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(54) **METHOD FOR DETECTING PANCREATIC CANCER**

(57) This invention relates to a method for detecting pancreatic cancer using novel tumor markers. Specifically, the invention provides a method for detecting pancreatic cancer comprising measuring the presence or an amount of a polypeptide having an reactivity of binding via an antigen-antibody reaction to an antibody against

CAPRIN-1 protein in a sample separated from a subject, and to a reagent or kit for detecting pancreatic cancer comprising a CAPRIN-1 protein or a fragment thereof, an antibody against the same, or a polynucleotide encoding the same.

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Description

Technical Field

5 **[0001]** The present invention relates to a method for detecting pancreatic cancer using CAPRIN-1 as a tumor marker.

Background Art

10 **[0002]** It is reported that there are over 10,000 patients with refractory pancreatic cancer in Japan, the occurrence thereof is increasing year after year, and it is assumed that the number of patients will continue to increase. Even if pancreatic cancer were surgically removed, small cancer cells have often infiltrated and metastasized to other organs. Accordingly, pancreatic cancer often relapses, and the 5-year survival rate is as low as 9%; that is, the prognosis of pancreatic cancer is very poor. For the purpose of preventing postoperative recurrence, Gemcitabine, an anticancer agent, has been employed. However, the primary purpose of Gemcitabine administration is pain relief, and tumor size reduction or survival advantage can hardly be expected. At some hospitals, another anti-cancer agent, TS-1, which is currently used for gastric cancer, is used, although it is difficult to expect any therapeutic effects.

15 **[0003]** In order to improve the prognosis for pancreatic cancer, early detection is important, as it is with other cancers; however, early detection is difficult because pancreatic cancer shows substantially no initial symptoms. To date, methods of detecting pancreatic cancer using carcinoembryonic antigen (CEA) and glycoproteins (CA19-9 and Dupan-2) in biological samples as tumor markers of pancreatic cancer have been actively employed. However, the levels of such tumor markers do not become elevated unless pancreatic cancer advances, and such markers occasionally show normal values in the progressive stage. Accordingly, such tumor markers are not considered to be sufficient for accurate detection of pancreatic cancer. In addition, most tumor markers that are currently known are present in very small amounts in the body fluids (at the pg/ml level). In order to detect such small amounts of markers, accordingly, detection techniques with high sensitivity or special techniques are necessary. Under such circumstances, a novel technique for detecting pancreatic cancer in a simple manner with high sensitivity is expected to be applicable to diagnosis of pancreatic cancer. It is necessary to undergo periodic thorough examinations in order to detect early-stage pancreatic cancer. Accordingly, a method of detecting cancer that can be carried out in a simple manner with the use of blood serum or urine samples without the imposition of physical or financial burdens on either healthy individuals without pancreatic cancer or patients with cancer has been awaited.

20 **[0004]** Also, pancreatic cancer is refractory in dogs. Although a lump can be observed in the abdominal region of a dog afflicted with pancreatic cancer, the major symptoms are rapid energy loss, unsteady gait, and gait abnormalities resulting from hypoglycemia. In most cases, the development of cancer would not be detected until such symptoms are observed. In addition, pancreatic cancer is often likely already to be in the advanced stage when such symptoms are observed. In addition to surgical removal of pancreatic cancer, accordingly, therapeutic techniques are limited to supportive therapy and administration of anticancer agents. As with the case of human patients, early detection is important for dogs afflicted with pancreatic cancer in order to effectively treat such pancreatic cancer. As with the case for humans, there were no diagnostic agents for dogs in the past that allowed detection of pancreatic cancer at the early stage in a simple manner. In the field of veterinary medicine, detection techniques such as radiographic techniques by means of X-rays, CT, or MRI have not yet become common. At present, detection is carried out by palpation, simple blood testing, and X-ray photography, and diagnosis is heavily dependent on the experience of veterinary doctors. If a simple means for cancer detection with high sensitivity that can be applied to diagnosis of pancreatic cancer in dogs is provided, adequate treatment can be performed, which has great advantages for dog owners and veterinary doctors.

25 **[0005]** Cytoplasmic- and proliferation-associated protein 1 (CAPRIN-1) is an intracellular protein that is expressed when normal cells in the resting phase are activated or undergo cell division. CAPRIN-1 is also known to be involved in the control of the transport and translation of mRNAs through formation of cytoplasmic stress granules and RNA in a cell. Also, genes encoding CAPRIN-1 proteins are demonstrated to be expressed specifically in canine and human testis and malignant tumor cells, FCM analysis of breast cancer cells with the use of antibodies against CAPRIN-1 demonstrates CAPRIN-1 expression on the surfaces of breast cancer cells, and immunohistochemical staining using breast cancer tissues demonstrates CAPRIN-1 expression at high level in breast cancer cells. In addition, it has been reported that the antibodies mentioned above would damage breast cancer cells through the functions of lymphocytes, and that antibodies against CAPRIN-1 exert potent antitumor effects in cancer-carrying mouse models into which breast cancer cells had been transplanted (Patent Literature 1). Also, it has been reported that cancers such as breast cancer could be diagnosed by measuring either antibodies induced in the body of a subject against CAPRIN-1 present in the blood serum or polypeptides that undergo antigen-antibody reactions with CAPRIN-1 (Patent Literature 2). Up to the present, however, there have been no reports of the fact that pancreatic cancer can be diagnosed by measuring either antibodies against CAPRIN-1 induced in the blood serum of a patient with pancreatic cancer or polypeptides that undergo antigen-antibody reactions with the CAPRIN-1.

Prior Art Literatures

Patent Literature

5 **[0006]**

Patent Literature 1: International Publication No. WO 2010/016526

Patent Literature 2: International Publication No. WO 2010/016527

10 Summary of the Invention

Problem to Be Attained by the Invention

15 **[0007]** It is an object of the present invention to provide a means for detecting pancreatic cancer that is useful for diagnosis of pancreatic cancer.

Means for Solving the Problem

20 **[0008]** The present inventors have conducted concentrated studies. As a result, the present inventors have now found that pancreatic cancer can be diagnosed, examined, or detected on the basis of CAPRIN-1 expression in pancreatic cancer, measurement (or assay) of antibodies against CAPRIN-1 induced in the blood serum of a patient with pancreatic cancer with the use of a CAPRIN-1 protein, and binding of the antibodies produced with the use of such proteins to CAPRIN-1 in the pancreatic cancer tissue. This has led to the completion of the present invention.

25 **[0009]** Specifically, the present invention provides a method for detecting pancreatic cancer comprising measuring a CAPRIN-1 expression in a sample separated from a subject. The term "detecting" as used herein can be used interchangeably with the term "examining" or "evaluating." Also, the present invention provides a reagent or kit for detecting pancreatic cancer comprising a polypeptide that undergoes an antigen-antibody reaction with an antibody against CAPRIN-1 induced or elicited in the body of a subject. Further, the present invention provides a reagent or kit for detecting pancreatic cancer comprising an antibody that undergoes an antigen-antibody reaction with CAPRIN-1 or an antigen-binding fragment of the antibody. Furthermore, the present invention provides a reagent or kit for detecting pancreatic cancer comprising a polynucleotide hybridizing specifically to a partial sequence comprising at least 15 to 19 nucleotides or at least 20 to 30 nucleotides of the nucleotide sequence represented by any of SEQ ID NOs: 1, 3, 5, 7, 9, 11, 13, ...,29. The "reagent or kit for detecting pancreatic cancer" used herein can also be referred to as a "reagent or kit for pancreatic cancer detection."

35 **[0010]** Specifically, the present invention has the features described below.

(1) A method for detecting pancreatic cancer comprising measuring the presence or amount of a polypeptide having a reactivity of specifically binding to an antibody against a CAPRIN-1 protein via an antigen-antibody reaction, or the presence or amount of a nucleic acid encoding the polypeptide in a sample separated from a subject.

40 (2) The method according to (1), wherein the polypeptide to be measured is a CAPRIN-1 protein consisting of an amino acid sequence represented by any of the even-numbered SEQ ID NOs: 2 to 30 or a polypeptide consisting of an amino acid sequence having 85-90% or higher sequence identity with the CAPRIN-1 protein.

(3) The method according to (1) or (2), wherein the subject is a human or a dog.

45 (4) The method according to (3), wherein the subject is a dog and the polypeptide to be measured comprises an amino acid sequence represented by any of the even-numbered SEQ ID NOs: 2 to 30.

(5) The method according to (4), wherein the subject is a dog and the polypeptide to be measured comprises the amino acid sequence represented by SEQ ID NO: 6, 8, 10, 12, or 14.

(6) The method according to (3), wherein the subject is a human and the polypeptide to be measured comprises the amino acid sequence represented by SEQ ID NO: 2 or 4.

50 (7) The method according to any of (1) to (6), wherein the presence or amount of the polypeptide is determined by immunologically measuring an antibody induced against the polypeptide to be measured in the body of a subject, which can be contained in the sample.

