctcta gttgggtgat cagacactgt gaaccaaatg  
 31381 ttcacacaaa agacagcaaa catccaatta agattatatt ttgactcttc tggaaatcta  
 31441 ttaactgagg aatcagactt aaaaattcca cttaaaaata aatcttctac agcgaccagt  
 31501 gaaactgtag ccagcagcaa agcctttatg ccaagtacta cagcttatcc cttcaacacc  
 31561 actactaggg atagtgaaaa ctacattcat ggaatatgtt actacatgac tagttatgat  
 31621 agaagtctat ttcccttgaa catttctata atgctaaaca gccgtatgat ttcttccaat  
 31681 gttgcctatg ccatacaatt tgaatggaat ctaaatgcaa gtgaatctcc agaaagcaac  
 31741 atagctacgc tgaccacatc cccctttttc ttttcttaca ttacagaaga cgacaactaa  
 31801 aataaagttt aagtgttttt atttaaaatc acaaaattcg agtagttatt ttgcctccac  
 31861 ctcccattt gacagaatac accaatctct ccccacgcac agctttaaac atttggatac  
 31921 cattagagat agacattggt ttagattcca cattccaac agtttcagag cgagccaatc  
 31981 tggggtcagt gatagataaa aatccatcgc gatagtcttt taaagcgctt tcacagtcca  
 32041 actgctgcgg atgcgactcc ggagtttggg tcacgggtcat ctggaagaag aacgatggga  
 32101 atcataatcc gaaaacggta tccggacgatt gtgtctcatc aaaccacaa gcagccgctg  
 32161 tctgcgtcgc tccgtgcgac tgctgtttat gggatcaggg tccacagttt cctgaagcat  
 32221 gattttaata gcccttaaca tcaactttct ggtgcgatgc gcgcagcaac gcattctgat  
 32281 ttcactcaaa tctttgcagt aggtacaaca cattattaca atattgttta ataaaccata  
 32341 attaaaagcg ctccagccaa aactcatatc tgatataatc gccctgcat gaccatcata  
 32401 ccaaagttta atataaatta aatgacgttc cctcaaaaac aactaccca catacatgat  
 32461 ctcttttggc atgtgcatat taacaatctg tctgtaccat ggacaacggt ggttaatcat  
 32521 gcaacccaat ataaccttcc ggaaccacac tgccaacacc gctccccag ccatgcattg  
 32581 aagtgaacc tgctgattac aatgacaatg aagaacccaa ttctctcgac cgtgaatcac  
 32641 ttgagaatga aaaatatcta tagtggcaca acatagacat aatgcatgc atcttctcat  
 32701 aatttttaac tcttcaggat ttagaaacat atcccaggga ataggaagct cttgcagaac  
 32761 agtaaagctg gcagaacaag gaagaccacg aacacaactt aactatgca tagtcatagt  
 32821 atcacaatct ggcaacagcg ggtgggtctt agtcatagaa gctcgggttt cattttctc  
 32881 acaacgtggg aactgggctc tgggtgtaagg gtgatgtctg gcgcatgatg tcgagcgtgc  
 32941 gcgcaacctt gtcataatgg agttgcttcc tgacattctc gtattttgta tagcaaaacg  
 33001 cggccctggc agaacacact cttcttcgcc ttctatcctg ccgcttagcg tgttccgtgt  
 33061 gatagttcaa gtacagccac actcttaagt tggcaaaag aatgctggct tcagttgtaa  
 33121 tcaaaactcc atcgcatcta attgttctga ggaaatcatc cacggtagca tatgcaaatc  
 33181 ccaaccaagc aatgcaactg gattgcgttt caagcaggag aggagaggga agagacggaa  
 33241 gaacatggtt aatttttatt ccaaacgatc tcgcagtact tcaaattgta gatcgcgcag  
 33301 atggcatctc tcgccccac tgtgttgggtg aaaaagcaca gctaaatcaa aagaaatgcg  
 33361 attttcaagg tgctcaacgg tggcttccaa caaagcctcc acgcgcatat ccaagaacaa  
 33421 aagaatacca aaagaaggag cattttctaa ctctcaatc atcatattac attcctgcac

