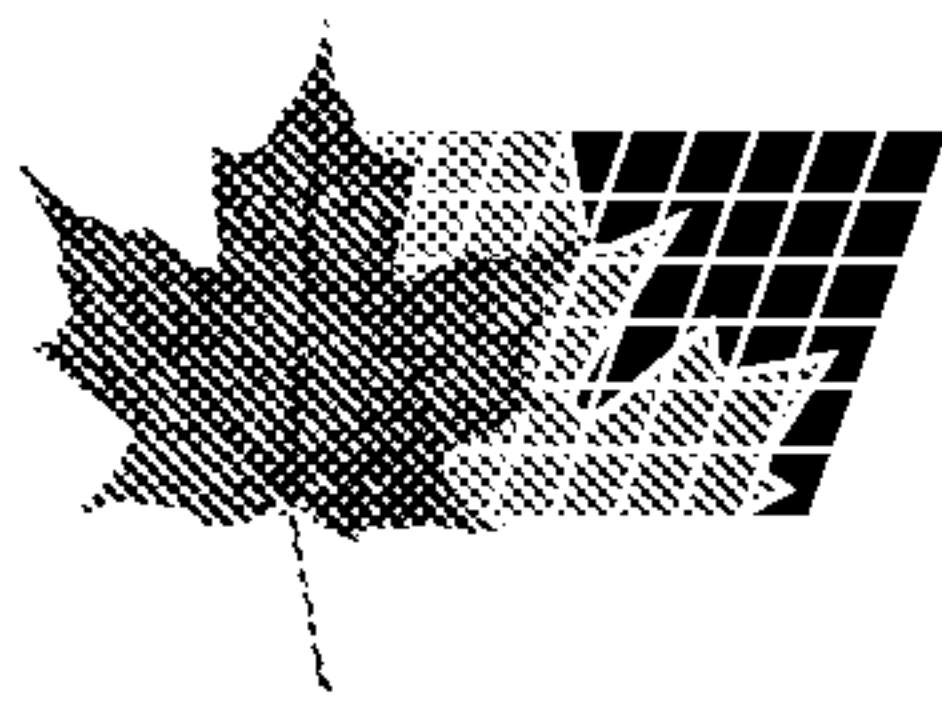




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(54) **PROTEINES RESSEMBLANT AU FACTEUR H DU
COMPLEMENT HUMAIN ET ADN_C CODANT CES
PROTEINES**

(54) **PROTEINS ALIKE TO HUMAN COMPLEMENT FACTOR H
AND cDNAs ENCODING THESE PROTEINS**

(57) La présente invention se rapporte à de l'ADN_C codant pour des protéines semblables au facteur H du complément humain, ainsi qu'aux protéines codées par cet ADN_C humain. L'ADN_C humain de la présente invention peut être utilisé en tant que sonde moléculaire pour le diagnostic génétique et en tant que sources de gènes pour la thérapie génique.

(57) The present invention provides human cDNAs coding for proteins alike to human complement factor H and proteins encoded by these human cDNAs. The human cDNAs of the present invention can be utilized as probes for the gene diagnosis and gene sources for the gene therapy.

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(54) Title: PROTEINS ALIKE TO HUMAN COMPLEMENT FACTOR H AND cDNAs ENCODING THESE PROTEINS (57) Abstract The present invention provides human cDNAs coding for proteins alike to human complement factor H and proteins encoded by these human cDNAs. The human cDNAs of the present invention can be utilized as probes for the gene diagnosis and gene sources for the gene therapy.		

DESCRIPTION

Proteins Alike to Human Complement
Factor H and cDNAs Encoding these Proteins

5

FIELD OF THE INVENTION

The present invention relates to proteins alike to human complement factor H and cDNAs coding for these proteins. The proteins of the present invention can be employed as pharmaceuticals. The human cDNAs of the present invention can be utilized as probes for the gene diagnosis and gene sources for the gene therapy. Furthermore, the cDNAs can be utilized as gene sources for large-scale production of the proteins encoded by said cDNAs. Eucaryotic cells wherein expression vectors of said cDNAs are introduced can be utilized for secretory production of the proteins alike to human complement factor H.

BACKGROUND ART

Human complement factor H is one of glycoproteins that are abundantly contained in the serum and are involved in the regulation of enzymatic processing of the major complement component C3. Cloning of the cDNA of the human complement factor H, followed by analysis of its amino acid sequence, has revealed that this protein has 20 short consensus repeats (SCR), each consisting of about 60 amino acid residues [Ripoche, J. et al., Biochem. J. 29: 593-602 (1988)]. Recently, several cDNAs of proteins, which are related to complement factor H and possess plural SCRs, have been reported and it has been recognized that

these proteins are organized as one family [Estaller, C. et al., J. Immunol. 146: 3190-3196 (1991); Skerka, C. et al., J. Biol. Chem. 266: 12015-12020 (1991); Skerka, C. et al., J. Immunol. 148: 3313-3318 (1992); Skerka, C. et al., J. Biol. Chem. 268: 2904-2908
5 (1993). These proteins related to complement factor H have been considered to play an important role in the immunological mechanism involving the compliment reaction.

DISCLOSURE OF INVENTION

10 The object of the present invention is to provide novel proteins alike to human complement factor H and cDNAs coding for these proteins.

As the result of intensive studies, the present inventors have been successful in cloning of human cDNAs coding for proteins
15 alike to human complement factor H, thereby completing the present invention. In other words, the present invention provides proteins alike to human complement factor H, namely proteins containing the amino acid sequence represented by Sequence No. 1. Moreover, the present invention provides cDNAs coding for the above-
20 mentioned proteins and containing the base sequence represented by Sequence No. 2 or Sequence No. 3 as well as transformation eucaryotic cells that are capable of expressing said cDNAs.

BRIEF DESCRIPTION OF DRAWINGS

25 Figure 1: A figure depicting the hydrophobicity/hydrophilicity profile of the protein encoded by clone HP01265.

Figure 2: A figure illustrating the results of the dot matrix analysis of the amino acid sequence of the protein encoded by clone HP01265.

5 BEST MODE FOR CARRYING OUT THE INVENTION

The proteins of the present invention can be obtained, for example, by a method for isolation from human organs, cell lines, etc., a method for preparation of peptides by the chemical synthesis, or a method for production with the recombinant DNA
10 technology using the DNAs coding for the proteins alike to human complement factor H of the present invention, wherein the method for obtainment by the recombinant DNA technology is employed preferably. For instance, in vitro expression can be achieved by preparation of an RNA by the in vitro transcription from a vector
15 having one of cDNAs of the present invention, followed by in vitro translation using this RNA as a template. Also, recombination of the translation region into a suitable expression vector by the method known in the art leads to expression of a large amount of the encoded protein by using prokaryotic cells such as
20 *Escherichia coli*, *Bacillus subtilis*, etc., and eucaryotic cells such as yeasts, insect cells, mammalian cells, etc.

In the case in which a protein of the present invention is expressed by prokaryotic cells such as *Escherichia coli* etc., a recombinant expression vector bearing the translation region
25 in the cDNA of the present invention is constructed in an expression vector having an origin, a promoter, a ribosome-binding site, a cDNA-cloning site, a terminator etc., which can be replicated in

the prokaryotic cells, and, after transformation of the host cells with said expression vector, the thus-obtained transformant is incubated, whereby the protein encoded by said cDNA can be produced on a large scale in the prokaryotic cells. Alternatively, a fusion
5 protein with another protein can be expressed. Only a protein portion coding for said cDNA can be obtained by cleavage of said fusion protein with an appropriate protease. Said fusion protein, provided that it possesses the activity alike to human complement factor H, shall come within the scope of the present invention.

10 In the case in which a protein of the present invention is produced in eucaryotic cells, the protein of the present invention can be produced by extracellular secretion, when the translation region of said cDNA is subjected to recombination to an expression vector for eucaryotic cells that has a promoter,
15 a splicing region, a poly(A) addition site, etc., followed by introduction into the eucaryotic cells. The expression vector is exemplified by pKA1, pED6dpc2, pCDM8, pSVK3, pMSG, pSVL, pBK-CMV, pBK-RSV, EBV vector, pRS, pYES2, and so on. Examples of eucaryotic cells to be used in general include mammalian culture cells such
20 as simian kidney cells COS7, Chinese hamster ovary cells CHO, etc., budding yeasts, fission yeasts, silkworm cells, *Xenopus laevis* egg cells, and so on, but any eucaryotic cells may be used, provided that they are capable of effecting secretory expression of the present proteins.

25 The proteins of the present invention include peptide fragments (more than 5 amino acid residues) containing any partial amino acid sequence in the amino acid sequence represented by

Sequence No. 1. These peptide fragments can be utilized as antigens for preparation of antibodies. Hereupon, the proteins of the present invention are secreted in an extracellular manner. Since a portion capable of binding sugar chains exists in the amino acid
5 sequence, proteins where sugar chains are added can be obtained by expression in appropriate eucaryotic cells. Accordingly, such proteins or peptides wherein sugar chains are added shall come within the scope of the present invention.

The DNAs of the present invention include all DNAs coding
10 for the above-mentioned proteins. Said DNAs can be obtained by using a method by chemical synthesis, a method by cDNA cloning, and so on.

The human cDNAs of the present invention can be cloned from cDNA libraries of the human cell origin. These cDNA libraries are
15 constructed by using as templates poly(A)⁺ RNAs extracted from human cells. The human cells may be cells delivered from the human body, for example, by the operation or may be the culture cells. A poly(A)⁺ RNA isolated from the liver is used in Examples. The cDNAs can be synthesized by using any method selected from the
20 Okayama-Berg method [Okayama, H. and Berg, P., Mol. Cell. Biol. 2: 161-170 (1982)], the Gubler-Hoffman method [Gubler, U. and Hoffman, J. Gene 25: 263-269 (1983)], and so on, but it is preferred to use the capping method [Kato, S. et al., Gene 150: 243-250 (1994)] in order to obtain a full-length clone in an effective
25 manner. The identification of the cDNAs is carried out by the determination of the whole base sequence by the sequencing, the search of known proteins having sequences analogous to the amino

acid sequence predicted from the base sequence, expression of proteins by in vitro translation, and expression by cultured cells.

The cDNAs of the present invention are characterized by
5 containing the base sequence represented by Sequence No. 2 or Sequence No. 3. For example, that represented by Sequence No. 3 possesses a 2033-bp base sequence with a 1737-bp open reading frame. This open reading frame codes for a protein consisting of 578 amino acid residues. This protein is partially identical with the
10 protein alike to human complement factor H in the amino acid sequence level.

Hereupon, the same clones as the cDNAs of the present invention can be easily obtained by screening of the human cDNA libraries constructed from the human cells by the use of an
15 oligonucleotide probe synthesized on the basis of the cDNA base sequence described in Sequence No. 3.

In general, the polymorphism due to the individual difference is frequently observed in human genes. Accordingly, any cDNA that is subjected to insertion or deletion of one or plural
20 nucleotides and/or substitution with other nucleotides in Sequence No. 2 or Sequence No. 3 shall come within the scope of the present invention.

In a similar manner, any protein that is formed by these modifications comprising insertion or deletion of one or plural
25 amino acids and/or substitution with other amino acids shall come within the scope of the present invention, as far as the protein possesses the activity of the protein alike to human complement

factor H.

The cDNAs of the present invention include cDNA fragments (more than 10 bp) containing any partial base sequence in the base sequence represented by Sequence No. 3. Also, DNA fragments
5 consisting of a sense chain and an anti-sense chain shall come within this scope. These DNA fragments can be utilized as the probes for the gene diagnosis.

In addition to the activities and uses described above, the polynucleotides and proteins of the present invention may
10 exhibit one or more of the uses or biological activities (including those associated with assays cited herein) identified below. Uses or activities described for proteins of the present invention may be provided by administration or use of such proteins or by
administration or use of polynucleotides encoding such proteins
15 (such as, for example, in gene therapies or vectors suitable for introduction of DNA).

Research Uses and Utilities

The polynucleotides provided by the present invention can be used by the research community for various purposes. The
20 polynucleotides can be used to express recombinant protein for analysis, characterization or therapeutic use; as markers for tissues in which the corresponding protein is preferentially expressed (either constitutively or at a particular stage of tissue differentiation or development or in disease states); as
25 molecular weight markers on Southern gels; as chromosome markers or tags (when labeled) to identify chromosomes or to map related gene positions; to compare with endogenous DNA sequences in

patients to identify potential genetic disorders; as probes to hybridize and thus discover novel, related DNA sequences; as a source of information to derive PCR primers for genetic fingerprinting; as a probe to "subtract-out" known sequences in the process of discovering other novel polynucleotides; for selecting and making oligomers for attachment to a "gene chip" or other support, including for examination of expression patterns; to raise anti-protein antibodies using DNA immunization techniques; and as an antigen to raise anti-DNA antibodies or elicit another immune response. Where the polynucleotide encodes a protein which binds or potentially binds to another protein (such as, for example, in a receptor-ligand interaction), the polynucleotide can also be used in interaction trap assays (such as, for example, that described in Gyuris et al., Cell 75:791-803 (1993)) to identify polynucleotides encoding the other protein with which binding occurs or to identify inhibitors of the binding interaction.