(8) The method according to any of (1) to (6), wherein the presence or amount of a nucleic acid encoding the polypeptide is determined by measuring a nucleic acid encoding such polypeptide contained in the sample.

55 (9) The method according to (8), wherein the presence or amount of the nucleic acid in the sample is measured using a polynucleotide specifically hybridizing to a partial sequence comprising at least 15 to 19 nucleotides, preferably at least 20 to 25 nucleotides, and more preferably at least 30 nucleotides of the nucleotide sequence in the nucleic acid or a sequence complementary thereto.

(10) The method according to (9), wherein the subject is a dog and the polynucleotide specifically hybridizes to a partial sequence comprising at least 15 to 19 nucleotides, preferably at least 20 to 25 nucleotides, and more preferably at least 30 nucleotides in the nucleotide sequence represented by SEQ ID NO: 5, 7, 9, 11, or 13 or a sequence complementary thereto.

(11) The method according to (9), wherein the subject is a human and the polynucleotide specifically hybridizes to a partial sequence comprising at least 15 to 19 nucleotides, preferably at least 20 to 25 nucleotides, and more preferably at least 30 nucleotides in the nucleotide sequence represented by SEQ ID NO: 1 or 3 or a sequence complementary thereto.

(12) The method according to any of (1) to (6), wherein the presence or amount of the polypeptide is determined by measuring the polypeptide contained in the sample.

(13) The method according to (12), wherein the assay is an immunological assay.

(14) The method according to any of (1) to (13), wherein the sample is blood, serum, blood plasma, ascites fluid, pleural effusion, tissues, or cells.

(15) A reagent or kit for detecting pancreatic cancer comprising one or more polypeptides having a reactivity of binding via an antigen-antibody reaction to an antibody induced against a CAPRIN-1 protein in the body of a subject.

(16) A reagent or kit for detecting pancreatic cancer comprising one or more antibodies that undergo an antigen-antibody reaction with a polypeptide having a reactivity of binding via an antigen-antibody reaction to an antibody against a CAPRIN-1 protein and produced in the body of a subject or to an antigen-binding fragment of the antibody.

(17) The reagent or kit according to (15) or (16), wherein the CAPRIN-1 protein has an amino acid sequence represented by any of the even-numbered SEQ ID NOs: 2 to 30.

(18) The reagent or kit according to (16) or (17), wherein the antibody or an antigen-binding fragment thereof that undergoes an antigen-antibody reaction with the polypeptide is an antibody or an antigen-binding fragment thereof that binds to the surface of a pancreatic cancer cell.

(19) The reagent or kit according to any of (16) to (18), wherein the antibody or an antigen-binding fragment thereof that undergoes an antigen-antibody reaction with the polypeptide comprises an antibody or a fragment thereof having an immunological reactivity with a polypeptide consisting of an amino acid sequence comprising at least 7 to 12 continuous amino acid residues within the region of amino acid residue Nos. 50 to 98 or amino acid residue Nos. 233 to 344 of the amino acid sequence represented by any of the even-numbered SEQ IDS NO: 2 to 30 except for SEQ ID NOs: 6 and 18 or with a polypeptide comprising the polypeptide as a partial sequence.

(20) The reagent or kit according to any of (16) to (19), wherein the antibody or an antigen-binding fragment that undergoes an antigen-antibody reaction with the polypeptide is one or more antibodies selected from the group consisting of: an antibody binding to a polypeptide comprising the amino acid sequence represented by SEQ ID NO: 43 or antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 44 and 45 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 44 and 46 or an antigen-binding fragment of each thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 44 and 47 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 44 and 48 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 49 and 50 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 51 and 52 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 53 and 54 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 55 and 56 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 57 and 58 or an antigen-binding fragment thereof; and a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 59 and 60 or an antigen-binding fragment thereof.

(21) A reagent or kit for detecting pancreatic cancer comprising one or more polynucleotides that specifically hybridize to a partial sequence comprising at least 15 to 19 nucleotides, preferably at least 20 to 25 nucleotides, and more preferably at least 30 nucleotides in the nucleotide sequence represented by any of the odd-numbered SEQ ID NOs: 1 to 29 and encoding a CAPRIN-1 protein or in a sequence complementary to the nucleotide sequence.

(22) A method for detecting pancreatic cancer comprising measuring the presence or amount of a CAPRIN-1 protein, an antibody against a CAPRIN-1 protein, or a nucleic acid encoding the CAPRIN-1 protein in a sample from a subject using at least one reagent or kit according to any of (15) to (21).

(23) The method for detecting pancreatic cancer according to any of (1) to (14), comprising measuring the presence or an amount of a CAPRIN-1 protein, an antibody against a CAPRIN-1 protein, or a nucleic acid encoding the CAPRIN-1 protein in a sample from a subject using at least one reagent or kit according to any one of (15) to (21).

[0011] According to the present invention, a novel method for detecting pancreatic cancer is provided. As specifically described in the examples below, a recombinant polypeptide prepared based on the amino acid sequence of CAPRIN-

1 (or Caprin-1) is capable of specifically reacting with an antibody that exists in the serum of a patient with pancreatic cancer. As such, the pancreatic cancer existing in a subject can be detected by measuring the antibody in a sample by the method of the present invention. Also, pancreatic cancer existing in a subject can be detected by measuring (or assaying) CAPRIN-1 itself. As described in the examples below, in addition, high levels of CAPRIN-1 gene expression are observed specifically in testis and pancreatic cancer cells of subjects (hereinafter, such expression product is occasionally referred to as a "nucleic acid encoding a CAPRIN-1 (protein)"). Therefore, pancreatic cancer can also be detected by measuring a nucleic acid. Further, the presence or amount of CAPRIN-1 (expression) in the pancreatic cancer tissue can be measured with the use of an antibody against CAPRIN-1. Patients with pancreatic cancer may be subjected to such measurement in advance, so that the patients to whom a CAPRIN-1-targeted therapeutic agent (e.g., antibody medicine) is applicable can be selected.

Embodiments for Carrying out the Invention

[0012] According to the method of the present invention, the presence or an amount of CAPRIN-1 (expression) is measured using a sample separated from a subject. Examples of methods for measuring the presence or an amount of CAPRIN-1 (expression) include: a method for immunologically measuring an antibody against CAPRIN-1 contained in a sample (the first method); a method for immunologically measuring CAPRIN-1 itself contained in a sample (the second method); and a method for measuring a nucleic acid encoding CAPRIN-1 contained in a sample, such as mRNA or cDNA synthesized from mRNA (the third method). In the present invention, the presence or an amount of CAPRIN-1 (expression) may be measured by any of the above methods. In the present invention, the term "measuring" is intended to include any of the following meaning: detecting, qualitatively measuring, quantitatively measuring, and semi-quantitatively measuring.

[0013] The amino acid sequence represented by SEQ ID NOs: 6, 8, 10, 12, or 14 is an amino acid sequence of canine CAPRIN-1. Canine CAPRIN-1 having such an amino acid sequence has been identified as a polypeptide binding to an antibody specifically existing in the serum derived from a cancer-bearing dog (see Example 1). An antibody against CAPRIN-1 having the amino acid sequence represented by SEQ ID NO: 6, 8, 10, 12, or 14 is specifically induced or elicited in the body of a cancer-bearing dog. Specifically, canine pancreatic cancer can be detected by measuring the above antibody against CAPRIN-1 having the amino acid sequence represented by SEQ ID NO: 6, 8, 10, 12, or 14 by the first method. Also, canine pancreatic cancer can be detected by measuring CAPRIN-1 itself as an antigen having the amino acid sequence represented by SEQ ID NO: 6, 8, 10, 12, or 14 by the second method. Since the CAPRIN-1 gene is expressed at significantly high levels in pancreatic cancer cells, canine pancreatic cancer can be detected by measuring the nucleic acid in accordance with the third method.

[0014] The term "having an amino acid sequence" used herein refers to amino acid residues aligned in a given order. Therefore, for example, the expression "polypeptide having the amino acid sequence represented by SEQ ID NO: 2" refers to a polypeptide having 709 amino acid residues, which consists of the amino acid sequence of Met Pro Ser Ala ... (partially omitted) ... Gln Gln Val Asn represented by SEQ ID NO: 2. Also, the expression "polypeptide having the amino acid sequence represented by SEQ ID NO: 2" may also be abbreviated as "the polypeptide of SEQ ID NO: 2," for example. The same applies to the expression "having a/ nucleotide sequence." In this case, the term "having" may be substituted with the expression "comprising" or "consisting of."

[0015] Also, the term "polypeptide" used herein refers to a molecule that is formed via a peptide bond of a plurality of amino acids. Examples of such molecule include not only polypeptide molecules with large numbers of constituent amino acids, but also low-molecular-weight molecules (oligopeptides) with small numbers of amino acids and full-length proteins. The present invention further encompasses full-length CAPRIN-1 proteins each having an amino acid sequence represented by any of the even-numbered SEQ ID NOs: 2 to 30 (i.e., SEQ ID NOs: 2, 4, 6, ..., 26, 28, and 30).