FIG. 11A-9



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33481 cattcccaga taatthttcag ctttccagcc ttgaattatt cgtgtcagtt cttgtggtaa  
33541 atccaatcca cacattacaa acaggtcccg gagggcgccc tccaccacca ttcttaaaca  
33601 caccctcata atgacaaaat atcttgctcc tgtgtcacct gtagcgaatt gagaatggca  
33661 acatcaattg acatgccctt ggctctaagt tcttctttaa gttctagttg taaaaactct  
33721 ctcatattat caccaaactg cttagccaga agcccccgga gaacaagagc aggggacgct  
33781 acagtgcagt acaagcgcag acctcccaa ttggctccag caaaaacaag attggaataa  
33841 gcatattggg aaccaccagt aatatcatcg aagttgctgg aaatataatc aggcagagtt  
33901 tcttgtagaa attgaataaa agaaaaattt gccaaaaaaa cattcaaac ctctgggatg  
33961 caaatgcaat aggttaccgc gctgcgctcc aacattgtta gttttgaatt agtctgcaaa  
34021 aataaaaaaa aaacaagcgt catatcatag tagcctgacg aacaggtgga taaatcagtc  
34081 tttccatcac aagacaagcc acagggctctc cagctcgacc ctcgtaaac ctgtcatcgt  
34141 gattaaacaa cagcaccgaa agttcctcgc ggtgaccagc atgaataagt cttgatgaag  
34201 catacaatcc agacatgtta gcatcagtta aggagaaaaa acagccaaca tagcctttgg  
34261 gtataattat gcttaatcgt aagtatagca aagccacccc tcgcgatac aaagtaaaag  
34321 gcacaggaga ataaaaaata taattatthc tctgctgctg tttaggcaac gtcgcccccg  
34381 gtcctctaa atacacatac aaagcctcat cagccatggc ttaccagaga aagtacagcg  
34441 ggcacacaaa ccacaagctc taaagtcact ctccaacctc tccacaatat atatacacia  
34501 gccctaaact gacgtaatgg gactaaagtg taaaaaatcc cgccaacccc aacacacacc  
34561 ccgaaactgc gtcaccaggg aaaagtacag tttcacttcc gcaatcccaa caagcgtcac  
34621 ttctctttc tcacggtagc tcacatccca ttaacttaca acgtcatttt cccacggccg  
34681 cgccgccctt tttaacggtt aaccccacag ccaatcacca cacggcccac actttttaa  
34741 atcacctcat ttacatattg gcaccattcc atctataagg tatattattg atgatg

FIG. 11A-10



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Vaccine T=0, 4 wks	Vaccine T=24 wks	Monkey ID	Pre		T=4 wks		T=8 wks		T=24 wks		T=28 wks		T=32 wks	
			Mock	Gag <sup>a</sup>	Mock	Gag	Mock	Gag	Mock	Gag	Mock	Gag	Mock	Gag
Ad34ΔE1gagΔE4Ad5Orf6, 10 <sup>11</sup> vp	Ad35ΔE1gagΔE4Ad5Orf6, 10 <sup>10</sup> vp	00D016	4	6	1	84	5	334	5	99	0	306	3	244
Ad34ΔE1gagΔE4Ad5Orf6, 10 <sup>11</sup> vp	Ad35ΔE1gagΔE4Ad5Orf6, 10 <sup>10</sup> vp	00D044	1	1	8	79	0	374	8	136	0	493	1	253
Ad34ΔE1gagΔE4Ad5Orf6, 10 <sup>11</sup> vp	Ad35ΔE1gagΔE4Ad5Orf6, 10 <sup>10</sup> vp	00D064	4	6	1	125	8	655	6	145	0	351	1	236
Naïve		00D087	1	1	3	3	8	54	6	8	5	5	3	0

FIG. 12



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Vaccine (T=0, 4 Wks)	Vaccine (T=24 Wk)	Monkey ID	IFN- $\gamma$ <sup>+</sup> CD4 <sup>+</sup> CD3 <sup>+</sup> per 10 <sup>6</sup> Lymphocytes		IFN- $\gamma$ <sup>+</sup> CD8 <sup>+</sup> CD3 <sup>+</sup> per 10 <sup>6</sup> Lymphocytes	
			Mock	Gag	Mock	Gag
Ad34 $\Delta$ E1gag $\Delta$ E4Ad5Orf6, 10 <sup>11</sup> vp	Ad35 $\Delta$ E1gag $\Delta$ E4Ad5Orf6, 10 <sup>10</sup> vp	00D016	62	433	176	1288
Ad34 $\Delta$ E1gag $\Delta$ E4Ad5Orf6, 10 <sup>11</sup> vp	Ad35 $\Delta$ E1gag $\Delta$ E4Ad5Orf6, 10 <sup>10</sup> vp	00D044	136	593	323	1871
Ad34 $\Delta$ E1gag $\Delta$ E4Ad5Orf6, 10 <sup>11</sup> vp	Ad35 $\Delta$ E1gag $\Delta$ E4Ad5Orf6, 10 <sup>10</sup> vp	00D064	188	785	292	992

FIG. 13