The proteins provided by the present invention can similarly be used in assay to determine biological activity, including in a panel of multiple proteins for high-throughput screening; to raise antibodies or to elicit another immune response; as a reagent (including the labeled reagent) in assays designed to quantitatively determine levels of the protein (or its receptor) in biological fluids; as markers for tissues in which the corresponding protein is preferentially expressed (either constitutively or at a particular stage of tissue differentiation or development or in a disease state); and, of course, to isolate

correlative receptors or ligands. Where the protein binds or potentially binds to another protein (such as, for example, in a receptor-ligand interaction), the protein can be used to identify the other protein with which binding occurs or to identify
5 inhibitors of the binding interaction. Proteins involved in these binding interactions can also be used to screen for peptide or small molecule inhibitors or agonists of the binding interaction.

Any or all of these research utilities are capable of being developed into reagent grade or kit format for commercialization
10 as research products.

Methods for performing the uses listed above are well known to those skilled in the art. References disclosing such methods include without limitation "Molecular Cloning: A Laboratory Manual", 2d ed., Cold Spring Harbor Laboratory Press, Sambrook,
15 J., E.F. Fritsch and T. Maniatis eds., 1989, and "Methods in Enzymology: Guide to Molecular Cloning Techniques", Academic Press, Berger, S.L. and A.R. Kimmel eds., 1987.

Nutritional Uses

Polynucleotides and proteins of the present invention can
20 also be used as nutritional sources or supplements. Such uses include without limitation use as a protein or amino acid supplement, use as a carbon source, use as a nitrogen source and use as a source of carbohydrate. In such cases the protein or polynucleotide of the invention can be added to the feed of a
25 particular organism or can be administered as a separate solid or liquid preparation, such as in the form of powder, pills, solutions, suspensions or capsules. In the case of microorganisms,

the protein or polynucleotide of the invention can be added to the medium in or on which the microorganism is cultured.

Cytokine and Cell Proliferation/Differentiation Activity

A protein of the present invention may exhibit cytokine,
5 cell proliferation (either inducing or inhibiting) or cell
differentiation (either inducing or inhibiting) activity or may
induce production of other cytokines in certain cell populations.
Many protein factors discovered to date, including all known
cytokines, have exhibited activity in one or more factor dependent
10 cell proliferation assays, and hence the assays serve as a
convenient confirmation of cytokine activity. The activity of
a protein of the present invention is evidenced by any one of a
number of routine factor dependent cell proliferation assays for
cell lines including, without limitation, 32D, DA2, DA1G, T10,
15 B9, B9/11, BaF3, MC9/G, M+ (preB M+), 2E8, RB5, DA1, 123, T1165,
HT2, CTLL2, TF-1, Mo7e and CMK.

The activity of a protein of the invention may, among other
means, be measured by the following methods:

Assays for T-cell or thymocyte proliferation include
20 without limitation those described in: Current Protocols in
Immunology, Ed by J. E. Coligan, A.M. Kruisbeek, D.H. Margulies,
E.M. Shevach, W Strober, Pub. Greene Publishing Associates and
Wiley-Interscience (Chapter 3, In Vitro assays for Mouse
Lymphocyte Function 3.1-3.19; Chapter 7, Immunologic studies in
25 Humans); Takai et al., J. Immunol. 137:3494-3500, 1986;
Bertagnolli et al., J. Immunol. 145:1706-1712, 1990; Bertagnolli
et al., Cellular Immunology 133:327-341, 1991; Bertagnolli, et

al., J. Immunol. 149:3778-3783, 1992; Bowman et al., J. Immunol. 152: 1756-1761, 1994.

Assays for cytokine production and/or proliferation of spleen cells, lymph node cells or thymocytes include, without
5 limitation, those described in: Polyclonal T cell stimulation, Kruisbeek, A.M. and Shevach, E.M. In Current Protocols in Immunology. J.E.e.a. Coligan eds. Vol 1 pp. 3.12.1-3.12.14, John Wiley and Sons, Toronto. 1994; and Measurement of mouse and human Interferon γ , Schreiber, R.D. In Current Protocols in Immunology.
10 J.E.e.a. Coligan eds. Vol 1 pp. 6.8.1-6.8.8, John Wiley and Sons, Toronto. 1994.

Assays for proliferation and differentiation of hematopoietic and lymphopoietic cells include, without limitation, those described in: Measurement of Human and Murine Interleukin
15 2 and Interleukin 4, Bottomly, K., Davis, L.S. and Lipsky, P.E. In Current Protocols in Immunology. J.E.e.a. Coligan eds. Vol 1 pp. 6.3.1-6.3.12, John Wiley and Sons, Toronto. 1991; deVries et al., J. Exp. Med. 173:1205-1211, 1991; Moreau et al., Nature 336:690-692, 1988; Greenberger et al., Proc. Natl. Acad. Sci.
20 U.S.A. 80:2931-2938, 1983; Measurement of mouse and human interleukin 6-Nordan, R. In Current Protocols in Immunology. J.E.e.a. Coligan eds. Vol 1 pp. 6.6.1-6.6.5, John Wiley and Sons, Toronto. 1991; Smith et al., Proc. Natl. Acad. Sci. U.S.A. 83:1857-1861, 1986; Measurement of human Interleukin 11 - Bennett,
25 F., Giannotti, J., Clark, S.C. and Turner, K. J. In Current Protocols in Immunology. J.E.e.a. Coligan eds. Vol 1 pp. 6.15.1 John Wiley and Sons, Toronto. 1991; Measurement of mouse and human

Interleukin 9 - Ciarletta, A., Giannotti, J., Clark, S.C. and Turner, K.J. In Current Protocols in Immunology. J.E.e.a. Coligan eds. Vol 1 pp. 6.13.1, John Wiley and Sons, Toronto. 1991.

Assays for T-cell clone responses to antigens (which will
5 identify, among others, proteins that affect APC-T cell interactions as well as direct T-cell effects by measuring proliferation and cytokine production) include, without limitation, those described in: Current Protocols in Immunology, Ed by J. E. Coligan, A.M. Kruisbeek, D.H. Margulies, E.M. Shevach,
10 W Strober, Pub. Greene Publishing Associates and Wiley-Interscience (Chapter 3, In Vitro assays for Mouse Lymphocyte Function; Chapter 6, Cytokines and their cellular receptors; Chapter 7, Immunologic studies in Humans); Weinberger et al., Proc. Natl. Acad. Sci. USA 77:6091-6095, 1980; Weinberger
15 et al., Eur. J. Immun. 11:405-411, 1981; Takai et al., J. Immunol. 137:3494-3500, 1986; Takai et al., J. Immunol. 140:508-512, 1988.

Immune Stimulating or Suppressing Activity

A protein of the present invention may also exhibit immune stimulating or immune suppressing activity, including without
20 limitation the activities for which assays are described herein. A protein may be useful in the treatment of various immune deficiencies and disorders (including severe combined immunodeficiency (SCID)), e.g., in regulating (up or down) growth and proliferation of T and/or B lymphocytes, as well as effecting
25 the cytolytic activity of NK cells and other cell populations. These immune deficiencies may be genetic or be caused by viral (e.g., HIV) as well as bacterial or fungal infections, or may result

from autoimmune disorders. More specifically, infectious diseases caused by viral, bacterial, fungal or other infection may be treatable using a protein of the present invention, including infections by HIV, hepatitis viruses, herpesviruses, mycobacteria, Leishmania spp., malaria spp. and various fungal infections such as candidiasis. Of course, in this regard, a protein of the present invention may also be useful where a boost to the immune system generally may be desirable, i.e., in the treatment of cancer.

Autoimmune disorders which may be treated using a protein of the present invention include, for example, connective tissue disease, multiple sclerosis, systemic lupus erythematosus, rheumatoid arthritis, autoimmune pulmonary inflammation, Guillain-Barre syndrome, autoimmune thyroiditis, insulin dependent diabetes mellitus, myasthenia gravis, graft-versus-host disease and autoimmune inflammatory eye disease. Such a protein of the present invention may also be useful in the treatment of allergic reactions and conditions, such as asthma (particularly allergic asthma) or other respiratory problems. Other conditions, in which immune suppression is desired (including, for example, organ transplantation), may also be treatable using a protein of the present invention.

Using the proteins of the invention it may also be possible to immune responses, in a number of ways. Down regulation may be in the form of inhibiting or blocking an immune response already in progress or may involve preventing the induction of an immune response. The functions of activated T cells may be inhibited

by suppressing T cell responses or by inducing specific tolerance in T cells, or both. Immunosuppression of T cell responses is generally an active, non-antigen-specific, process which requires continuous exposure of the T cells to the suppressive agent.

- 5 Tolerance, which involves inducing non-responsiveness or anergy in T cells, is distinguishable from immunosuppression in that it is generally antigen-specific and persists after exposure to the tolerizing agent has ceased. Operationally, tolerance can be demonstrated by the lack of a T cell response upon reexposure to
10 specific antigen in the absence of the tolerizing agent.

Down regulating or preventing one or more antigen functions (including without limitation B lymphocyte antigen functions (such as , for example, B7)), e.g., preventing high level lymphokine synthesis by activated T cells, will be useful in
15 situations of tissue, skin and organ transplantation and in graft-versus-host disease (GVHD). For example, blockage of T cell function should result in reduced tissue destruction in tissue transplantation. Typically, in tissue transplants, rejection of the transplant is initiated through its recognition as foreign
20 by T cells, followed by an immune reaction that destroys the transplant. The administration of a molecule which inhibits or blocks interaction of a B7 lymphocyte antigen with its natural ligand(s) on immune cells (such as a soluble, monomeric form of a peptide having B7-2 activity alone or in conjunction with a
25 monomeric form of a peptide having an activity of another B lymphocyte antigen (e.g., B7-1, B7-3) or blocking antibody), prior to transplantation can lead to the binding of the molecule to the

natural ligand(s) on the immune cells without transmitting the corresponding costimulatory signal. Blocking B lymphocyte antigen function in this matter prevents cytokine synthesis by immune cells, such as T cells, and thus acts as an immunosuppressant.

5 Moreover, the lack of costimulation may also be sufficient to anergize the T cells, thereby inducing tolerance in a subject. Induction of long-term tolerance by B lymphocyte antigen-blocking reagents may avoid the necessity of repeated administration of these blocking reagents. To achieve sufficient

10 immunosuppression or tolerance in a subject, it may also be necessary to block the function of a combination of B lymphocyte antigens.

The efficacy of particular blocking reagents in preventing organ transplant rejection or GVHD can be assessed using animal

15 models that are predictive of efficacy in humans. Examples of appropriate systems which can be used include allogeneic cardiac grafts in rats and xenogeneic pancreatic islet cell grafts in mice, both of which have been used to examine the immunosuppressive effects of CTLA4Ig fusion proteins in vivo as described in Lenschow

20 et al., Science 257:789-792 (1992) and Turka et al., Proc. Natl. Acad. Sci USA, 89:11102-11105 (1992). In addition, murine models of GVHD (see Paul ed., Fundamental Immunology, Raven Press, New York, 1989, pp. 846-847) can be used to determine the effect of blocking B lymphocyte antigen function in vivo on the development

25 of that disease.

Blocking antigen function may also be therapeutically useful for treating autoimmune diseases. Many autoimmune

disorders are the result of inappropriate activation of T cells that are reactive against self tissue and which promote the production of cytokines and autoantibodies involved in the pathology of the diseases. Preventing the activation of autoreactive T cells may reduce or eliminate disease symptoms. Administration of reagents which block costimulation of T cells by disrupting receptor:ligand interactions of B lymphocyte antigens can be used to inhibit T cell activation and prevent production of autoantibodies or T cell-derived cytokines which may be involved in the disease process. Additionally, blocking reagents may induce antigen-specific tolerance of autoreactive T cells which could lead to long-term relief from the disease. The efficacy of blocking reagents in preventing or alleviating autoimmune disorders can be determined using a number of well-characterized animal models of human autoimmune diseases. Examples include murine experimental autoimmune encephalitis, systemic lupus erythmatosis in MRL/lpr/lpr mice or NZB hybrid mice, murine autoimmune collagen arthritis, diabetes mellitus in NOD mice and BB rats, and murine experimental myasthenia gravis (see Paul ed., Fundamental Immunology, Raven Press, New York, 1989, pp. 840-856).

Upregulation of an antigen function (preferably a B lymphocyte antigen function), as a means of up regulating immune responses, may also be useful in therapy. Upregulation of immune responses may be in the form of enhancing an existing immune response or eliciting an initial immune response. For example, enhancing an immune response through stimulating B lymphocyte

antigen function may be useful in cases of viral infection. In addition, systemic viral diseases such as influenza, the commoncold, and encephalitis might be alleviated by the administration of stimulatory forms of B lymphocyte
5 antigens systemically.