[0016] The term "subject" used herein refers to vertebrates, including mammals and birds, preferably mammals, and more preferably humans, dogs, cows, and horses.

[0017] The term "sample" used herein refers to a biological sample subjected to examination aimed at detection of pancreatic cancer. Examples of the sample include body fluids, tissues, and cells separated from a subject. Examples of body fluids include, but are not limited to, blood, serum, blood plasma, ascites fluid, and pleural effusion. Tissues or cells of the pancreas that is suspected of being afflicted with cancer are within the scope of the "sample."

[0018] In the method of the present invention, the targets to be measured are to not only canine CAPRIN-1 of SEQ ID NO: 6, 8, 10, 12, or 14, but also CAPRIN-1 of other mammals (hereinafter, which may also be referred to as a "homolog" (or "ortholog") for canine CAPRIN-1). When it is simply referred to as "CAPRIN-1," CAPRIN-1 from another mammal, including a human, is also a target to be measured, in addition to CAPRIN-1 from a dog. As specifically described in the examples below, the human CAPRIN-1 gene expression level is significantly high in human pancreatic cancer cells, whereas no antibody against human CAPRIN-1 gene is detected in a healthy human body. As such, pancreatic cancer of a mammal other than a dog can be detected by measuring CAPRIN-1 expression in the mammal. An example of CAPRIN-1 of a mammal other than a dog to be measured by the method of the present invention is, but

is not limited to, human CAPRIN-1. Nucleotide sequences encoding human CAPRIN-1 and amino acid sequences therefor are represented by SEQ ID NOs: 1 and 3 and SEQ ID NOs: 2 and 4 in the Sequence Listing. Sequence identity between human CAPRIN-1 and canine CAPRIN-1 is 94% for nucleotide sequence and is 98% for amino acid sequence. The sequence identity of the amino acid sequences of CAPRIN-1 is as high as 98% between genetically distant mammals, such as a dog and a human. Therefore, it is considered that the sequence identity is about 85% or higher between a dog and a mammal other than a human; that is, canine CAPRIN-1 and its homolog. CAPRIN-1, the expression of which is to be measured by the method of the present invention, preferably has 85% or higher, more preferably 90% or higher, and further preferably 95% or higher sequence identity with the amino acid sequence of canine CAPRIN-1 represented by SEQ ID NO: 6, 8, 10, 12, or 14, although the sequence identities are not limited thereto.

[0019] In the first method, the antibody that can be present in a sample can be easily measured by immunoassay using an antigenic substance that undergoes an antigen-antibody reaction with the antibody. Immunoassay itself is a well-known conventional method as specifically described below. As an antigenic substance for immunoassay, for example, the canine CAPRIN-1 protein of SEQ ID NO: 6, 8, 10, 12, or 14 that induces the antibody within the body of a cancer-bearing dog or a fragment containing an epitope of such protein can be used. Further, the antibody has a cross-reactivity. A molecule other than an antigenic substance that actually serves as an immunogen can also bind, via an antigen-antibody reaction, to an antibody induced against an immunogen, as long as the molecule has a structure analogous to an epitope of the immunogen. Between a protein from a certain type of mammal and a homolog thereof from another mammal, in particular, the identity of their amino acid sequences is high, and epitope structures are often analogous to each other. As specifically described in the examples below, the canine CAPRIN-1 of SEQ ID NO: 6, 8, 10, 12, or 14 undergoes an antigen-antibody reaction with an antibody induced against the canine CAPRIN-1 within the body of a cancer-bearing dog. Also, human CAPRIN-1 undergoes an antigen-antibody reaction with the antibody induced within the body of a cancer-bearing dog. Accordingly, CAPRIN-1 from any mammal can be used as an antigen for immunoassay in accordance with the first method of the present invention.

[0020] When an antigenic substance is a protein or the like having a complicated structure and a high molecular weight, in general, a plurality of sites having different structures are present on the molecule. Therefore, a plurality of types of antibodies capable of recognizing and binding to different sites of such antigenic substances are produced in the body of a subject. Specifically, an antibody that is produced in the subject against an antigenic substance such as a protein is a polyclonal antibody that is a mixture of a plurality of types of antibodies. An antibody now found by the present inventors is also a polyclonal antibody that is specifically present in the serum obtained from a cancer-bearing subject and specifically binds, via an antigen-antibody reaction, to a recombinant CAPRIN-1 protein. The term "polyclonal antibody" used in the present invention refers to an antibody that exists in the serum obtained from a subject containing an antigenic substance and is induced against such antigenic substance.

[0021] In the examples below, polypeptides of SEQ ID NO: 6 and SEQ ID NO: 8 (both, canine CAPRIN-1) and the polypeptide of SEQ ID NO: 2 (human CAPRIN-1) were prepared as antigens for immunoassay of specific antibodies in cancer-bearing living animals. The reactivity between these polypeptides and the antibodies in the serum obtained from a cancer-bearing subject was then confirmed. However, the antibodies mentioned above are polyclonal antibodies, and they naturally bind to polypeptides consisting of homologs of SEQ ID NO: 6, 8, and 2. Even in the case of a fragment of such a polypeptide, it can bind to an antibody contained in the serum obtained from a cancer-bearing subject since some polyclonal antibodies are capable of recognizing the structure of the fragment. That is, both the polypeptide (that is, the full-length CAPRIN-1 protein) of the homolog of SEQ ID NO: 6, 8, or 2 and a fragment thereof can be similarly used for assay of a polyclonal antibody contained specifically in the serum of a cancer-bearing subject, and they are useful for cancer detection. Accordingly, a polypeptide to be used as an antigen for immunoassay in the first method of the present invention is not limited to a polypeptide alone consisting of the full-length region of a CAPRIN-1 protein (e.g., SEQ ID NO: 6, 8, or 2). It can be a polypeptide fragment consisting of at least 7 to 12, and preferably at least 8, 9, or 10 continuous amino acids of the amino acid sequence of a CAPRIN-1 protein that undergoes an antigen-antibody reaction with a polyclonal antibody against the CAPRIN-1 protein (hereinafter, it may be referred to as a "specifically reactive partial polypeptide" for convenience). It is known in the art that a polypeptide comprising about 7 to 12 or more amino acid residues can exert antigenicity. If the number of amino acid residues is too low, however, such polypeptide is highly likely to cross-react with an antibody against a protein other than the CAPRIN-1 protein that exists in the sample. In view of enhancing the accuracy of immunoassay, accordingly, the number of amino acid residues of a polypeptide fragment is preferably 20 or more, 30 or more, and 50 or more, more preferably 100 or more or 150 or more, further preferably 300 or more, and even further preferably 600 or more. The number of amino acid residues may be 1,000 or more, or 1,500 or more.

[0022] Preferable examples of the polypeptides to be used as antigens include the polypeptides of the even-numbered SEQ ID NOs: 2 to 30 or fragments thereof comprising epitopes (e.g., a polypeptide fragment comprising about 7 to 12 or more amino acid residues).

[0023] Nucleotide sequences of polynucleotides encoding proteins consisting of the amino acid sequences of the even-numbered SEQ ID NOs: 2 to 30 (i.e., SEQ ID NOs: 2, 4, 6...28, and 30) are represented by the odd-numbered

SEQ ID NOs: 1 to 29 (i.e., SEQ ID NOs: 1, 3, 5...27, and 29).

[0024] In general, it is well known in the art that protein antigens retain antigenicity almost equivalent to that of the original protein even if a small number of amino acid residues have been substituted, deleted, added, or inserted in the amino acid sequence of the protein. Therefore, a polypeptide having a sequence derived from the amino acid sequence of a CAPRIN-1 protein by substitution, deletion, and/or insertion of a small number of (preferably one or several) amino acid residues, having 80% or higher, 85-90% or higher, preferably 90% or higher, more preferably 95% or higher, further preferably 98% or higher, and still further preferably 99% or higher sequence identity with the original sequence, and specifically binding via an antigen-antibody reaction to an antibody against CAPRIN-1 (hereinafter, the same may be referred to as a "specifically reactive modified polypeptide" for convenience) can be used for cancer detection in a manner similar to the case of the polypeptides described above. Preferably, a specifically reactive modified polypeptide has an amino acid sequence derived from the amino acid sequence of a CAPRIN-1 protein by substitution, deletion, and/or insertion of one or several amino acid residues. The term "several" used herein refers to an integer of 2 to 10, preferably an integer of 2 to 6, and further preferably an integer of 2 to 4.