Alternatively, anti-viral immune responses may be enhanced in an infected patient by removing T cells from the patient, costimulating the T cells in vitro with viral antigen-pulsed APCs either expressing a peptide of the present invention or together
10 with a stimulatory form of a soluble peptide of the present invention and reintroducing the in vitro activated T cells into the patient. Another method of enhancing anti-viral immune responses would be to isolate infected cells from a patient, transfect them with a nucleic acid encoding a protein of the present
15 invention as described herein such that the cells express all or a portion of the protein on their surface, and reintroduce the transfected cells into the patient. The infected cells would now be capable of delivering a costimulatory signal to, and thereby activate, T cells in vivo.

20 In another application, up regulation or enhancement of antigen function (preferably B lymphocyte antigen function) may be useful in the induction of tumor immunity. Tumor cells (e.g., sarcoma, melanoma, lymphoma, leukemia, neuroblastoma, carcinoma) transfected with a nucleic acid encoding at least one peptide of
25 the present invention can be administered to a subject to overcome tumor-specific tolerance in the subject. If desired, the tumor cell can be transfected to express a combination of peptides. For

example, tumor cells obtained from a patient can be transfected ex vivo with an expression vector directing the expression of a peptide having B7-2-like activity alone, or in conjunction with a peptide having B7-1-like activity and/or B7-3-like activity.

5 The transfected tumor cells are returned to the patient to result in expression of the peptides on the surface of the transfected cell. Alternatively, gene therapy techniques can be used to target a tumor cell for transfection in vivo.

The presence of the peptide of the present invention having

10 the activity of a B lymphocyte antigen(s) on the surface of the tumor cell provides the necessary costimulation signal to T cells to induce a T cell mediated immune response against the transfected tumor cells. In addition, tumor cells which lack MHC class I or MHC class II molecules, or which fail to reexpress sufficient

15 amounts of MHC class I or MHC class II molecules, can be transfected with nucleic acid encoding all or a portion of (e.g., a cytoplasmic-domain truncated portion) of an MHC class I α chain protein and β_2 microglobulin protein or an MHC class II α chain protein and an MHC class II β chain protein to thereby express MHC

20 class I or MHC class II proteins on the cell surface. Expression of the appropriate class I or class II MHC in conjunction with a peptide having the activity of a B lymphocyte antigen (e.g., B7-1, B7-2, B7-3) induces a T cell mediated immune response against the transfected tumor cell. Optionally, a gene encoding an

25 antisense construct which blocks expression of an MHC class II associated protein, such as the invariant chain, can also be cotransfected with a DNA encoding a peptide having the activity

of a B lymphocyte antigen to promote presentation of tumor associated antigens and induce tumor specific immunity. Thus, the induction of a T cell mediated immune response in a human subject may be sufficient to overcome tumor-specific tolerance
5 in the subject.

The activity of a protein of the invention may, among other means, be measured by the following methods:

Suitable assays for thymocyte or splenocyte cytotoxicity include, without limitation, those described in: Current
10 Protocols in Immunology, Ed by J. E. Coligan, A.M. Kruisbeek, D.H. Margulies, E.M. Shevach, W Strober, Pub. Greene Publishing Associates and Wiley-Interscience (Chapter 3, In Vitro assays for Mouse Lymphocyte Function 3.1-3.19; Chapter 7, Immunologic studies in Humans); Herrmann et al., Proc. Natl. Acad. Sci. USA
15 78:2488-2492, 1981; Herrmann et al., J. Immunol. 128:1968-1974, 1982; Handa et al., J. Immunol. 135:1564-1572, 1985; Takai et al., J. Immunol. 137:3494-3500, 1986; Takai et al., J. Immunol. 140:508-512, 1988; Herrmann et al., Proc. Natl. Acad. Sci. USA 78:2488-2492, 1981; Herrmann et al., J. Immunol. 128:1968-1974,
20 1982; Handa et al., J. Immunol. 135:1564-1572, 1985; Takai et al., J. Immunol. 137:3494-3500, 1986; Bowman et al., J. Virology 61:1992-1998; Takai et al., J. Immunol. 140:508-512, 1988; Bertagnolli et al., Cellular Immunology 133:327-341, 1991; Brown et al., J. Immunol. 153:3079-3092, 1994.

25 Assays for T-cell-dependent immunoglobulin responses and isotype switching (which will identify, among others, proteins that modulate T-cell dependent antibody responses and that affect

Th1/Th2 profiles) include, without limitation, those described in: Maliszewski, J. Immunol. 144:3028-3033, 1990; and Assays for B cell function: In vitro antibody production, Mond, J.J. and Brunswick, M. In Current Protocols in Immunology. J.E.e.a. Coligan
5 eds. Vol 1 pp. 3.8.1-3.8.16, John Wiley and Sons, Toronto. 1994.

Mixed lymphocyte reaction (MLR) assays (which will identify, among others, proteins that generate predominantly Th1 and CTL responses) include, without limitation, those described in: Current Protocols in Immunology, Ed by J. E. Coligan, A.M.
10 Kruisbeek, D.H. Margulies, E.M. Shevach, W Strober, Pub. Greene Publishing Associates and Wiley-Interscience (Chapter 3, In Vitro assays for Mouse Lymphocyte Function 3.1-3.19; Chapter 7, Immunologic studies in Humans); Takai et al., J. Immunol. 137:3494-3500, 1986; Takai et al., J. Immunol. 140:508-512, 1988;
15 Bertagnolli et al., J. Immunol. 149:3778-3783, 1992.

Dendritic cell-dependent assays (which will identify, among others, proteins expressed by dendritic cells that activate naive T-cells) include, without limitation, those described in: Guery et al., J. Immunol. 134:536-544, 1995; Inaba et al., Journal
20 of Experimental Medicine 173:549-559, 1991; Macatonia et al., Journal of Immunology 154:5071-5079, 1995; Porgador et al., Journal of Experimental Medicine 182:255-260, 1995; Nair et al., Journal of Virology 67:4062-4069, 1993; Huang et al., Science 264:961-965, 1994; Macatonia et al., Journal of Experimental
25 Medicine 169:1255-1264, 1989; Bhardwaj et al., Journal of Clinical Investigation 94:797-807, 1994; and Inaba et al., Journal of Experimental Medicine 172:631-640, 1990.

Assays for lymphocyte survival/apoptosis (which will identify, among others, proteins that prevent apoptosis after superantigen induction and proteins that regulate lymphocyte homeostasis) include, without limitation, those described in:

5 Darzynkiewicz et al., Cytometry 13:795-808, 1992; Gorczyca et al., Leukemia 7:659-670, 1993; Gorczyca et al., Cancer Research 53:1945-1951, 1993; Itoh et al., Cell 66:233-243, 1991; Zacharchuk, Journal of Immunology 145:4037-4045, 1990; Zamai et al., Cytometry 14:891-897, 1993; Gorczyca et al., International Journal of

10 Oncology 1:639-648, 1992.

Assays for proteins that influence early steps of T-cell commitment and development include, without limitation, those described in: Antica et al., Blood 84:111-117, 1994; Fine et al., Cellular Immunology 155:111-122, 1994; Galy et al., Blood

15 85:2770-2778, 1995; Toki et al., Proc. Nat. Acad Sci. USA 88:7548-7551, 1991.

Hematopoiesis Regulating Activity

A protein of the present invention may be useful in regulation of hematopoiesis and, consequently, in the treatment

20 of myeloid or lymphoid cell deficiencies. Even marginal biological activity in support of colony forming cells or of factor-dependent cell lines indicates involvement in regulating hematopoiesis, e.g. in supporting the growth and proliferation of erythroid progenitor cells alone or in combination with other

25 cytokines, thereby indicating utility, for example, in treating various anemias or for use in conjunction with irradiation/chemotherapy to stimulate the production of erythroid

precursors and/or erythroid cells; in supporting the growth and proliferation of myeloid cells such as granulocytes and monocytes/macrophages (i.e., traditional CSF activity) useful, for example, in conjunction with chemotherapy to prevent or treat
5 consequent myelo-suppression; in supporting the growth and proliferation of megakaryocytes and consequently of platelets thereby allowing prevention or treatment of various platelet disorders such as thrombocytopenia, and generally for use in place of or complimentary to platelet transfusions; and/or in supporting
10 the growth and proliferation of hematopoietic stem cells which are capable of maturing to any and all of the above-mentioned hematopoietic cells and therefore find therapeutic utility in various stem cell disorders (such as those usually treated with transplantation, including, without limitation, aplastic anemia
15 and paroxysmal nocturnal hemoglobinuria), as well as in repopulating the stem cell compartment post irradiation/chemotherapy, either in-vivo or ex-vivo (i.e., in conjunction with bone marrow transplantation or with peripheral progenitor cell transplantation (homologous or heterologous)) as
20 normal cells or genetically manipulated for gene therapy.

The activity of a protein of the invention may, among other means, be measured by the following methods:

Suitable assays for proliferation and differentiation of various hematopoietic lines are cited above.

25 Assays for embryonic stem cell differentiation (which will identify, among others, proteins that influence embryonic differentiation hematopoiesis) include, without limitation,

those described in: Johansson et al. Cellular Biology 15:141-151, 1995; Keller et al., Molecular and Cellular Biology 13:473-486, 1993; McClanahan et al., Blood 81:2903-2915, 1993.

Assays for stem cell survival and differentiation (which
5 will identify, among others, proteins that regulate lympho-
hematopoiesis) include, without limitation, those described in:
Methylcellulose colony forming assays, Freshney, M.G. In Culture
of Hematopoietic Cells. R.I. Freshney, et al. eds. Vol pp. 265-268,
Wiley-Liss, Inc., New York, NY. 1994; Hirayama et al., Proc. Natl.
10 Acad. Sci. USA 89:5907-5911, 1992; Primitive hematopoietic colony
forming cells with high proliferative potential, McNiece, I.K.
and Briddell, R.A. In Culture of Hematopoietic Cells. R.I.
Freshney, et al. eds. Vol pp. 23-39, Wiley-Liss, Inc., New York,
NY. 1994; Neben et al., Experimental Hematology 22:353-359, 1994;
15 Cobblestone area forming cell assay, Ploemacher, R.E. In Culture
of Hematopoietic Cells. R.I. Freshney, et al. eds. Vol pp. 1-21,
Wiley-Liss, Inc., New York, NY. 1994; Long term bone marrow
cultures in the presence of stromal cells, Spooncer, E., Dexter,
M. and Allen, T. In Culture of Hematopoietic Cells. R.I. Freshney,
20 et al. eds. Vol pp. 163-179, Wiley-Liss, Inc., New York, NY. 1994;
Long term culture initiating cell assay, Sutherland, H.J. In
Culture of Hematopoietic Cells. R.I. Freshney, et al. eds. Vol
pp. 139-162, Wiley-Liss, Inc., New York, NY. 1994.

Tissue Growth Activity

25 A protein of the present invention also may have utility
in compositions used for bone, cartilage, tendon, ligament and/or
nerve tissue growth or regeneration, as well as for wound healing

and tissue repair and replacement, and in the treatment of burns, incisions and ulcers.

A protein of the present invention, which induces cartilage and/or bone growth in circumstances where bone is not normally
5 formed, has application in the healing of bone fractures and cartilage damage or defects in humans and other animals. Such a preparation employing a protein of the invention may have prophylactic use in closed as well as open fracture reduction and also in the improved fixation of artificial joints. De novo bone
10 formation induced by an osteogenic agent contributes to the repair of congenital, trauma induced, or oncologic resection induced craniofacial defects, and also is useful in cosmetic plastic surgery.

A protein of this invention may also be used in the treatment
15 of periodontal disease, and in other tooth repair processes. Such agents may provide an environment to attract bone-forming cells, stimulate growth of bone-forming cells or induce differentiation of progenitors of bone-forming cells. A protein of the invention may also be useful in the treatment of osteoporosis or
20 osteoarthritis, such as through stimulation of bone and/or cartilage repair or by blocking inflammation or processes of tissue destruction (collagenase activity, osteoclast activity, etc.) mediated by inflammatory processes.

Another category of tissue regeneration activity that may
25 be attributable to the protein of the present invention is tendon/ligament formation. A protein of the present invention, which induces tendon/ligament-like tissue or other tissue

formation in circumstances where such tissue is not normally formed, has application in the healing of tendon or ligament tears, deformities and other tendon or ligament defects in humans and other animals. Such a preparation employing a tendon/ligament-like tissue inducing protein may have prophylactic use in preventing damage to tendon or ligament tissue, as well as use in the improved fixation of tendon or ligament to bone or other tissues, and in repairing defects to tendon or ligament tissue. De novo tendon/ligament-like tissue formation induced by a composition of the present invention contributes to the repair of congenital, trauma induced, or other tendon or ligament defects of other origin, and is also useful in cosmetic plastic surgery for attachment or repair of tendons or ligaments. The compositions of the present invention may provide an environment to attract tendon or ligament-forming cells, stimulate growth of tendon- or ligament-forming cells, induce differentiation of progenitors of tendon- or ligament-forming cells, or induce growth of tendon/ligament cells or progenitors ex vivo for return in vivo to effect tissue repair. The compositions of the invention may also be useful in the treatment of tendinitis, carpal tunnel syndrome and other tendon or ligament defects. The compositions may also include an appropriate matrix and/or sequestering agent as a carrier as is well known in the art.