[0025] The term "sequence identity" used herein with reference to amino acid sequences is determined by aligning two amino acid sequences to be compared, so that as many amino acid residues match as possible, dividing the number of amino acid residues that match by the total number of amino acid residues, and then expressing the results in percentage terms (%). Upon the above alignment, gaps are inserted as appropriate into one or both of the sequences to be compared, according to need. Such sequence alignment can be performed using a well-known program or algorithm, such as BLAST, FASTA, or CLUSTAL W (Karlín and Altschul, Proc. Natl. Acad. Sci. U.S.A., 87: 2264-2268, 1993; Altschul et al., Nucleic Acids Res., 25: 3389-3402, 1997).

[0026] Twenty types of amino acids constituting naturally occurring proteins can be divided into groups of amino acids having properties analogous to each other: neutral amino acids having side chains with low polarity (Gly, Ile, Val, Leu, Ala, Met, and Pro); neutral amino acids having hydrophilic side chains (Asn, Gln, Thr, Ser, Tyr, and Cys); acidic amino acids (Asp and Glu); basic amino acids (Arg, Lys, and His); and aromatic amino acids (Phe, Tyr, Trp, and His). It is known that substitution among these amino acids (that is, conservative substitution) rarely alters the properties of the resulting polypeptide. When amino acid residues of CAPRIN-1 are to be substituted, accordingly, substitution is performed between members of the same group, so that the possibility of maintaining binding with the corresponding antibody becomes higher. In the present invention, however, the above variant may involve non-conservative substitution, as long as immunity-inducing activity equivalent to or almost equivalent to that of an unmodified polypeptide is imparted.

[0027] A polypeptide (hereinafter, which may be referred to as a "specifically reactive addition polypeptide" for convenience) that contains, as a partial sequence, the above polypeptide to be used in the present invention (e.g., prepared by addition of another (poly)peptide to one end or both ends of a polypeptide to be used in the present invention) and specifically binds via an antigen-antibody reaction to an antibody against CAPRIN-1 can also be used for detection of pancreatic cancer in a manner similar to the cases of the above polypeptides.

[0028] The polypeptides used in the present invention can be synthesized in accordance with a chemical synthesis method such as the Fmoc method (the fluorenylmethyloxycarbonyl method) or the tBoc method (the t-butyloxy-carbonyl method) (the Japanese Biochemical Society (ed.), Seikagaku Jikken Koza (Biochemical Experimental Lecture Series) 1, Tanpakushitsu no Kagaku (Protein Chemistry) IV, Kagaku Shushoku to Peptide Gousei (Chemical Modification and Peptide Synthesis), TOKYO KAGAKU DOZIN CO., LTD, Japan, 1981). Also, the polypeptides can be synthesized by a conventional method using various commercially available peptide synthesizers. Alternatively, the polypeptides can be easily prepared by known genetic engineering techniques (e.g., Sambrook et al., Molecular Cloning, 2nd Edition, Current Protocols in Molecular Biology, 1989, Cold Spring Harbor Laboratory Press; Ausubel et al., Short Protocols in Molecular Biology, 3rd Edition, A Compendium of Methods from Current Protocols in Molecular Biology, 1995, John Wiley & Sons). From RNA extracted from a tissue expressing a gene encoding the human CAPRIN-1 of SEQ ID NO: 2 or a homolog thereof, for example, cDNA of the gene is prepared via RT-PCR, the full-length sequence or a desired partial sequence of the cDNA is incorporated into an expression vector, and the vector is then introduced into a host cell. Thus, a polypeptide of interest can be obtained. The nucleotide sequences of cDNAs encoding canine CAPRIN-1 of SEQ ID NOs: 6, 8, 10, 12, and 14 are shown in SEQ ID NOs: 5, 7, 9, 11, and 13, respectively. The nucleotide sequences of human homologs thereof; that is, the cDNAs encoding human CAPRIN-1 of SEQ ID NOs: 2 and 4, are shown in SEQ ID NOs: 1 and 3, respectively. Accordingly, primers used for RT-PCR can be easily designed with reference to these nucleotide sequences. As described below, also, a gene encoding CAPRIN-1 of a non-human mammal can be amplified using primers designed with reference to the nucleotide sequences of the odd-numbered SEQ ID NOs: 1 to 29. Accordingly, cDNA encoding, for example, feline CAPRIN-1 can be easily prepared by techniques similar to the above techniques. RNA extraction, RT-PCR, incorporation of cDNA into a vector, and introduction of a vector into a host cell can be performed by, for example, well-known methods as described below. Also, vectors and host cells used herein are well known, and various vectors and host cells are commercially available.

[0029] The above host cells may be any cells, as long as they can express the above polypeptides. Examples of prokaryotic host cells include *Escherichia coli*. Examples of eukaryotic host cells include cultured mammalian cells, such

as monkey kidney cells (COS1), Chinese hamster ovary cells (CHO), the human embryonic kidney cell line (HEK293), and the mouse embryonic skin cell line (NIH3T3), budding yeast, fission yeast, silkworm cells, and Xenopus oocytes.

[0030] When prokaryotic cells are used as host cells, an expression vector having a replication origin in prokaryotic cells, a promoter, a ribosome-binding site, a multi-cloning site, a terminator, a drug-resistance gene, an auxotrophic complementary gene, and the like is used. Examples of expression vectors for *Escherichia coli* include pUC vectors, pBluescriptII, pET expression systems, and pGEX expression systems. DNA encoding the above polypeptide is incorporated into such an expression vector, prokaryotic host cells are transformed with the vector, and the thus obtained transformant is cultured. Thus, the polypeptide encoded by the DNA can be expressed in the prokaryotic host cells. At this time, the polypeptide can also be expressed as a fusion protein with another protein. DNA encoding the above polypeptide can be obtained by preparing cDNA by, for example, RT-PCR, as described above. Alternatively, such DNA can be synthesized by a conventional technique using a commercially available nucleic acid synthesizer, as described below. The nucleotide sequences of cDNAs of the genes encoding CAPRIN-1 of SEQ ID NOs: 2 and 4 are shown in SEQ ID NOs: 1 and 3 in the Sequence Listing, respectively.

[0031] When eukaryotic cells are used as host cells, an expression vector for eukaryotic cells having a promoter, a splicing region, a poly(A) additional site, and the like is used. Examples of such an expression vector include pKA1, pCDM8, pSVK3, pMSG, pSVL, pBK-CMV, pBK-RSV, an EBV vector, pRS, pcDNA3, and pYES2. As described above, DNA encoding a polypeptide used in the present invention is incorporated into such an expression vector, eukaryotic host cells are transformed with the vector, and the thus obtained transformant is cultured. Thus, the polypeptide encoded by the above DNA can be expressed in eukaryotic host cells. When pIND/V5-His, pFLAG-CMV-2, pEGFP-N1, pEGFP-C1, or the like is used as an expression vector, the above polypeptide can be expressed as a fusion protein with various tags, such as His tags (e.g., (His)₆ to (His)₁₀), a FLAG tag, a myc tag, an HA tag, or GFP.

[0032] An expression vector can be introduced into a host cell in accordance with a well-known technique, such as electroporation, a calcium phosphate method, a liposome method, a DEAE dextran method, microinjection, viral infection, lipofection, or binding with a cell-membrane-permeable peptide.

[0033] A polypeptide of interest can be isolated and purified from host cells using known isolation techniques in combination. Examples of such techniques include treatment using a denaturing agent such as urea or a surfactant, ultrasonication, enzymatic digestion, salting-out, solvent fractionation and precipitation, dialysis, centrifugation, ultrafiltration, gel filtration, SDS-PAGE, isoelectric focusing, ion exchange chromatography, hydrophobic chromatography, affinity chromatography, and reverse phase chromatography.

[0034] Polypeptides obtained by the above methods include polypeptides in the form of fusion proteins with any other proteins. Examples of such fusion proteins include a fusion protein with glutathione-S-transferase (GST) and a fusion protein with a His tag. Polypeptides in the form of such fusion proteins are also within the scope of the above described specifically reactive addition polypeptides, and such polypeptides can be used for the first detection method of the present invention. Further, polypeptides expressed in transformed cells may occasionally be subjected to various types of modification within cells after translation. Such a post-translationally modified polypeptide can be used in the first detection method of the present invention, as long as it is capable of specifically binding, via an antigen-antibody reaction, to an antibody against a CAPRIN-1 protein. Examples of such post-translational modification include the removal of N-terminal methionine, N-terminal acetylation, glycosylation, limited proteolysis by intracellular protease, myristoylation, isoprenylation, and phosphorylation.

[0035] An antibody in a sample can be easily measured by immunoassay using the above polypeptide as an antigen. Immunoassay itself is well known in the art. Immunoassay is classified into the sandwich method, the competition method, the agglutination method, the Western blot method, and the like based on types of reactions. Also, immunoassay is classified based on labels into radioimmunoassay, fluorescence immunoassay, enzyme immunoassay, and biotin immunoassay, for example. Immunoassay of the above antibody can be performed using any of these methods. Sandwich ELISA or the agglutination method are preferably employed as an immunoassay technique for the above antibody in the method of the present invention, since the procedures of these methods are convenient and require no extensive apparatus and the like, although techniques are not limited thereto. When an enzyme is used as a label for an antibody, such enzyme is not particularly limited, as long as it satisfies conditions such that: the turnover number is high; it remains stable even if it is bound to an antibody; and it specifically causes the color development of the substrate. Enzymes that can be used for general enzyme immunoassay, such as peroxidase, β -galactosidase, alkaline phosphatase, glucose oxidase, acetylcholine esterase, glucose-6-phosphorylation dehydrogenase, and malic acid dehydrogenase, can be used. In addition, enzyme-inhibiting substances, coenzymes, and the like can be used. Binding of these enzymes with antibodies can be performed by known methods involving the use of a cross-linking agent, such as a maleimide compound, the biotin-(strept)avidin system, or the like. As a substrate, a known substance can be used depending on the type of an enzyme to be used. When peroxidase is used as an enzyme, for example, 3,3',5,5'-tetramethylbenzidine can be used. When alkaline phosphatase is used, for example, para-nitrophenol can be used. A radioisotope that is generally used for radioimmunoassay, such as ¹²⁵I or ³H, can be used. A fluorescent dye that is used for general fluorescent antibody techniques, such as fluorescence isothiocyanate (FITC), tetramethylrhodamine isothiocyanate (TRITC), or a

cyanine fluorescence dye (e.g., Cy3 or Cy5), can be used.

5 [0036] There is no need to explain the above immunoassay techniques herein since these techniques are well known; however, briefly, for example, the sandwich method comprises immobilizing the above polypeptide used as an antigen on a solid phase, allowing the polypeptide to react with a sample such as serum, washing, allowing an appropriate
10 secondary antibody to react with an antibody from the sample, again washing, and then measuring the secondary antibody bound to the solid phase. By immobilizing an antigenic polypeptide on a solid phase, an unbound secondary antibody can be easily removed. Accordingly, it is preferable as an embodiment of the method for detecting cancer of the present invention. As a secondary antibody, an anti-canine IgG antibody, for example, can be used if a sample is from a dog. A secondary antibody is labeled in advance with a labeling substance exemplified above, so that the secondary
15 antibody bound to a solid phase can be measured. The amount of the secondary antibody thus measured corresponds to the amount of the above antibody in the serum sample. When an enzyme is used as a labeling substance, the amount of the antibody can be measured by adding a substrate that is degraded in order to develop color by enzymatic action and then optically measuring the amount of the substrate degraded. When a radioisotope is used as a labeling substance, the amount of radiation from the radioisotope can be measured using a scintillation counter or the like.

20 [0037] In the second method of the present invention, CAPRIN-1 that can be contained in a sample obtained from a subject is measured. As described above, the amount of an antibody that undergoes an antigen-antibody reaction with CAPRIN-1 of a dog, a human, or the like is significantly higher in subjects with pancreatic cancer, compared with healthy subjects. This indicates that the amount of CAPRIN-1 accumulated as an antigen is significantly high in pancreatic cancer cells. In the case of healthy subjects, the CAPRIN-1 expression level is below the detection limit, or CAPRIN-1
25 expression in tissue is weak and it might occur merely within cells. Pancreatic cancer can also be detected by directly measuring CAPRIN-1, as specifically described in the examples below. Therefore, pancreatic cancer can be detected in a subject by measuring CAPRIN-1 itself, as in the case of the first method.

30 [0038] A polypeptide in a sample can be easily measured by well-known immunoassay techniques. Specifically, an antibody that undergoes an antigen-antibody reaction with CAPRIN-1 or an antigen-binding fragment thereof is prepared, and immunoassay is carried out using the same. Thus, the presence of CAPRIN-1 in the sample can be measured. As described above, an antibody has cross-reactivity. With the use of an antibody that undergoes an antigen-antibody
35 reaction with canine CAPRIN-1 of SEQ ID NO: 6 or an antigen-binding fragment thereof, accordingly, not only the canine CAPRIN-1 of SEQ ID NO: 6, but also its homologs in other mammals (e.g., the human CAPRIN-1 of SEQ ID NO: 2 or 4) can be measured. The immunoassay technique itself is a well-known conventional technique, as described above.

40 [0039] This study reveals that CAPRIN-1 is a cell membrane protein that is expressed on the surface of pancreatic cancer cells. A subject with cancer contains many proteases in cancer tissues. Accordingly, the portion of the CAPRIN-1 sequence expressed outside the cancer cells is degraded and separated from the cancer cells, and such portion is larger in amount than the portion of the CAPRIN-1 sequence expressed in the cancer cells. If an antibody capable of binding to the surfaces of pancreatic cancer cells is used as an antibody against CAPRIN-1 in the measurement, or if
45 an antigen-binding fragment thereof is used, accordingly, a larger amount of CAPRIN-1 can be detected, and pancreatic cancer can be diagnosed with higher sensitivity.

50 [0040] In the present invention, accordingly, use of an antibody that binds to a portion expressed on the surface of a pancreatic cancer cell of a CAPRIN-1 protein molecule is preferable. An example of a partial peptide of a CAPRIN-1 protein expressed on the surface of a pancreatic cancer cell is a polypeptide consisting of an amino acid sequence of 7 to 12 or more continuous amino acid residues within the region of amino acid residues (aa) 50 to 98 or amino acid
55 residues (aa) 233 to 305 in any of the amino acid sequences represented by even-numbered SEQ ID NOs: 2 to 30 in the Sequence Listing, excluding SEQ ID NO: 6 and SEQ ID NO: 18. A specific example thereof is, but is not limited to, the amino acid sequence represented by SEQ ID NO: 43 or SEQ ID NO: 61 (in the amino acid sequence represented by SEQ ID NO: 61, a region of the amino acid sequence represented by SEQ ID NO: 62 or SEQ ID NO: 63 is preferable) or an amino acid sequence having 80% or higher, preferably 85% or higher, more preferably 90% or higher, and further preferably 95% or higher sequence identity with the relevant amino acid sequence. In addition, all antibodies binding to these polypeptides fall within the scope of the antibodies used in the present invention. Specific examples include an antibody binding to a polypeptide comprising the amino acid sequence represented by SEQ ID NO: 43 or antigen-binding
60 fragment thereof, a monoclonal antibody having the amino acid sequences represented by SEQ ID NOs: 44 and 45 or an antigen-binding fragment thereof, a monoclonal antibody having the amino acid sequences represented by SEQ ID NOs: 44 and 46 or an antigen-binding fragment thereof, a monoclonal antibody having the amino acid sequences represented by SEQ ID NOs: 44 and 47 or an antigen-binding fragment thereof, a monoclonal antibody having the amino acid sequences represented by SEQ ID NOs: 44 and 48 or an antigen-binding fragment thereof, a monoclonal antibody having the amino acid sequences represented by SEQ ID NOs: 49 and 50 or an antigen-binding fragment thereof, a monoclonal antibody having the amino acid sequences represented by SEQ ID NOs: 51 and 52 or an antigen-binding
65 fragment thereof, a monoclonal antibody having the amino acid sequences represented by SEQ ID NOs: 53 and 54 or an antigen-binding fragment thereof, a monoclonal antibody having the amino acid sequences represented by SEQ ID NOs: 55 and 56 or an antigen-binding fragment thereof, a monoclonal antibody having the amino acid sequences

represented by SEQ ID NOs: 57 and 58 or an antigen-binding fragment thereof, and a monoclonal antibody having the amino acid sequences represented by SEQ ID NOs: 59 and 60 or an antigen-binding fragment thereof.

5 [0041] The term "antigen-binding fragment" used herein refers to an antibody fragment capable of binding to an antigen, such as an Fab fragment, an F(ab)₂ fragment, or an Fv fragment, contained in an antibody molecule. An antibody may be a polyclonal antibody or a monoclonal antibody. For immunoassay, a monoclonal antibody with high reproducibility is preferable. Methods for preparing a polyclonal antibody and a monoclonal antibody using a polypeptide as an immunogen are well known and can be easily performed in a conventional manner. For example, an animal is immunized with CAPRIN-1 or a fragment thereof alone or CAPRIN-1 or a fragment thereof bound to a carrier protein, such as keyhole limpet hemocyanin (KLH), casein, or serum albumin, as an immunogen, together with an adjuvant, and an antibody against CAPRIN-1 can then be induced. Antibody-producing cells, such as splenocytes or lymphocytes, collected from the immunized animal are fused to myeloma cells to prepare hybridomas, and hybridomas producing an antibody that binds to CAPRIN-1 are selected and then grown. Thus, a monoclonal antibody whose corresponding antigen is CAPRIN-1 can be obtained from the culture supernatant. The method described above is a well-known conventional method.