25 The protein of the present invention may also be useful for proliferation of neural cells and for regeneration of nerve and brain tissue, i.e. for the treatment of central and peripheral

nervous system diseases and neuropathies, as well as mechanical and traumatic disorders, which involve degeneration, death or trauma to neural cells or nerve tissue. More specifically, a protein may be used in the treatment of diseases of the peripheral nervous system, such as peripheral nerve injuries, peripheral neuropathy and localized neuropathies, and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager syndrome. Further conditions which may be treated in accordance with the present invention include mechanical and traumatic disorders, such as spinal cord disorders, head trauma and cerebrovascular diseases such as stroke. Peripheral neuropathies resulting from chemotherapy or other medical therapies may also be treatable using a protein of the invention.

Proteins of the invention may also be useful to promote better or faster closure of non-healing wounds, including without limitation pressure ulcers, ulcers associated with vascular insufficiency, surgical and traumatic wounds, and the like.

It is expected that a protein of the present invention may also exhibit activity for generation or regeneration of other tissues, such as organs (including, for example, pancreas, liver, intestine, kidney, skin, endothelium), muscle (smooth, skeletal or cardiac) and vascular (including vascular endothelium) tissue, or for promoting the growth of cells comprising such tissues. Part of the desired effects may be by inhibition or modulation of fibrotic scarring to allow normal tissue to regenerate. A protein of the invention may also exhibit angiogenic activity.

A protein of the present invention may also be useful for gut protection or regeneration and treatment of lung or liver fibrosis, reperfusion injury in various tissues, and conditions resulting from systemic cytokine damage.

5 A protein of the present invention may also be useful for promoting or inhibiting differentiation of tissues described above from precursor tissues or cells; or for inhibiting the growth of tissues described above.

The activity of a protein of the invention may, among other
10 means, be measured by the following methods:

Assays for tissue generation activity include, without limitation, those described in: International Patent Publication No. WO95/16035 (bone, cartilage, tendon); International Patent Publication No. WO95/05846 (nerve, neuronal); International
15 Patent Publication No. WO91/07491 (skin, endothelium).

Assays for wound healing activity include, without limitation, those described in: Winter, Epidermal Wound Healing, pps. 71-112 (Maibach, HI and Rovee, DT, eds.), Year Book Medical Publishers, Inc., Chicago, as modified by Eaglstein and Mertz,
20 J. Invest. Dermatol 71:382-84 (1978).

Activin/Inhibin Activity

A protein of the present invention may also exhibit activin- or inhibin-related activities. Inhibins are characterized by their ability to inhibit the release of follicle stimulating
25 hormone (FSH), while activins and are characterized by their ability to stimulate the release of follicle stimulating hormone (FSH). Thus, a protein of the present invention, alone or in

heterodimers with a member of the inhibin α family, may be useful as a contraceptive based on the ability of inhibins to decrease fertility in female mammals and decrease spermatogenesis in male mammals. Administration of sufficient amounts of other inhibins
5 can induce infertility in these mammals. Alternatively, the protein of the invention, as a homodimer or as a heterodimer with other protein subunits of the inhibin- β group, may be useful as a fertility inducing therapeutic, based upon the ability of activin molecules in stimulating FSH release from cells of the
10 anterior pituitary. See, for example, United States Patent 4,798,885. A protein of the invention may also be useful for advancement of the onset of fertility in sexually immature mammals, so as to increase the lifetime reproductive performance of domestic animals such as cows, sheep and pigs.

15 The activity of a protein of the invention may, among other means, be measured by the following methods:

Assays for activin/inhibin activity include, without limitation, those described in: Vale et al., Endocrinology 91:562-572, 1972; Ling et al., Nature 321:779-782, 1986; Vale et
20 al., Nature 321:776-779, 1986; Mason et al., Nature 318:659-663, 1985; Forage et al., Proc. Natl. Acad. Sci. USA 83:3091-3095, 1986.

Chemotactic/Chemokinetic Activity

A protein of the present invention may have chemotactic or chemokinetic activity (e.g., act as a chemokine) for mammalian
25 cells, including, for example, monocytes, fibroblasts, neutrophils, T-cells, mast cells, eosinophils, epithelial and/or endothelial cells. Chemotactic and chemokinetic proteins can be

used to mobilize or attract a desired cell population to a desired site of action. Chemotactic or chemokinetic proteins provide particular advantages in treatment of wounds and other trauma to tissues, as well as in treatment of localized infections. For
5 example, attraction of lymphocytes, monocytes or neutrophils to tumors or sites of infection may result in improved immune responses against the tumor or infecting agent.

A protein or peptide has chemotactic activity for a particular cell population if it can stimulate, directly or
10 indirectly, the directed orientation or movement of such cell population. Preferably, the protein or peptide has the ability to directly stimulate directed movement of cells. Whether a particular protein has chemotactic activity for a population of cells can be readily determined by employing such protein or
15 peptide in any known assay for cell chemotaxis.

The activity of a protein of the invention may, among other means, be measured by the following methods:

Assays for chemotactic activity (which will identify proteins that induce or prevent chemotaxis) consist of assays that
20 measure the ability of a protein to induce the migration of cells across a membrane as well as the ability of a protein to induce the adhesion of one cell population to another cell population. Suitable assays for movement and adhesion include, without limitation, those described in: Current Protocols in Immunology,
25 Ed by J.E. Coligan, A.M. Kruisbeek, D.H. Margulies, E.M. Shevach, W.Strober, Pub. Greene Publishing Associates and Wiley-Interscience (Chapter 6.12, Measurement of alpha and beta

Chemokines 6.12.1-6.12.28; Taub et al. J. Clin. Invest. 95:1370-1376, 1995; Lind et al. APMIS 103:140-146, 1995; Muller et al Eur. J. Immunol. 25: 1744-1748; Gruber et al. J. of Immunol. 152:5860-5867, 1994; Johnston et al. J. of Immunol. 153: 1762-1768, 5 1994.

Hemostatic and Thrombolytic Activity

A protein of the invention may also exhibit hemostatic or thrombolytic activity. As a result, such a protein is expected to be useful in treatment of various coagulation disorders 10 (including hereditary disorders, such as hemophilias) or to enhance coagulation and other hemostatic events in treating wounds resulting from trauma, surgery or other causes. A protein of the invention may also be useful for dissolving or inhibiting formation of thromboses and for treatment and prevention of 15 conditions resulting therefrom (such as, for example, infarction of cardiac and central nervous system vessels (e.g., stroke).

The activity of a protein of the invention may, among other means, be measured by the following methods:

Assay for hemostatic and thrombolytic activity include, 20 without limitation, those described in: Linet et al., J. Clin. Pharmacol. 26:131-140, 1986; Burdick et al., Thrombosis Res. 45:413-419, 1987; Humphrey et al., Fibrinolysis 5:71-79 (1991); Schaub, Prostaglandins 35:467-474, 1988.

Receptor/Ligand Activity

25 A protein of the present invention may also demonstrate activity as receptors, receptor ligands or inhibitors or agonists of receptor/ligand interactions. Examples of such receptors and

ligands include, without limitation, cytokine receptors and their ligands, receptor kinases and their ligands, receptor phosphatases and their ligands, receptors involved in cell-cell interactions and their ligands (including without limitation, 5 cellular adhesion molecules (such as selectins, integrins and their ligands) and receptor/ligand pairs involved in antigen presentation, antigen recognition and development of cellular and humoral immune responses). Receptors and ligands are also useful for screening of potential peptide or small molecule inhibitors 10 of the relevant receptor/ligand interaction. A protein of the present invention (including, without limitation, fragments of receptors and ligands) may themselves be useful as inhibitors of receptor/ligand interactions.

The activity of a protein of the invention may, among other 15 means, be measured by the following methods:

Suitable assays for receptor-ligand activity include without limitation those described in: Current Protocols in Immunology, Ed by J.E. Coligan, A.M. Kruisbeek, D.H. Margulies, E.M. Shevach, W. Strober, Pub. Greene Publishing Associates and 20 Wiley-Interscience (Chapter 7.28, Measurement of Cellular Adhesion under static conditions 7.28.1-7.28.22), Takai et al., Proc. Natl. Acad. Sci. USA 84:6864-6868, 1987; Bierer et al., J. Exp. Med. 168:1145-1156, 1988; Rosenstein et al., J. Exp. Med. 169:149-160 1989; Stoltenborg et al., J. Immunol. Methods 25 175:59-68, 1994; Stitt et al., Cell 80:661-670, 1995.

Anti-Inflammatory Activity

Proteins of the present invention may also exhibit

anti-inflammatory activity. The anti-inflammatory activity may be achieved by providing a stimulus to cells involved in the inflammatory response, by inhibiting or promoting cell-cell interactions (such as, for example, cell adhesion), by inhibiting
5 or promoting chemotaxis of cells involved in the inflammatory process, inhibiting or promoting cell extravasation, or by stimulating or suppressing production of other factors which more directly inhibit or promote an inflammatory response. Proteins exhibiting such activities can be used to treat inflammatory
10 conditions including chronic or acute conditions), including without limitation inflammation associated with infection (such as septic shock, sepsis or systemic inflammatory response syndrome (SIRS)), ischemia-reperfusion injury, endotoxin lethality, arthritis, complement-mediated hyperacute rejection, nephritis,
15 cytokine or chemokine-induced lung injury, inflammatory bowel disease, Crohn's disease or resulting from over production of cytokines such as TNF or IL-1. Proteins of the invention may also be useful to treat anaphylaxis and hypersensitivity to an antigenic substance or material.

20 Tumor Inhibition Activity

In addition to the activities described above for immunological treatment or prevention of tumors, a protein of the invention may exhibit other anti-tumor activities. A protein may inhibit tumor growth directly or indirectly (such as, for example,
25 via ADCC). A protein may exhibit its tumor inhibitory activity by acting on tumor tissue or tumor precursor tissue, by inhibiting

formation of tissues necessary to support tumor growth (such as, for example, by inhibiting angiogenesis), by causing production of other factors, agents or cell types which inhibit tumor growth, or by suppressing, eliminating or inhibiting factors, agents or
5 cell types which promote tumor growth

Other Activities

A protein of the invention may also exhibit one or more of the following additional activities or effects: inhibiting the growth, infection or function of, or killing, infectious agents,
10 including, without limitation, bacteria, viruses, fungi and other parasites; effecting (suppressing or enhancing) bodily characteristics, including, without limitation, height, weight, hair color, eye color, skin, fat to lean ratio or other tissue pigmentation, or organ or body part size or shape (such as, for
15 example, breast augmentation or diminution, change in bone form or shape); effecting biorhythms or circadian cycles or rhythms; effecting the fertility of male or female subjects; effecting the metabolism, catabolism, anabolism, processing, utilization, storage or elimination of dietary fat, lipid, protein, carbohydrate,
20 vitamins, minerals, cofactors or other nutritional factors or component(s); effecting behavioral characteristics, including, without limitation, appetite, libido, stress, cognition (including cognitive disorders), depression (including depressive disorders) and violent behaviors; providing analgesic effects or other pain
25 reducing effects; promoting differentiation and growth of embryonic stem cells in lineages other than hematopoietic lineages;

hormonal or endocrine activity; in the case of enzymes, correcting deficiencies of the enzyme and treating deficiency-related diseases; treatment of hyperproliferative disorders (such as, for example, psoriasis); immunoglobulin-like activity (such as, for example, the ability to bind antigens or complement); and the ability to act as an antigen in a vaccine composition to raise an immune response against such protein or another material or entity which is cross-reactive with such protein.

10 Examples

The present invention is embodied in more detail by the following examples, but this embodiment is not intended to restrict the present invention. The basic operations and the enzyme reactions with regard to the DNA recombination are carried out according to the literature ["Molecular Cloning. A Laboratory Manual", Cold Spring Harbor Laboratory, 1989]. Unless otherwise stated, restrictive enzymes and a variety of modification enzymes to be used were those available from TAKARA SHUZO. The manufacturer's instructions were used for the buffer compositions as well as for the reaction conditions, in each of the enzyme reactions. The cDNA synthesis was carried out according to the literature [Kato, S. et al., Gene 150: 243-250 (1994)].

(1) Preparation of Poly(A)⁺ RNA

After about 1 g of the liver extracted by an operation was homogenized in 20 ml of a 5.5 M guanidinium thiocyanate solution, total mRNA was prepared according to the literature [Okayama, H. et al., "Method in Enzymology", Vol. 164, Academic Press, 1987].

This was subjected to chromatography on oligo(dT)-cellulose column washed with a 20 mM Tris-hydrochloride buffer solution (pH 7.6), 0.5 M NaCl, and 1 mM EDTA to obtain a poly(A)⁺ RNA according to the above-described literature.

5 (2) Construction of cDNA Library

Ten micrograms of the above-mentioned poly(A)⁺ RNA were dissolved in a 100 mM Tris-hydrochloride buffer solution (pH 8), one unit of an RNase-free, bacterial alkaline phosphatase was added, and the reaction was run at 37°C for one hour. After the
10 reaction solution was subjected to phenol extraction, followed by ethanol precipitation, the resulting pellet was dissolved in a solution containing 50 mM sodium acetate (pH 6), 1 mM EDTA, 0.1% 2-mercaptoethanol, and 0.01% Triton X-100. Thereto was added one unit of a tobacco-origin acid pyrophosphatase (Epicentre
15 Technologies) and a total 100 µl volume of the resulting mixture was reacted at 37°C for one hour. After the reaction solution was subjected to phenol extraction, followed by ethanol precipitation, the resulting pellet was dissolved in water to obtain a solution of a decapped poly(A)⁺ RNA.

20 The decapped poly(A)⁺ RNA and 3 nmol of a chimeric DNA-RNA oligonucleotide (5'-dG-dG-dG-dG-dA-dA-dT-dT-dC-dG-dA-G-G-A-3') were dissolved in a solution containing 50 mM Tris-hydrochloride buffer solution (pH 7.5), 0.5 mM ATP, 5 mM MgCl₂, 10 mM 2-mercaptoethanol, and 25% polyethylene glycol, whereto was
25 added 50 units of T4RNA ligase and a total 30 µl volume of the resulting mixture was reacted at 20°C for 12 hours. After the reaction solution was subjected to phenol extraction, followed

by ethanol precipitation, the resulting pellet was dissolved in water to obtain a chimeric-oligo-capped poly(A)⁺ RNA.

After digestion of vector pKA1 (Japanese Patent Kokai Publication No. 1992-117292) developed by the present inventors
5 with KpnI, about 60 dT tails were added using a terminal transferase. A vector primer to be used below was prepared by digestion of this product with EcoRV to remove a dT tail at one side.

After 6 µg of the previously-prepared chimeric-oligo-capped poly(A)⁺ RNA was annealed with 1.2 µg of the vector primer,
10 the resulting product was dissolved in a solution containing 50 mM Tris-hydrochloride buffer solution (pH 8.3), 75 mM KCl, 3 mM MgCl₂, 10 mM dithiothreitol, and 1.25 mM dNTP (dATP + dCTP + dGTP + dTTP), 200 units of a reverse transcriptase (GIBCO-BRL) were added, and the reaction in a total 20 µl volume was run at 42°C
15 for one hour. After the reaction solution was subjected to phenol extraction, followed by ethanol precipitation, the resulting pellet was dissolved in a solution containing 50 mM Tris-hydrochloride buffer solution (pH 7.5), 100 mM NaCl, 10 mM MgCl₂, and 1 mM dithiothreitol. Thereto were added 100 units of EcoRI
20 and a total 20 µl volume of the resulting mixture was reacted at 37°C for one hour. After the reaction solution was subjected to phenol extraction, followed by ethanol precipitation, the resulting pellet was dissolved in a solution containing 20 mM Tris-hydrochloride buffer solution (pH 7.5), 100 mM KCl, 4 mM MgCl₂,
25 10 mM (NH₄)₂SO₄, and 50 µg/ml of the bovine serum albumin. Thereto were added 60 units of an *Escherichia coli* DNA ligase and the resulting mixture was reacted at 16°C for 16 hours. To the reaction

solution were added 2 μ l of 2 mM dNTP, 4 units of *Escherichia coli* DNA polymerase I, and 0.1 unit of *Escherichia coli* RNase H and the resulting mixture was reacted at 12°C for one hour and then at 22°C for one hour.

5 Next, the cDNA-synthesis reaction solution was used for transformation of *Escherichia coli* DH12S (GIBCO-BRL). The transformation was carried out by the electroporation method. A portion of the transformant was sprayed on the 2xYT agar culture medium containing 100 μ g/ml ampicillin and the mixture was
10 incubated at 37°C overnight. A colony formed on the agar medium was picked up at random and inoculated on 2 ml of the 2xYT culture medium containing 100 μ g/ml ampicillin. After incubation at 37°C overnight, the culture mixture was centrifuged to separate the mycelia, from which a plasmid DNA was prepared by the alkaline
15 lysis method. The plasmid DNA was subjected to double digestion with EcoRI and NotI, followed by 0.8% agarose gel electrophoresis, to determine the size of the cDNA insert. Furthermore, using the thus-obtained plasmid as a template, the sequence reaction was carried out by using an M13 universal primer labeled with a
20 fluorescent dye and a Taq polymerase (a kit of Applied Biosystems) and then the product was examined with a fluorescent DNA sequencer (Applied Biosystems) to determine an about 400-bp base sequence at the 5'-terminus of the cDNA. The sequence data were filed as the homo/protein cDNA bank database.

25 (3) cDNA Cloning

Clone HP01265 was obtained as the result of a large-scale sequencing of cDNA clones selected from the above-mentioned cDNA

library. The present clone has a structure consisting of a 77-bp 5'-nontranslation region, a 1737-bp open reading frame, and a 219-bp 3'-nontranslation region (Sequence No. 3). The open reading frame codes for a protein consisting of 578 amino acid residues and there existed in the N-terminus a hydrophobic region that is considered to be a secretory signal sequence. Figure 1 depicts the hydrophobicity/hydrophilicity profile, obtained by the Kyte-Doolittle method, of this protein of the present invention.

The dot matrix analysis of the thus-obtained amino acid sequence revealed that two highly analogous domains are connected in a direct manner in the present protein, as shown in Figure 2. Table 1 shows the comparison between the two domain sequences. Therein, the marks of -, *, and . represent a gap, an amino acid residue identical with the protein of the present invention, and an amino acid residue analogous to the protein of the present invention, respectively. A sequence consisting of 245 residues from lysine at position 22 to isoleucine at position 266 had an 89.8% analogy to a sequence from lysine at position 269 to isoleucine at position 513. Furthermore, examination of the dot matrix has revealed the presence of 9 repeated sequences in the entire region except for the N-terminus. It has been revealed that the unit in the repeated sequences consists of a sequence of about 60 residues containing 4 molecules of cystine and is the SCR characteristic of the protein relating to human complement factor H.

Table 1

1'	MLLLINVILTLWVSCANGQEVKPCDFPEIQHGGLYYKSLRRLYFPAAAGQSYS
	****.***** **.. ** ***,*.*****
5	241" QSYVELQGSKYVTCSNGDWSEPPRCISMKPCEFPEIQHGHLYYENTRRPYFPVATGQSYS
	54' YYCDQNFVTPSGSYWDYIHCTQDGWSPTVPCLRTCSKSDVEIENGFI SESSSIYILNEET
	*****.*****.*****.*****.*.
	301" YYCDQNFVTPSGSYWDYIHCTQDGLPTVPCLRTCSKSDIEIENGFI SESSSIYILNKEI
	114' QYNCKPGYATADGNSSGSITCLQNGWSTQPICIKFCDMPVFENSRAKSNGMWFKLHDTLD
10	**.*.*****.*****.*****.*****.
	361" QYKCKPGYATADGNSSGSITCLQNGWSAQPICIKFCDMPVFENSRAKSNGMRFKLHDTLD
	174' YECYDGYESSYGNTTDSIVCGEDGWSHLPTCYNSSESCGPPPI SN GDTTSFPQKVYLPW
	***** ***,*****.*****.*****.*****.***** ***,*
	421" YECYDGYEISYGNTTGSIVCGEDGWSHFPTCYNSSEKCGPPPI SN GDTTSFLLKVYVPQ
15	234' SRVEYQCQSYVELQGSKYVTCSNGDWSEPPRCISMKPCEFP--EIQHGHLYYENTRRPYF
	*****.*****.***** **
	481" SRVEYQCQSYVELQGSNYVTCSNGEWSEPPRCI--HPCIITEENMNKNNIQLKGKSDIKY
	292' PVATGQSYSYYCDQNFVTPSGSYWDYIHCTQDGLPTVPCLRTCSKSDIEIENGFI SESS
	. **.. .. *.** . *
20	539" YAKTGDTIEFMCKLGY-NANTSVLSFQAVCREGIVEYPRCE

The search of the protein data base using this sequence has revealed the presence of a sequence that is partly identical with the amino acid sequence of protein 4 relating to human complement factor H (GenBank Registration No. X98337). Table 2 shows the comparison of the amino acid sequences between the protein alike to human complement factor H (HS) of the present invention and protein 4 relating to human complement factor H (H4). Therein, the marks of - and * represent a gap and an amino acid

residue identical with the protein of the present invention,
 respectively. In other words, it has been revealed that, in the
 present protein, an amino acid sequence consisting of 247 residues
 is inserted after position 57 of protein 4 relating to human
 5 complement factor H.

Table 2

	HS	MLLLINVILTLWVSCANGQEVKPCDFPEIQHGGLYYKSLRRLYFPAAAGQSYSYYCDQNF

10	H4	MLLLINVILTLWVSCANGQEVKPCDFPEIQHGGLYYKSLRRLYFPAAAGQSYSYYCD---
	HS	VTPSGSYWDYIHCTQDGWSPTVPCLRTCSKSDVEIENGFI SESSSIYILNEETQYNCKPG
	H4	-----
	HS	YATADGNSSGSITCLQNGWSTQPIKFCMPVFENSRAKSNGMWFKLHDTLDYECYDGY
15	H4	-----
	HS	ESSYGNTTDSIVCGEDGWSHLPTCYNSSSESCGPPPPISNGDTSFPQKVYLPWSRVEYQC
	H4	-----
20	HS	QSYVELQGSKYVTCSNGDWSEPPRCISMKPCEFPEIQHGHLYYENTRRPYFPVATGQSYS
	H4	-----
	HS	YYCDQNFVTPSGSYWDYIHCTQDGLPTVPCLRTCSKSDIEIENGFI SESSSIYILNKEI

25	H4	----QNFVTPSGSYWDYIHCTQDGLPTVPCLRTCSKSDIEIENGFI SESSSIYILNKEI
	HS	QYKCKPGYATADGNSSGSITCLQNGWSAQPIKFCMPVFENSRAKSNGMRFKLHDTLD

	H4	QYKCKPGYATADGNSSGSITCLQNGWSAQPIKFCMPVFENSRAKSNGMRFKLHDTLD
	HS	YECYDGYEISYGNTTGSIVCGEDGWSHFPTCYNSEKCGPPPPISNGDTSFLLKVYVPQ
30		*****
	H4	YECYDGYEISYGNTTGSIVCGEDGWSHFPTCYNSEKCGPPPPISNGDTSFLLKVYVPQ
	HS	SRVEYQCQSYVELQGSNYVTCSNGEWSEPPRCIHPCIITEENMNKNNIQLKGKSDIKYYA

H4 SRVEYQCQSYVELQGSNYVTCSNGEWSEPPRCIHPCIITEENMNKNNIQLKGKSDIKYYA
 HS KTGDTIEFMCKLGYNANTSVLSFQAVCREGIVEYPRCE

5 H4 KTGDTIEFMCKLGYNANTSVLSFQAVCREGIVEYPRCE

There exist in the amino acid sequence of this protein seven sites where N-glycosylation is likely to occur (Asn-Ser-Ser at position 127, Asn-Thr-Thr at position 186, Asn-Ser-Ser at position 206, Asn-Ser-Ser at position 374, Asn-Thr-Thr at position 433, Asn-Ser-Ser at position 453, and Asn-Thr-Ser at position 557).

(4) Protein Synthesis by In Vitro Translation

Vector pHP01265 bearing the cDNA of the present invention was used for in vitro translation with a T₁T rabbit reticulocyte lysate kit (Promega). In this case, [³⁵S]methionine was added to label the expression product with a radioisotope. Each of the reactions was carried out according to the protocols attached to the kit. Two micrograms of the plasmid pHP01265 was reacted at 30°C for 90 minutes in a total 100 µl volume of the reaction solution containing 50 µl of T₁T rabbit reticulocyte lysate, 4 µl of a buffer solution (attached to kit), 2 µl of an amino acid mixture (methionine-free), 8 µl of [³⁵S]methionine (Amersham) (0.37 MBq/µl), 2 µl of T7RNA polymerase, and 80 U of RNasin. To 3 µl of the resulting reaction solution was added 2 µl of the SDS sampling buffer (125 mM Tris-hydrochloric acid buffer, pH 6.8, 120 mM 2-mercaptoethanol, 2% SDS solution, 0.025% bromophenol blue, and 20% glycerol) and the resulting mixture was heated at 95°C

for 3 minutes and then subjected to SDS-polyacrylamide gel electrophoresis. Determination of the molecular weight of the translation product by carrying out the autoradiograph indicated that the cDNA of the present invention yielded the translation
5 product with the molecular mass of about 65 kDa. This value is consistent with the molecular weight of 65,308 predicted for the putative protein from the base sequence represented by Sequence No. 1, thereby indicating that this cDNA certainly codes for the protein represented by Sequence No. 1.