10 [0042] In the third method of the present invention, a nucleic acid encoding CAPRIN-1 (e.g., mRNA or cDNA synthesized from mRNA) that can be contained in a sample obtained from a living organism is measured. As specifically described in the examples below, a nucleic acid encoding the canine CAPRIN-1 of SEQ ID NO: 6, 8, 10, 12, or 14 or human CAPRIN-1 of SEQ ID NO: 2 or 4 is expressed at a significantly high level in pancreatic cancer cells. Therefore, a cancer existing in the living body can be detected by measuring such nucleic acid in a sample.

15 [0043] mRNA in a sample can be quantitatively measured by, for example, a conventional method, such as real-time detection RT-PCR using the mRNA as a template. Such mRNA can generally be quantitatively measured based on staining intensity or the like in the conventional Northern blot method. The cDNA sequences encoding CAPRIN-1 of the even-numbered SEQ ID NOs: 2 to 30 are represented by the odd-numbered SEQ ID NOs: 1 to 29, respectively. Accordingly, a polynucleotide specifically hybridizing to a partial region in the nucleotide sequence represented by any of the odd-numbered SEQ ID NOs: 1 to 29 (hereinafter, referred to as a "polynucleotide for cancer detection") is prepared based on these sequences, and such polynucleotide is used as a probe or a primer for nucleic acid amplification to determine the amount of the mRNA in a sample. If a polynucleotide is capable of specifically hybridizing to a partial region in the nucleotide sequence represented by any of the odd-numbered SEQ ID NOs: 1 to 29, mRNA encoding CAPRIN-1 in mammals other than dogs and humans can also be measured, as described later. In the present invention, a polynucleotide may be RNA or DNA.

20 [0044] The term "specifically hybridizing to" used herein refers to a situation in which said polynucleotide hybridizes only to a target partial region and does not substantially hybridize to any other regions under stringent hybridization conditions.

25 [0045] The term "under stringent hybridization conditions" used herein refers to conditions employed for annealing in general PCR or detection using a probe. In the case of PCR using Taq polymerase, for example, a reaction is performed at an appropriate annealing temperature ranging from about 54°C to 60°C using a general buffer, such as a buffer containing 50 mM KCl, 10 mM Tris-HCl (pH 8.3 to 9.0), and 1.5 mM MgCl₂. In the case of Northern hybridization, for example, a reaction is performed using a general hybridization solution such as 5x SSPE, 50% formamide, 5x Denhardt's solution, and 0.1-0.5% SDS, or 0.1-5x SSC and 0.1-0.5% SDS at an appropriate hybridization temperature ranging from about 42-65°C. After hybridization, washing is performed with, for example, 0.1-0.2x SSC and 0.1% SDS. Appropriate annealing temperatures or hybridization temperatures are not limited to the above examples, and such temperatures are determined based on the T_m value for a polynucleotide for cancer detection that is used as a primer or a probe and the empirical rules of experimenters. A person skilled in the art can easily determine such temperature range.

30 [0046] The expression "does not substantially hybridize to" used herein refers to a situation in which said polynucleotide does not hybridize to a target partial region at all or a extremely low amount of the polynucleotide hybridizes to a target partial region, that is, in a relatively negligible amount, even when it hybridizes to a target partial region. An example of the polynucleotide specifically hybridizing under such conditions is a polynucleotide having a certain level or higher of a sequence identity with the nucleotide sequence of a target partial region. For example, such polynucleotide has 70% or higher, preferably 80% or higher, more preferably 85% or higher, further preferably 90% or higher, even further preferably 93% or higher, still further preferably 95% or higher, and particularly preferably 98% or higher sequence identity. Most preferably, the polynucleotide has a nucleotide sequence identical to the nucleotide sequence of a target partial region. Sequence identity is defined in the same manner as the sequence identity of the amino acid sequence described above. Even if a terminus of a polynucleotide for cancer detection contains a region that does not hybridize thereto, in the case of a probe, it can be used for detection as long as a hybridizing region occupies about a half or more of the entire probe. In the case of a primer, it can be used for detection as long as a hybridizing region occupies about a half or more of the entire primer and is located on the 3' terminal side, since this allows a normal annealing and extension reaction to take place. When a terminus of a polynucleotide for cancer detection contains a non-hybridizing region, as described above, sequence identity with a target nucleotide sequence is calculated focusing only on the

hybridizing region without taking a non-hybridizing region into consideration.

[0047] In the present invention, the term "partial sequence" (or "partial region") refers to a part of a nucleotide sequence represented by any of the odd-numbered SEQ ID NOs: 1 to 29. Specifically, the partial sequence comprises at least 15 to 19 continuous nucleotides, preferably 18 or more continuous nucleotides, more preferably at least 20 or 25 continuous nucleotides, and further preferably at least 30, 40, or 50 continuous nucleotides. The expression "the nucleotide sequence represented by SEQ ID NO: 5" used herein refers to, in addition to the nucleotide sequence actually shown in SEQ ID NO: 5, a sequence complementary thereto. Accordingly, the expression "a polynucleotide having the nucleotide sequence represented by SEQ ID NO: 5" refers to, for example, a single-stranded polynucleotide having the nucleotide sequence actually represented by SEQ ID NO: 5, a single-stranded polynucleotide having a nucleotide sequence complementary to the nucleotide sequence represented by SEQ ID NO: 5, or a double-stranded polynucleotide consisting of the two single-stranded polynucleotides. When a polynucleotide to be used in the present invention is prepared or a polynucleotide encoding a polypeptide to be used in the present invention is prepared, any of the nucleotide sequences is appropriately selected, and a person skilled in the art can readily perform such selection.

[0048] The number of nucleotides in a polynucleotide for cancer detection is preferably 18 or more in view of ensuring specificity. When the polynucleotide is used as a probe, it preferably comprises 18 or more nucleotides, and it further preferably comprises 20 nucleotides to the full length of the coding region. When the polynucleotide is used as a primer, it preferably comprises 18 to 50 nucleotides. A preferable example of the polynucleotide for cancer detection is a polynucleotide comprising 18 or more continuous nucleotides in a nucleotide sequence represented by any of the odd-numbered SEQ ID NOs: 1 to 29.

[0049] It is apparent to a person skilled in the art who refers to the description of the present invention that: a polynucleotide specifically hybridizing to a partial region in SEQ ID NO: 5, 7, 9, 11, or 13 is used for measuring the amount of a nucleic acid (e.g., mRNA or cDNA synthesized from mRNA) encoding the canine CAPRIN-1 protein of SEQ ID NO: 6, 8, 10, 12, or 14, respectively; and a polynucleotide specifically hybridizing to a partial region in SEQ ID NO: 1 or 3 is used for measuring the amount of a nucleic acid (e.g., mRNA or cDNA synthesized from mRNA) encoding the human CAPRIN-1 protein of SEQ ID NO: 2 or 4, respectively. However, a protein from a given mammal and a homolog thereof from another mammal generally share a high sequence identity even at the nucleotide sequence level. Thus, the sequence identity among the nucleotide sequences of SEQ ID NOs: 1 to 13 is also as high as 94% to 100%. Accordingly, a polynucleotide specifically hybridizing to a partial region of the sequence of SEQ ID NO: 5 can also specifically hybridize to a partial region corresponding to the partial region of any of the odd-numbered SEQ ID NOs: 1 to 29.

[0050] In fact, a pair of primers having the nucleotide sequences represented by SEQ ID NO: 33 and 34 specifically hybridize to both a partial region of any of the odd-numbered SEQ ID NOs: 1 to 29 and a partial region of the sequence of SEQ ID NO: 5, as described in the examples below. Thus, both mRNA encoding the canine CAPRIN-1 of SEQ ID NO: 6 and mRNA encoding a homolog thereof can be measured. With the use of a polynucleotide specifically hybridizing to a partial region of the sequence of SEQ ID NO: 5, accordingly, not only mRNA encoding the canine CAPRIN-1 of SEQ ID NO: 6 but also mRNA encoding the human CAPRIN-1 of SEQ ID NO: 2 or 4 can be measured. Similarly, mRNA encoding CAPRIN-1 of another mammal such as a cat can also be measured. When a polynucleotide for cancer detection is designed, it is more desirable to select partial regions having particularly high sequence identity from among the odd-numbered SEQ ID NOs: 1 to 29 (and identical nucleotide sequences are preferable). If there is a particularly high sequence identity having partial region between canine CAPRIN-1 and human CAPRIN-1, a region showing very high sequence identity with such region is expected to be also present in a homolog gene of another animal species. Through selection of such a partial region, accuracy for measuring mRNA encoding CAPRIN-1 of an animal species other than dogs or humans can be increased.

[0051] A method for measuring a nucleic acid in test object using a polynucleotide specifically hybridizing to a partial region of the nucleic acid as a probe or a primer(s) for nucleic acid amplification method such as PCR is well known. Examples of such method include, in addition to RT-PCR as specifically described in the examples below, Northern blot and *in situ* hybridization. When the amount of mRNA is measured in the present invention, any such well-known measurement method can be employed.