10 (5) Functional Verification of Secretory Signal Sequence

It was verified by the method described in the literature [Yokoyama-Kobayashi, M. et al., Gene 163: 193-196 (1995)] that the N-terminal hydrophobic region in the secretory protein clone candidate obtained in the above-mentioned steps functions as a
15 secretory signal sequence. Vector pHP01265 was cleaved at the restriction enzyme site PvuII existing at the downstream of the portion expected for encoding the secretory signal sequence. Digestion with HindIII was further carried out and a DNA fragment containing the SV40 promoter and a cDNA encoding the secretory
20 signal sequence at the downstream of the promoter was separated by agarose gel electrophoresis. The resulting fragment was inserted between HindIII and PvuII in pSSD3 (DDBJ/EMBL/GenBank Registration No. AB007632), thereby constructing a vector expressing a fusion protein of the secretory signal sequence of
25 the target cDNA and the urokinase protease domain.

After *Escherichia coli* (host: JM109) bearing the fusion-protein expression vector was incubated at 37°C for 2 hours

in 2 ml of the 2xYT culture medium containing 100 μ g/ml of
ampicillin, the helper phage M13K07 (50 μ l) was added and the
incubation was continued at 37 °C overnight. A supernatant
separated by centrifugation underwent precipitation with
5 polyethylene glycol to obtain single-stranded phage particles.
These particles were suspended in 100 μ l of 1 mM Tris-0.1 mM EDTA,
pH 8 (TE). Also, there were used as controls suspensions of
single-stranded phage particles prepared in the same manner from
pSSD3 and from the vector pKA1-UPA containing a full-length cDNA
10 of urokinase [Yokoyama-Kobayashi, M. et al., Gene 163: 193-196
(1995)].

The culture cells originating from the simian kidney, COS7,
were incubated at 37°C in the presence of 5% CO₂ in the Dulbecco's
modified Eagle's culture medium (DMEM) containing 10% fetal calf
15 albumin. Into a 6-well plate (Nunc Inc., 3 cm in the well diameter)
were inoculated 1×10^5 COS7 cells and incubation was carried out
at 37°C for 22 hours in the presence of 5% CO₂. After the culture
medium was removed, the cell surface was washed with a phosphate
buffer solution and then washed again with DMEM containing 50 mM
20 Tris-hydrochloric acid (pH 7.5) (TDMEM). To the resulting cells
was added a suspension of 1 μ l of the single-stranded phage
suspension, 0.6 ml of the DMEM culture medium, and 3 μ l of
TRANSFECTAM™ (IBF Inc.) and the resulting mixture was incubated
at 37°C for 3 hours in the presence of 5% CO₂. After the sample
25 solution was removed, the cell surface was washed with TDMEM, 2
ml per well of DMEM containing 10% fetal calf albumin was added,
and the incubation was carried out at 37°C for 2 days in the presence

of 5% CO₂.

To 10 ml of 50 mM phosphate buffer solution (pH 7.4) containing 2% bovine fibrinogen (Miles Inc.), 0.5% agarose, and 1 mM calcium chloride were added 10 units of human thrombin (Mochida
5 Pharmaceutical Co., Ltd.) and the resulting mixture was solidified in a plate of 9 cm in diameter to prepare a fibrin plate. Ten microliters of the culture supernatant of the transfected COS7 cells were spotted on the fibrin plate, which was incubated at 37°C for 15 hours. Except for the case in which pSSD3 was used as
10 the control, each of the samples formed a clear circle to identify that urokinase was secreted in the culture medium. In other words, it has been indicated that the inserted cDNA fragment codes for the amino acid sequence that functions as the secretory signal sequence. Application of the (-3,-1) rule, a method for predicting
15 the cleavage site in the secretory signal sequence, allows to expect that the maturation protein starts from glutamine at position 19.

(6) Expression by COS7

Escherichia coli bearing pHP01265 of the present invention
20 was infected with helper phage M13K07 and single-stranded phage particles were obtained by the above-mentioned procedure. The thus-obtained phage was used for introducing pHP01265 in the culture cells originating from the simian kidney, COS7. After incubation at 37°C for 2 days in the presence of 5% CO₂, the
25 incubation was continued for one hour in the culture medium containing [³⁵S]cystine. Concentration of the culture medium, followed by subjecting to SDS-PAGE, allowed to observe the

presence of a band of about 120 kDa wherein the present protein was formed by the secretory expression, which did not exist in the culture supernatant from the COS7 cells only.

5 INDUSTRIAL APPLICABILITY

The present invention provides human cDNAs coding for proteins alike to human complement factor H and proteins encoded by these human cDNAs. The human cDNAs of the present invention can be utilized as probes for the gene diagnosis and gene sources
10 for the gene therapy. Furthermore, the cDNAs can be utilized as gene sources for large-scale production of the proteins encoded by said cDNAs. Eucaryotic cells wherein expression vectors of said cDNAs are introduced can be utilized for secretory production of the proteins alike to human complement factor H. Said recombinant
15 proteins can be employed as pharmaceuticals/research reagents, particularly as pharmaceuticals and research reagents for the immunological mechanism involving the complement reaction.

The present invention also provides genes corresponding to the polynucleotide sequences disclosed herein.
20 "Corresponding genes" are the regions of the genome that are transcribed to produce the mRNAs from which cDNA polynucleotide sequences are derived and may include contiguous regions of the genome necessary for the regulated expression of such genes. Corresponding genes may therefore include but are not limited to
25 coding sequences, 5' and 3' untranslated regions, alternatively spliced exons, introns, promoters, enhancers, and silencer or suppressor elements. The corresponding genes can be isolated in

accordance with known methods using the sequence information disclosed herein. Such methods include the preparation of probes or primers from the disclosed sequence information for identification and/or amplification of genes in appropriate
5 genomic libraries or other sources of genomic materials. An "isolated gene" is a gene that has been separated from the adjacent coding sequences, if any, present in the genome of the organism from which the gene was isolated.

Organisms that have enhanced, reduced, or modified
10 expression of the gene(s) corresponding to the polynucleotide sequences disclosed herein are provided. The desired change in gene expression can be achieved through the use of antisense polynucleotides or ribozymes that bind and/or cleave the mRNA transcribed from the gene (Albert and Morris, 1994, Trends
15 Pharmacol. Sci. 15(7): 250-254; Lavarosky et al., 1997, Biochem. Mol. Med. 62(1): 11-22; and Hampel, 1998, Prog. Nucleic Acid Res. Mol. Biol. 58: 1-39; all of which are incorporated by reference herein). Transgenic animals that have multiple copies of the gene(s) corresponding to the polynucleotide sequences disclosed
20 herein, preferably produced by transformation of cells with genetic constructs that are stably maintained within the transformed cells and their progeny, are provided. Transgenic animals that have modified genetic control regions that increase or reduce gene expression levels, or that change temporal or
25 spatial patterns of gene expression, are also provided (see European Patent No. 0 649 464 B1, incorporated by reference herein). In addition, organisms are provided in which the gene(s)

corresponding to the polynucleotide sequences disclosed herein have been partially or completely inactivated, through insertion of extraneous sequences into the corresponding gene(s) or through deletion of all or part of the corresponding gene(s). Partial
5 or complete gene inactivation can be accomplished through insertion, preferably followed by imprecise excision, of transposable elements (Plasterk, 1992, Bioessays 14(9): 629-633; Zwaal et al., 1993, Proc. Natl. Acad. Sci. USA 90(16): 7431-7435; Clark et al., 1994, Proc. Natl. Acad. Sci. USA 91(2): 719-722;
10 all of which are incorporated by reference herein), or through homologous recombination, preferably detected by positive/negative genetic selection strategies (Mansour et al., 1988, Nature 336: 348-352; U.S. Patent Nos. 5,464,764; 5,487,992; 5,627,059; 5,631,153; 5,614,396; 5,616,491; and 5,679,523; all
15 of which are incorporated by reference herein). These organisms with altered gene expression are preferably eukaryotes and more preferably are mammals. Such organisms are useful for the development of non-human models for the study of disorders involving the corresponding gene(s), and for the development of
20 assay systems for the identification of molecules that interact with the protein product(s) of the corresponding gene(s).

Where the protein of the present invention is membrane-bound (e.g., is a receptor), the present invention also provides for soluble forms of such protein. In such forms part
25 or all of the intracellular and transmembrane domains of the protein are deleted such that the protein is fully secreted from the cell in which it is expressed. The intracellular and

transmembrane domains of proteins of the invention can be identified in accordance with known techniques for determination of such domains from sequence information.

Proteins and protein fragments of the present invention
5 include proteins with amino acid sequence lengths that are at least 25% (more preferably at least 50%, and most preferably at least 75%) of the length of a disclosed protein and have at least 60% sequence identity (more preferably, at least 75% identity; most preferably at least 90% or 95% identity) with that disclosed
10 protein, where sequence identity is determined by comparing the amino acid sequences of the proteins when aligned so as to maximize overlap and identity while minimizing sequence gaps. Also included in the present invention are proteins and protein fragments that contain a segment preferably comprising 8 or more
15 (more preferably 20 or more, most preferably 30 or more) contiguous amino acids that shares at least 75% sequence identity (more preferably, at least 85% identity; most preferably at least 95% identity) with any such segment of any of the disclosed proteins.

Species homologs of the disclosed polynucleotides and
20 proteins are also provided by the present invention. As used herein, a "species homologue" is a protein or polynucleotide with a different species of origin from that of a given protein or polynucleotide, but with significant sequence similarity to the given protein or polynucleotide, as determined by those of skill
25 in the art. Species homologs may be isolated and identified by making suitable probes or primers from the sequences provided herein and screening a suitable nucleic acid source from the

desired species.

The invention also encompasses allelic variants of the disclosed polynucleotides or proteins; that is, naturally-occurring alternative forms of the isolated polynucleotide which
5 also encode proteins which are identical, homologous, or related to that encoded by the polynucleotides.

The invention also includes polynucleotides with sequences complementary to those of the polynucleotides disclosed herein.

The present invention also includes polynucleotides
10 capable of hybridizing under reduced stringency conditions, more preferably stringent conditions, and most preferably highly stringent conditions, to polynucleotides described herein. Examples of stringency conditions are shown in the table below: highly stringent conditions are those that are at least as
15 stringent as, for example, conditions A-F; stringent conditions are at least as stringent as, for example, conditions G-L; and reduced stringency conditions are at least as stringent as, for example, conditions M-R.

Table 3

Stringency Condition	Polynucleotide Hybrid	Hybrid Length (bp) [‡]	Hybridization Temperature and Buffer [†]	Wash Temperature and Buffer [†]
A	DNA : DNA	≥50	65°C; 1×SSC -or- 42°C; 1×SSC, 50% formamide	65°C; 0.3×SSC
B	DNA : DNA	<50	T _B *; 1×SSC	T _B *; 1×SSC
C	DNA : RNA	≥50	67°C; 1×SSC -or- 45°C; 1×SSC, 50% formamide	67°C; 0.3×SSC
D	DNA : RNA	<50	T _D *; 1×SSC	T _D *; 1×SSC
E	RNA : RNA	≥50	70°C; 1×SSC -or- 50°C; 1×SSC, 50% formamide	70°C; 0.3×SSC
F	RNA : RNA	<50	T _F *; 1×SSC	T _F *; 1×SSC
G	DNA : DNA	≥50	65°C; 4×SSC -or- 42°C; 4×SSC, 50% formamide	65°C; 1×SSC
H	DNA : DNA	<50	T _H *; 4×SSC	T _H *; 4×SSC
I	DNA : RNA	≥50	67°C; 4×SSC -or- 45°C; 4×SSC, 50% formamide	67°C; 1×SSC
J	DNA : RNA	<50	T _J *; 4×SSC	T _J *; 4×SSC
K	RNA : RNA	≥50	70°C; 4×SSC -or- 50°C; 4×SSC, 50% formamide	67°C; 1×SSC
L	RNA : RNA	<50	T _L *; 2×SSC	T _L *; 2×SSC
M	DNA : DNA	≥50	50°C; 4×SSC -or- 40°C; 6×SSC, 50% formamide	50°C; 2×SSC
N	DNA : DNA	<50	T _N *; 6×SSC	T _N *; 6×SSC
O	DNA : RNA	≥50	55°C; 4×SSC -or- 42°C; 6×SSC, 50% formamide	55°C; 2×SSC
P	DNA : RNA	<50	T _P *; 6×SSC	T _P *; 6×SSC
Q	RNA : RNA	≥50	60°C; 4×SSC -or- 45°C; 6×SSC, 50% formamide	60°C; 2×SSC
R	RNA : RNA	<50	T _R *; 4×SSC	T _R *; 4×SSC

‡ : The hybrid length is that anticipated for the hybridized region(s) of the hybridizing polynucleotides. When hybridizing a polynucleotide to a target polynucleotide of unknown sequence, the hybrid length is assumed to be that of the hybridizing polynucleotide. When polynucleotides of known sequence are hybridized, the hybrid length can be determined by aligning the sequences of the polynucleotides and identifying the region or regions of optimal sequence complementarity.

† : SSPE (1×SSPE is 0.15M NaCl, 10mM NaH₂PO₄, and 1.25mM EDTA, pH7.4) can be substituted for SSC (1×SSC is 0.15M NaCl and 15mM sodium citrate) in the hybridization and wash buffers; washes are performed for 15 minutes after hybridization is complete.

*T_B - T_R : The hybridization temperature for hybrids anticipated to be less than 50 base pairs in length should be 5-10°C less than the melting temperature (T_m) of the hybrid, where T_m is determined according to the following equations. For hybrids less than 18 base pairs in length, T_m(°C)=2(#of A + T bases) + 4(# of G + C bases). For hybrids between 18 and 49 base pairs in length, T_m(°C)=81.5 + 16.6(log₁₀[Na⁺]) + 0.41 (%G+C) - (600/N), where N is the number of bases in the hybrid, and [Na⁺] is the concentration of sodium ions in the hybridization buffer ([Na⁺] for 1×SSC=0.165M).

Additional examples of stringency conditions for polynucleotide hybridization are provided in Sambrook, J., E.F. Fritsch, and T. Maniatis, 1989, Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY, chapters 9 and 11, and Current Protocols in Molecular Biology, 1995, F.M. Ausubel et al., eds., John Wiley & Sons, Inc., sections 2.10 and 6.3-6.4, incorporated herein by reference.

Preferably, each such hybridizing polynucleotide has a length that is at least 25% (more preferably at least 50%, and most preferably at least 75%) of the length of the polynucleotide of the present invention to which it hybridizes, and has at least 60% sequence identity (more preferably, at least 75% identity; most preferably at least 90% or 95% identity) with the polynucleotide of the present invention to which it hybridizes, where sequence identity is determined by comparing the sequences of the hybridizing polynucleotides when aligned so as to maximize overlap and identity while minimizing sequence gaps.

Sequence Listing

<110> Sagami Chemical Research Center

5 <120> Proteins Alike to Human Complement Factor H and cDNAs Encoding these Proteins

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<151> 1997-10-06

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20 <211> 578

<212> PRT

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1

5

10

15

Asn Gly Gln Glu Val Lys Pro Cys Asp Phe Pro Glu Ile Gln His Gly

20

25

30

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Gly Leu Tyr Tyr Lys Ser Leu Arg Arg Leu Tyr Phe Pro Ala Ala Ala
 35 40 45
 Gly Gln Ser Tyr Ser Tyr Tyr Cys Asp Gln Asn Phe Val Thr Pro Ser
 50 55 60
 5 Gly Ser Tyr Trp Asp Tyr Ile His Cys Thr Gln Asp Gly Trp Ser Pro
 65 70 75 80
 Thr Val Pro Cys Leu Arg Thr Cys Ser Lys Ser Asp Val Glu Ile Glu
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 Thr Gln Tyr Asn Cys Lys Pro Gly Tyr Ala Thr Ala Asp Gly Asn Ser
 115 120 125
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 15 Cys Ile Lys Phe Cys Asp Met Pro Val Phe Glu Asn Ser Arg Ala Lys
 145 150 155 160
 Ser Asn Gly Met Trp Phe Lys Leu His Asp Thr Leu Asp Tyr Glu Cys
 165 170 175
 Tyr Asp Gly Tyr Glu Ser Ser Tyr Gly Asn Thr Thr Asp Ser Ile Val
 20 180 185 190
 Cys Gly Glu Asp Gly Trp Ser His Leu Pro Thr Cys Tyr Asn Ser Ser
 195 200 205
 Glu Ser Cys Gly Pro Pro Pro Pro Ile Ser Asn Gly Asp Thr Thr Ser
 210 215 220
 25 Phe Pro Gln Lys Val Tyr Leu Pro Trp Ser Arg Val Glu Tyr Gln Cys
 225 230 235 240
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 5 Pro Tyr Phe Pro Val Ala Thr Gly Gln Ser Tyr Ser Tyr Tyr Cys Asp
 290 295 300
 Gln Asn Phe Val Thr Pro Ser Gly Ser Tyr Trp Asp Tyr Ile His Cys
 305 310 315 320
 Thr Gln Asp Gly Trp Leu Pro Thr Val Pro Cys Leu Arg Thr Cys Ser
 10 325 330 335
 Lys Ser Asp Ile Glu Ile Glu Asn Gly Phe Ile Ser Glu Ser Ser Ser
 340 345 350
 Ile Tyr Ile Leu Asn Lys Glu Ile Gln Tyr Lys Cys Lys Pro Gly Tyr
 355 360 365
 15 Ala Thr Ala Asp Gly Asn Ser Ser Gly Ser Ile Thr Cys Leu Gln Asn
 370 375 380
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 Phe Glu Asn Ser Arg Ala Lys Ser Asn Gly Met Arg Phe Lys Leu His
 20 405 410 415
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 Asn Thr Thr Gly Ser Ile Val Cys Gly Glu Asp Gly Trp Ser His Phe
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 25 Pro Thr Cys Tyr Asn Ser Ser Glu Lys Cys Gly Pro Pro Pro Pro Ile
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 Ser Asn Gly Asp Thr Thr Ser Phe Leu Leu Lys Val Tyr Val Pro Gln
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Asn Tyr Val Thr Cys Ser Asn Gly Glu Trp Ser Glu Pro Pro Arg Cys
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5 Ile His Pro Cys Ile Ile Thr Glu Glu Asn Met Asn Lys Asn Asn Ile
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Gln Leu Lys Gly Lys Ser Asp Ile Lys Tyr Tyr Ala Lys Thr Gly Asp
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Thr Ile Glu Phe Met Cys Lys Leu Gly Tyr Asn Ala Asn Thr Ser Val
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Cys Glu

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agactatact ttccagcagc tgcaggacaa tcttattcct attactgtga tcaaaatttt 180

25 gtgactcett caggaagtta ctgggattac attcattgca cacaagatgg ttggtcacca 240

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	Phe Pro Ala Ala Ala Gly Gln Ser Tyr Ser Tyr Tyr Cys Asp Gln Asn	
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	Phe Val Thr Pro Ser Gly Ser Tyr Trp Asp Tyr Ile His Cys Thr Gln	
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	Asp Val Glu Ile Glu Asn Gly Phe Ile Ser Glu Ser Ser Ser Ile Tyr	
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Asn Ser Arg Ala Lys Ser Asn Gly Met Trp Phe Lys Leu His Asp Thr
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10 Leu Asp Tyr Glu Cys Tyr Asp Gly Tyr Glu Ser Ser Tyr Gly Asn Thr
175 180 185
aca gat tcc ata gtg tgt ggt gaa gat ggc tgg tcc cat ttg cca aca 686
Thr Asp Ser Ile Val Cys Gly Glu Asp Gly Trp Ser His Leu Pro Thr
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Cys Tyr Asn Ser Ser Glu Ser Cys Gly Pro Pro Pro Pro Ile Ser Asn
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gga gat acc acg tcc ttc ccg caa aaa gtg tat ctg cca tgg tca aga 782
Gly Asp Thr Thr Ser Phe Pro Gln Lys Val Tyr Leu Pro Trp Ser Arg
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gtc gag tac cag tgc cag tcc tac tat gaa ctt cag ggt tct aaa tat 830
Val Glu Tyr Gln Cys Gln Ser Tyr Tyr Glu Leu Gln Gly Ser Lys Tyr
240 245 250
gta aca tgt agt aat gga gac tgg tca gaa cca cca aga tgc ata tca 878
25 Val Thr Cys Ser Asn Gly Asp Trp Ser Glu Pro Pro Arg Cys Ile Ser
255 260 265
atg aaa cct tgt gag ttt cca gaa att caa cat gga cat cta tat tat 926

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Met Lys Pro Cys Glu Phe Pro Glu Ile Gln His Gly His Leu Tyr Tyr
270 275 280

gag aat acg cgt aga cca tac ttt cca gta gct aca gga caa tct tac 974
Glu Asn Thr Arg Arg Pro Tyr Phe Pro Val Ala Thr Gly Gln Ser Tyr

5 285 290 295

tcc tat tac tgt gac caa aat ttt gtg act cct tca gga agt tac tgg 1022
Ser Tyr Tyr Cys Asp Gln Asn Phe Val Thr Pro Ser Gly Ser Tyr Trp

300 305 310 315

gat tac att cac tgc aca caa gat ggg tgg ttg cca aca gtc cca tgc 1070
10 Asp Tyr Ile His Cys Thr Gln Asp Gly Trp Leu Pro Thr Val Pro Cys

320 325 330

ctc aga aca tgc tca aaa tca gat ata gaa att gaa aat gga ttc att 1118
Leu Arg Thr Cys Ser Lys Ser Asp Ile Glu Ile Glu Asn Gly Phe Ile

335 340 345

15 tct gaa tct tcc tct att tat att tta aat aaa gaa ata caa tat aaa 1166
Ser Glu Ser Ser Ser Ile Tyr Ile Leu Asn Lys Glu Ile Gln Tyr Lys

350 355 360

tgt aaa cca gga tat gca aca gca gat gga aat tct tca ggt tca att 1214
Cys Lys Pro Gly Tyr Ala Thr Ala Asp Gly Asn Ser Ser Gly Ser Ile

20 365 370 375

aca tgt ttg caa aat gga tgg tca gca caa cca att tgc att aaa ttt 1262
Thr Cys Leu Gln Asn Gly Trp Ser Ala Gln Pro Ile Cys Ile Lys Phe

380 385 390 395

tgt gat atg cct gtt ttt gag aat tcc aga gcc aag agt aat ggc atg 1310
25 Cys Asp Met Pro Val Phe Glu Asn Ser Arg Ala Lys Ser Asn Gly Met

400 405 410

cgg ttt aag ctc cat gac aca ttg gac tac gaa tgc tac gat gga tat 1358

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Arg Phe Lys Leu His Asp Thr Leu Asp Tyr Glu Cys Tyr Asp Gly Tyr
 415 420 425
 gaa atc agt tat gga aac acc aca ggt tcc ata gtg tgt ggt gaa gat 1406
 Glu Ile Ser Tyr Gly Asn Thr Thr Gly Ser Ile Val Cys Gly Glu Asp
 5 430 435 440
 ggg tgg tcc cat ttc cca aca tgt tat aat tct tca gaa aag tgt ggg 1454
 Gly Trp Ser His Phe Pro Thr Cys Tyr Asn Ser Ser Glu Lys Cys Gly
 445 450 455
 cct cct cca cct att agc aat ggt gat acc acc tcc ttt cta cta aaa 1502
 10 Pro Pro Pro Pro Ile Ser Asn Gly Asp Thr Thr Ser Phe Leu Leu Lys
 460 465 470 475
 gtg tat gtg cca cag tca aga gtc gag tac caa tgc cag tcc tac tat 1550
 Val Tyr Val Pro Gln Ser Arg Val Glu Tyr Gln Cys Gln Ser Tyr Tyr
 480 485 490
 15 gaa ctt cag ggt tct aat tat gta aca tgt agt aat gga gag tgg tgg 1598
 Glu Leu Gln Gly Ser Asn Tyr Val Thr Cys Ser Asn Gly Glu Trp Ser
 495 500 505
 gaa cca cca aga tgc ata cat cca tgt ata ata act gaa gaa aac atg 1646
 Glu Pro Pro Arg Cys Ile His Pro Cys Ile Ile Thr Glu Glu Asn Met
 20 510 515 520
 aat aaa aat aac ata cag tta aaa gga aaa agt gac ata aaa tat tat 1694
 Asn Lys Asn Asn Ile Gln Leu Lys Gly Lys Ser Asp Ile Lys Tyr Tyr
 525 530 535
 gca aaa aca ggg gat acc att gaa ttt atg tgt aaa ttg gga tat aat 1742
 25 Ala Lys Thr Gly Asp Thr Ile Glu Phe Met Cys Lys Leu Gly Tyr Asn
 540 545 550 555
 gcg aat aca tca gtt cta tca ttt caa gca gtg tgt agg gaa ggc ata 1790

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Ala Asn Thr Ser Val Leu Ser Phe Gln Ala Val Cys Arg Glu Gly Ile