[0052] A nucleic acid amplification method such as PCR is well known in the art, and reagent kits and apparatuses used therefor are commercially available, so that the method can be easily performed. Specifically, denaturation, annealing, and extension steps are each performed using a nucleic acid in test object (e.g., the cDNA of a gene encoding a protein having an amino acid sequence represented by any of the even-numbered SEQ ID NOs: 2 to 30) as a template and a pair of polynucleotides (primers) for cancer detection in a known buffer in the presence of thermostable DNA polymerase such as Taq polymerase or Pfu polymerase and dNTPs (here, N = A, T, C, and G) by varying the temperature of the reaction solution in each step. In general, the denaturation step is performed at 90°C to 95°C, the annealing step is performed at or near the T_m of the template and the primers (preferably within $\pm 4^\circ\text{C}$), and the extension step is performed at 72°C, which is the optimum temperature for thermostable DNA polymerase such as Taq polymerase or Pfu polymerase, or a temperature near the optimum temperature. The duration of each step is adequately set to between about 30 seconds and 2 minutes. This heating cycle is repeated about 25 to 40 times, for example, so that the template

nucleic acid region sandwiched between a pair of primers is amplified. The nucleic acid amplification method is not limited to PCR, and any other nucleic acid amplification methods well known in the art can be employed. When a nucleic acid amplification method is performed using a pair of polynucleotides for cancer detection as primers and a nucleic acid in test object as a template, as described above, the nucleic acid is amplified. If a sample does not contain the test nucleic acid, however, amplification does not take place. Accordingly, an amplification product may be detected so as to determine the presence or absence of the nucleic acid in the sample. An amplification product can be detected by a method that comprises subjecting a reaction solution after amplification to electrophoresis and then staining the band with ethidium bromide or the like or a method that comprises immobilizing an amplification product after electrophoresis on a solid phase such as a nylon membrane, performing hybridization with a labeling probe that specifically hybridizes to a nucleic acid, washing, and then detecting the label. Also, so-called real-time detection PCR is performed using a quencher fluorescent dye and a reporter fluorescent dye, and the amount of a nucleic acid in a specimen can thus be quantified. Since kits for real-time detection PCR are commercially available, real-time detection PCR can be easily performed. Further, semi-quantitative measurement of a nucleic acid in test object can also be carried out based on electrophoresis band intensity. A nucleic acid in test object may be either mRNA or cDNA reversely transcribed from mRNA. When mRNA is amplified as a nucleic acid, a NASBA method (the 3SR method or TMA method) using the above pair of primers can also be employed. The NASBA method is well known, and kits therefor are also commercially available, so that the method can be easily performed using the above pair of primers.

[0053] As a probe, a labeled probe that is prepared by labeling a polynucleotide for cancer detection with a fluorescent label, a radiolabel, a biotin label, or the like can be used. Methods for labeling a polynucleotide are well known. The presence or absence of a nucleic acid in a sample can be examined by immobilizing a nucleic acid or an amplification product thereof, performing hybridization with a labeled probe, washing, and then measuring the label bound to the solid phase. Alternatively, a polynucleotide for cancer detection is immobilized, a nucleic acid in test object is hybridized thereto, and the test nucleic acid bound to the solid phase can then be detected using the labeled probe or the like. In such a case, a polynucleotide for cancer detection bound to a solid phase is also referred to as a "probe." Methods for measuring a nucleic acid using a polynucleotide probe are also well known in the art. Such a method can be performed by, in a buffer, bringing a polynucleotide probe into contact with a nucleic acid in test object at T_m or near T_m (preferably, $\pm 4^\circ\text{C}$) for hybridization, washing, and then measuring the labeled probe hybridized or the template nucleic acid bound to the solid-phase probe. Examples of such method include well-known methods such as Northern blotting, *in situ* hybridization, and Southern blotting. In the present invention, any well-known method is applicable.

[0054] According to the detection method of the present invention, whether or not a subject animal (or a subject) is afflicted with pancreatic cancer is evaluated based on the presence or amount of CAPRIN-1 expression measured as described above. While pancreatic cancer can be detected only by measuring the presence or amount of CAPRIN-1 expression in a subject animal, it is preferable that the expression levels (the antibody level, polypeptide level, or mRNA level) of CAPRIN-1 in one or more samples of healthy subjects is examined and the determined value of a subject animal is compared with the standard value obtained from healthy subjects, in view of enhancing detection accuracy. To further enhance detection accuracy, CAPRIN-1 expression levels are measured for samples obtained from many patients found to have pancreatic cancer, so as to obtain a standard value for pancreatic cancer patients, and the determined value of a subject animal may then be compared with both the standard value for healthy subjects and the standard value for pancreatic cancer patients. The above standard values can be determined by, for example, quantifying the CAPRIN-1 expression level in each sample and calculating the mean value thereof. The standard value for healthy subjects and the same for pancreatic cancer patients can be determined in advance by measuring CAPRIN-1 expression levels in many healthy subjects and pancreatic cancer patients. When comparison with the standard value is performed in the method of the present invention, accordingly, a standard value determined in advance may be used.

[0055] The detection method of the present invention may comprise diagnosis based on other cancer antigens or cancer markers in combination. This can further enhance the accuracy of pancreatic cancer detection. When an antibody specifically existing in pancreatic cancer patients is measured by the method of the present invention, for example, another polypeptide that is often expressed in a cancer tissue can be used in combination as an antigen in a manner similar to that used for polypeptides described above. Also, the method of the present invention may be performed in combination with diagnosis using a previously known cancer marker.

[0056] Pancreatic cancer to be subjected to the method for detecting pancreatic cancer of the present invention is pancreatic cancer expressing CAPRIN-1. Examples of such cancer include, but are not limited to, pancreatic ductal carcinoma, invasive pancreatic ductal carcinoma, adenocarcinoma, acinar cell carcinoma, adenosquamous carcinoma, giant cell tumor, intraductal papillary-mucinous neoplasm (IPMN), mucinous cystic neoplasm (MCN), pancreatoblastoma, serous cystadenocarcinoma, solid-pseudopapillary tumor (SPT), gastrinomas (Zollinger-Ellison syndrome), glucagonomas, insulinomas, multiple endocrine neoplasia Type-1 (MEN1) (Wermer syndrome), nonfunctional islet cell tumor, somatostatinomas, and VIPomas. A subject in the method of the present invention is a mammal, preferably a human or a dog.

[0057] Examples of samples to be subjected to the method of the present invention include body fluids, such as blood,

serum, blood plasma, ascites fluid, and pleural effusion, tissues, and cells. In the first method and the second method, in particular, serum, blood plasma, ascites fluid, pleural effusion, tissue, and cell samples can be preferably used. In the third method comprising measuring a nucleic acid such as mRNA, tissue and cell samples are preferable.

[0058] One or more polypeptides to be used as antigens for immunoassay in the first method described above (i.e., the canine CAPRIN-1 of SEQ ID NO: 2 and a homolog thereof, a specifically reactive partial polypeptide, a specifically reactive modified polypeptide, and a specifically reactive addition polypeptide) can be provided as reagents or kits for detection of pancreatic cancer. Such a reagent may consist of the above polypeptide, or it may contain various additives useful for stabilization of the polypeptide, a buffer necessary for assay, secondary antibodies, substrates for enzymes, or the like, separately. Alternatively, such a reagent can be immobilized on a solid phase such as a plate or membrane. Preferable examples of such polypeptides are given above.

[0059] An antibody or an antigen-binding fragment thereof, which undergoes an antigen-antibody reaction with CAPRIN-1, used for immunoassays of CAPRIN-1 by the second method can also be provided in the form of a reagent for pancreatic cancer detection. The reagent for pancreatic cancer detection may consist of the above antibody or an antigen-binding fragment thereof. The reagent may contain various additives useful for stabilization of such antibody or an antigen-binding fragment thereof. Alternatively, a metal, such as manganese or iron, may be bound to the antibody or an antigen-binding fragment thereof. When such metal-bound antibody or an antigen-binding fragment thereof is administered to a living organism, the metal-bound antibody or antigen-binding fragment thereof is accumulated at an increased level at a site at which the antigen protein is present at a higher level. When such metal-bound antibody or antigen-binding fragment thereof is administered to a living organism, the metal-bound antibody or antigen-binding fragment thereof is accumulated at an increased level at a site at which the antigen protein is present at a higher level. Therefore, the metal is measured by MRI or the like, and the presence of cancer cells producing the antigen protein can be thus detected.

[0060] Furthermore, one or more the above polynucleotides for pancreatic cancer detection to be used for measuring a nucleic acid such as mRNA in the third method can also be provided as a reagent or kit for pancreatic cancer detection. In such a case, the reagent for pancreatic cancer detection may consist of the polynucleotide, or it may contain various additives useful for stabilization of the polypeptide, a buffer necessary for assay (e.g., a fluorescent label), and the like, separately. The polynucleotide for pancreatic cancer detection contained in the reagent is preferably a primer(s) or a probe(s). Conditions and preferable examples of the polynucleotide for pancreatic cancer detection are as described above.