560

565

570

gtg gaa tac ccc aga tgc gaa taaggcagc attggtaccc taaatgtatg 1840

Val Glu Tyr Pro Arg Cys Glu

5

575

tccaacttcc acttctcact cttatggtct caaagcttgc aaagatagct tctgatattg 1900

ttgtaatttc tactttatctt caaagaaaat taatataata gtttcaattt gcaacttaat 1960

atgttctcaa aaatatgtta aaacaaacta aattattgct tatgcttgta ctaaaataat 2020

aaaaactacc ctt 2033

10

<210> 4

<211> 578

<212> PRT

<213> Homo sapiens

15

<400> 4

Met Leu Leu Leu Ile Asn Val Ile Leu Thr Leu

1

5

10

Trp Val Ser Cys Ala Asn Gly Gln Glu Val Lys Pro Cys Asp Phe Pro

20

15

20

25

Glu Ile Gln His Gly Gly Leu Tyr Tyr Lys Ser Leu Arg Arg Leu Tyr

30

35

40

Phe Pro Ala Ala Ala Gly Gln Ser Tyr Ser Tyr Tyr Cys Asp Gln Asn

45

50

55

25

Phe Val Thr Pro Ser Gly Ser Tyr Trp Asp Tyr Ile His Cys Thr Gln

60

65

70

75

Asp Gly Trp Ser Pro Thr Val Pro Cys Leu Arg Thr Cys Ser Lys Ser

80

85

90

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Asp Val Glu Ile Glu Asn Gly Phe Ile Ser Glu Ser Ser Ser Ile Tyr
 95 100 105
 Ile Leu Asn Glu Glu Thr Gln Tyr Asn Cys Lys Pro Gly Tyr Ala Thr
 110 115 120
 5 Ala Asp Gly Asn Ser Ser Gly Ser Ile Thr Cys Leu Gln Asn Gly Trp
 125 130 135
 Ser Thr Gln Pro Ile Cys Ile Lys Phe Cys Asp Met Pro Val Phe Glu
 140 145 150 155
 Asn Ser Arg Ala Lys Ser Asn Gly Met Trp Phe Lys Leu His Asp Thr
 10 160 165 170
 Leu Asp Tyr Glu Cys Tyr Asp Gly Tyr Glu Ser Ser Tyr Gly Asn Thr
 175 180 185
 Thr Asp Ser Ile Val Cys Gly Glu Asp Gly Trp Ser His Leu Pro Thr
 190 195 200
 15 Cys Tyr Asn Ser Ser Glu Ser Cys Gly Pro Pro Pro Pro Ile Ser Asn
 205 210 215
 Gly Asp Thr Thr Ser Phe Pro Gln Lys Val Tyr Leu Pro Trp Ser Arg
 220 225 230 235
 Val Glu Tyr Gln Cys Gln Ser Tyr Tyr Glu Leu Gln Gly Ser Lys Tyr
 20 240 245 250
 Val Thr Cys Ser Asn Gly Asp Trp Ser Glu Pro Pro Arg Cys Ile Ser
 255 260 265
 Met Lys Pro Cys Glu Phe Pro Glu Ile Gln His Gly His Leu Tyr Tyr
 270 275 280
 25 Glu Asn Thr Arg Arg Pro Tyr Phe Pro Val Ala Thr Gly Gln Ser Tyr
 285 290 295
 Ser Tyr Tyr Cys Asp Gln Asn Phe Val Thr Pro Ser Gly Ser Tyr Trp
 300 305 310 315

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Asp Tyr Ile His Cys Thr Gln Asp Gly Trp Leu Pro Thr Val Pro Cys
 320 325 330
 Leu Arg Thr Cys Ser Lys Ser Asp Ile Glu Ile Glu Asn Gly Phe Ile
 335 340 345
 5 Ser Glu Ser Ser Ser Ile Tyr Ile Leu Asn Lys Glu Ile Gln Tyr Lys
 350 355 360
 Cys Lys Pro Gly Tyr Ala Thr Ala Asp Gly Asn Ser Ser Gly Ser Ile
 365 370 375
 Thr Cys Leu Gln Asn Gly Trp Ser Ala Gln Pro Ile Cys Ile Lys Phe
 10 380 385 390 395
 Cys Asp Met Pro Val Phe Glu Asn Ser Arg Ala Lys Ser Asn Gly Met
 400 405 410
 Arg Phe Lys Leu His Asp Thr Leu Asp Tyr Glu Cys Tyr Asp Gly Tyr
 415 420 425
 15 Glu Ile Ser Tyr Gly Asn Thr Thr Gly Ser Ile Val Cys Gly Glu Asp
 430 435 440
 Gly Trp Ser His Phe Pro Thr Cys Tyr Asn Ser Ser Glu Lys Cys Gly
 445 450 455
 Pro Pro Pro Pro Ile Ser Asn Gly Asp Thr Thr Ser Phe Leu Leu Lys
 20 460 465 470 475
 Val Tyr Val Pro Gln Ser Arg Val Glu Tyr Gln Cys Gln Ser Tyr Tyr
 480 485 490
 Glu Leu Gln Gly Ser Asn Tyr Val Thr Cys Ser Asn Gly Glu Trp Ser
 495 500 505
 25 Glu Pro Pro Arg Cys Ile His Pro Cys Ile Ile Thr Glu Glu Asn Met
 510 515 520
 Asn Lys Asn Asn Ile Gln Leu Lys Gly Lys Ser Asp Ile Lys Tyr Tyr
 525 530 535

WO 99/18200

13/13

PCT/JP98/04448

Ala Lys Thr Gly Asp Thr Ile Glu Phe Met Cys Lys Leu Gly Tyr Asn

540

545

550

555

Ala Asn Thr Ser Val Leu Ser Phe Gln Ala Val Cys Arg Glu Gly Ile

560

565

570

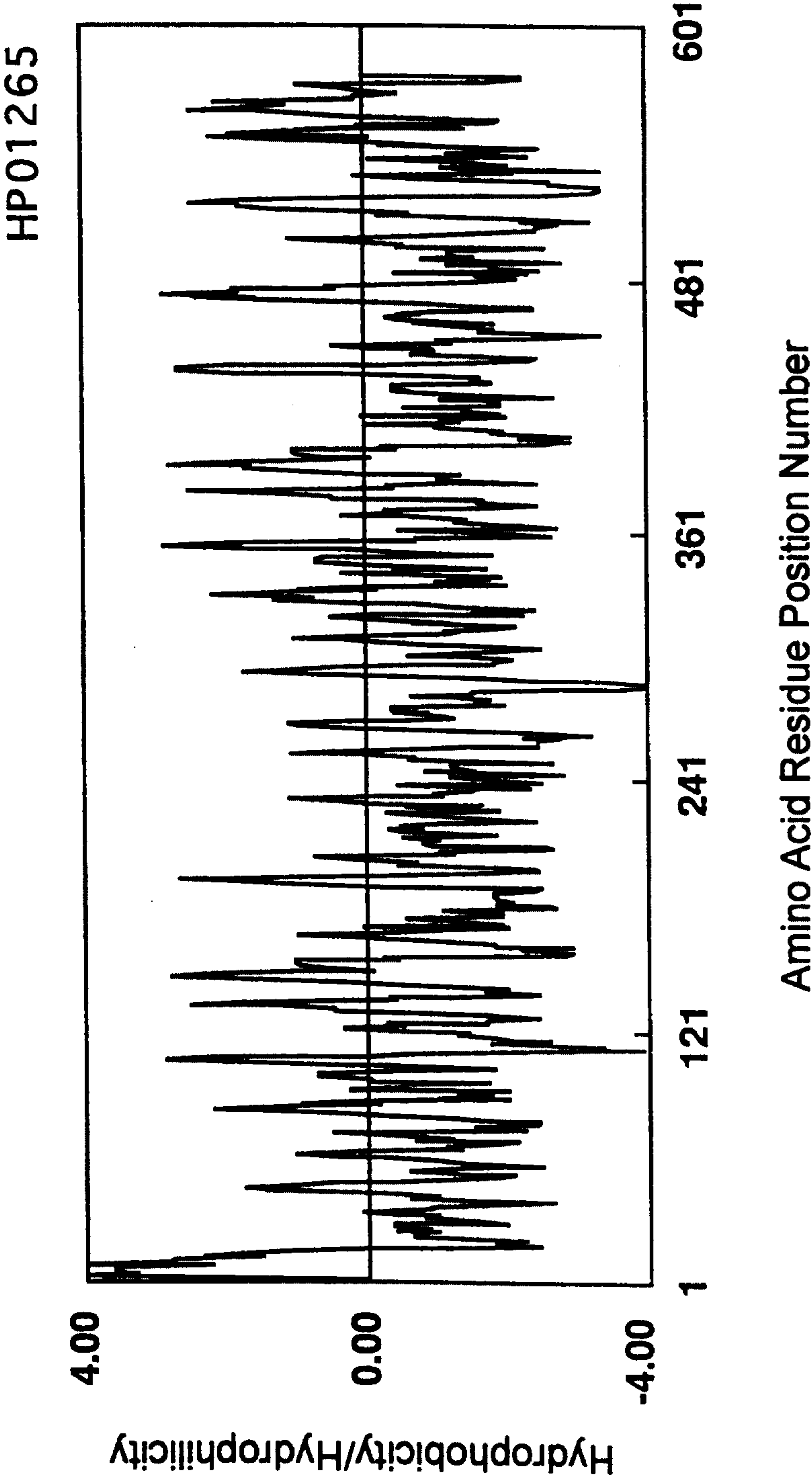
5 Val Glu Tyr Pro Arg Cys Glu

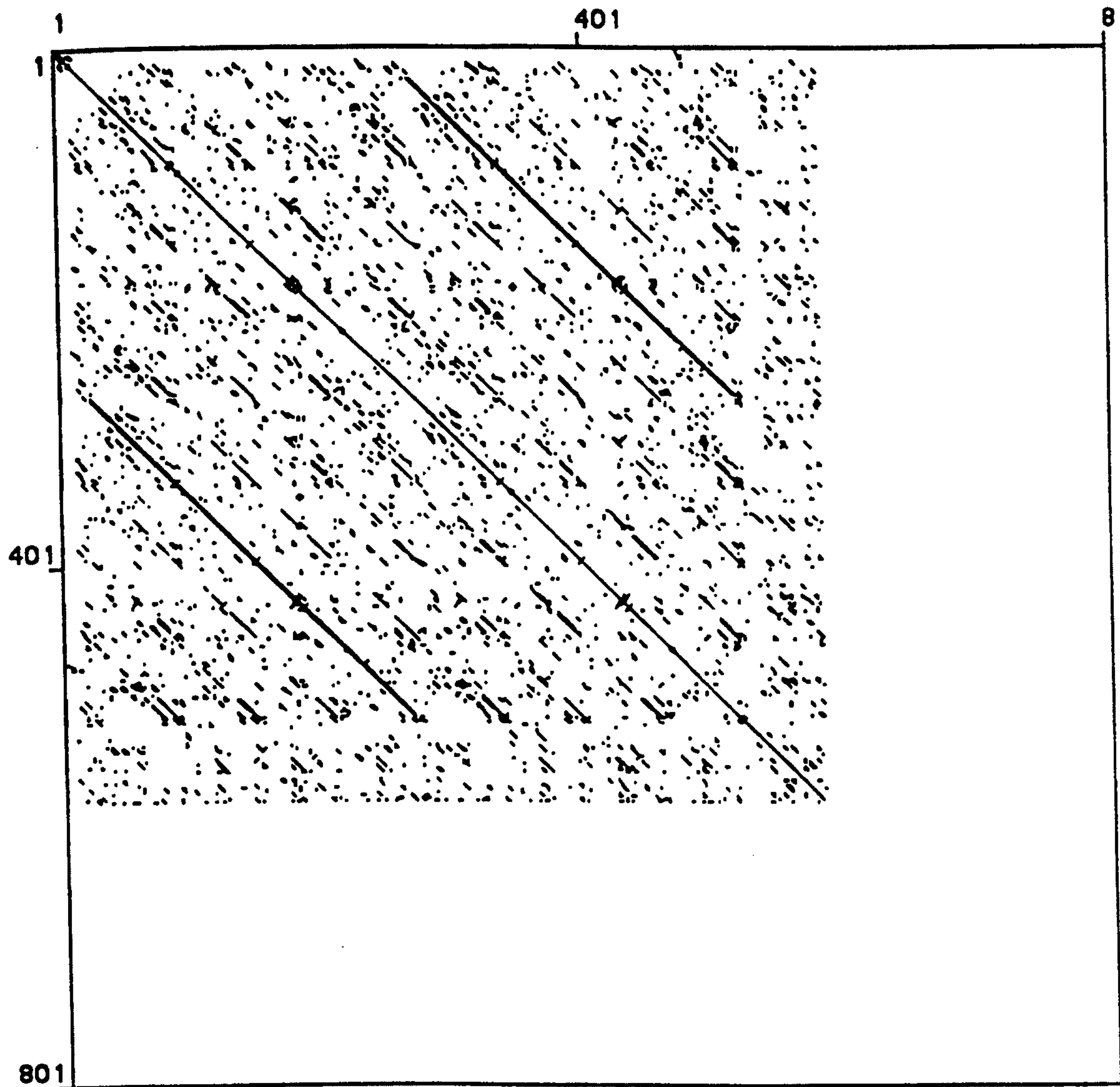
575

CLAIMS

1. A protein comprising the amino acid sequence represented by Sequence No. 1.
- 5 2. A DNA coding for the protein according to Claim 1.
3. A cDNA comprising the base sequence represented by Sequence No. 2.
4. The cDNA according to Claim 3, which comprises the base sequence represented by Sequence No. 3.
- 10 5. A vector capable of expressing the DNA according to any of Claims 2 to 4 in in vitro translation or an eukaryotic cell.
6. A transformation eukaryotic cell capable of expressing the DNA according to any of Claims 2 to 4 to produce the protein according to Claim 1.

Fig. 1





[H] : HP01265.PEP
[U] : HP01265.PEP
Unit Size to Compare : 5
Dot Plot Score : 2.00

Position : 1 - 578
Position : 1 - 578

Fig. 2