Examples

[0061] The present invention will be described in more detail with reference to the following examples, although the technical scope of the present invention is not limited to the examples.

[Example 1] Obtaining pancreatic cancer antigenic protein by SEREX method (1) Construction of cDNA library

[0062] Total RNA was extracted from a testis tissue of a healthy dog by the acid guanidium-phenol-chloroform method, and poly A RNA was purified using an Oligotex-dT30 mRNA purification kit (Takara Shuzo, Co., Ltd.) in accordance with the protocols attached to the kit.

[0063] A canine testis cDNA phage library was synthesized using the thus obtained mRNA (5 μ g). The cDNA phage library was constructed using cDNA Synthesis Kit, ZAP-cDNA Synthesis Kit, and ZAP-cDNA GigapackIII Gold Cloning Kit (STRATAGENE) in accordance with the protocols attached to the kits. The size of the thus constructed cDNA phage library was 7.73×10^5 pfu/ml.

(2) Screening of cDNA library using serum

[0064] Immunoscreening was performed using the canine testis cDNA phage library constructed above. Specifically, host *Escherichia coli* (XL1-Blue MRF') was infected with the phage on an NZY agarose plate ($\Phi 90 \times 15$ mm) so as to obtain 2,210 clones. *E. coli* cells were cultured at 42°C for 3 to 4 hours to form plaques. The plate was covered with a nitrocellulose membrane (Hybond C Extra: GE Healthcare Bio-Science) impregnated with IPTG (isopropyl- β -D-thiogalactoside) at 37°C for 4 hours, so that the protein was induced to express and then transferred to the membrane. Thereafter, the membrane was collected and then soaked in TBS (10 mM Tris-HCl, 150 mM NaCl, pH 7.5) containing 0.5% powdered skim milk, followed by shaking at 4°C overnight, so as to suppress nonspecific reactions. The filter was subjected to a reaction with a 500-fold diluted serum of a afflicted dog r at room temperature for 2 to 3 hours.

[0065] As the above serum of the afflicted dog, a serum collected from a canine afflicted with pancreatic cancer was used. The serum was stored at -80°C and then subjected to pre-treatment immediately before use. A method for serum pretreatment is as follows. Specifically, host *Escherichia coli* (XL1-Blue MRF') was infected with a λ ZAP Express phage

into which no foreign gene had been inserted, and culture was conducted overnight on a NZY plate medium at 37°C. Subsequently, buffer (0.2 M NaHCO₃, pH 8.3) containing 0.5 M NaCl was added to the plate, the plate was allowed to stand at 4°C for 15 hours, and a supernatant was then collected as an *Escherichia coli*/phage extract. The thus collected *Escherichia coli*/phage extract was then applied to an NHS-column (GE Healthcare Bio-Science), so that an *Escherichia coli*/phage-derived protein was immobilized. The serum of the afflicted dog was applied to the protein-immobilized column for reaction and *Escherichia coli* and an antibody adsorbed to the phage were then removed from the serum. The serum fraction that had passed through the column was diluted 500-fold with TBS containing 0.5% powdered skim milk. The diluted serum fraction was used as an immunoscreening material.

[0066] A membrane onto which the treated serum and the above fusion protein had been blotted was washed 4 times with TBS-T (0.05% Tween 20/TBS), and the membrane was then allowed to react with goat anti-canine IgG (Goat anti-Dog IgG-h+I HRP conjugated, BETHYL Laboratories) diluted 5000-fold with TBS containing 0.5% powdered skim milk as a secondary antibody at room temperature for 1 hour. Detection was performed via an enzymatic color development reaction using the NBT/BCIP reaction solution (Roche). Colonies that matched sites positive for the color development reaction were collected from the NZY agarose plate (Φ90 x 15 mm) and then dissolved in 500 μl of an SM buffer (100 mM NaCl, 10 mM MgClSO₄, 50 mM Tris-HCl, 0.01 % gelatin, pH 7.5). Until colonies positive for color development reaction were unified, secondary screening and tertiary screening were repeated by a method similar to the above, so that 30,940 phage clones reacting with the serum IgG were screened. Thus, 5 positive clones were isolated.

(3) Homology search for isolated antigen gene

[0067] For nucleotide sequence analysis of the 5 positive clones isolated by the above method, a procedure for conversion from phage vectors to plasmid vectors was performed. Specifically, 200 μl of a solution containing host *Escherichia coli* (XL1-Blue MRF') at the absorbance (OD₆₀₀) of 1.0 was prepared. The solution was mixed with 250 μl of a purified phage solution and 1 μl of ExAssist helper phage (STRATAGENE), the mixture was subjected to a reaction at 37°C for 15 minutes, 3 ml of LB medium was added thereto, and culture was then performed at 37°C for 2.5 to 3 hours. Immediately thereafter, the temperature of the solution was kept in a water bath at 70°C for 20 minutes, centrifugation was performed at 4°C and 1000 x g for 15 minutes, and the supernatant was then collected as a phagemid solution. Subsequently, 200 μl of a solution containing phagemid host *Escherichia coli* (SOLR) at the absorbance OD₆₀₀ of 1.0 was prepared. The resulting solution was mixed with 10 μl of a purified phage solution, followed by a reaction at 37°C for 15 minutes. The reaction product (50 ml) was seeded on LB agar medium containing ampicillin (final concentration: 50 μg/ml), and culture was conducted at 37°C overnight. Transformed SOLR single colony was collected and then cultured in LB medium containing ampicillin (final concentration: 50 μg/ml) at 37°C. Thereafter, plasmid DNA containing an insert of interest was purified using the QIAGEN plasmid Miniprep Kit (QIAGEN).

[0068] The purified plasmid was subjected to analysis of the full-length insert sequence by the primer walking method using the T3 primer represented by SEQ ID NO: 31 and the T7 primer represented by SEQ ID NO: 32. As a result of sequence analysis, the gene sequences represented by SEQ ID NOs: 5, 7, 9, 11, and 13 were obtained. A homology search program, BLAST search (<http://www.ncbi.nlm.nih.gov/BLAST/>), was performed using the nucleotide sequences and amino acid sequences (SEQ ID NOs: 6, 8, 10, 12, and 14) of the genes. As a result of this homology search with known genes, all of the 5 obtained genes were found to encode CAPRIN-1. The sequence identity among the 5 genes was 100% for nucleotide sequence and 99% for amino acid sequence in regions translated into proteins. Also, the sequence identity between the canine gene (any of SEQ ID NO: 5, 7, 9, 11, or 13) and a gene encoding a human homolog thereof was 94% for nucleotide sequence and 98% for amino acid sequence in regions translated into proteins. The nucleotide sequences of the human homolog are represented by SEQ ID NOs: 1 and 3 and the amino acid sequences of the same are represented by SEQ ID NOs: 2 and 4. Also, the sequence identity between the obtained canine gene and a gene encoding a cattle homolog was 94% for nucleotide sequence and 97% for amino acid sequence in regions translated into proteins. The nucleotide sequence of the cattle homolog is represented by SEQ ID NO: 15 and the amino acid sequence of the same is represented by SEQ ID NO: 16. The sequence identity between the gene encoding the human homolog and the gene encoding the cattle homolog was 94% for nucleotide sequences and ranged from 93% to 97% for amino acid sequence in regions translated into proteins. Also, the sequence identity between the obtained canine gene and a gene encoding an equine homolog was 93% for nucleotide sequence and 97% for amino acid sequence in regions translated into proteins. The nucleotide sequence of the equine homolog is represented by SEQ ID NO: 17 and the amino acid sequence of the same is represented by SEQ ID NO: 18. The sequence identity between the gene encoding the human homolog and the gene encoding the equine homolog was 93% for nucleotide sequence and 96% for amino acid sequence in regions translated into proteins. Also, the sequence identity between the obtained canine gene and a gene encoding the mouse homolog ranged from 87% to 89% in terms of nucleotide sequence and ranged from 95% to 97% for amino acid sequence in regions translated into proteins. The nucleotide sequences of the mouse homolog are represented by SEQ ID NOs: 19, 21, 23, 25, and 27 and the amino acid sequences of the same are represented by SEQ ID NOs: 20, 22, 24, 26, and 28. The sequence identity between the gene encoding the human

homolog and the gene encoding the mouse homolog ranged from 89% to 91% for nucleotide sequence and ranged from 95% to 96% for amino acid sequence in regions translated into proteins. Also, the sequence identity between the obtained canine gene and a gene encoding a chicken homolog was 82% for nucleotide sequence and 87% for amino acid sequence in regions translated into proteins. The nucleotide sequence of the chicken homolog is represented by SEQ ID NO: 29 and the amino acid sequence of the same is represented by SEQ ID NO: 30. The sequence identity between the gene encoding the human homolog and the gene encoding the chicken homolog ranged from 81 % to 82% for nucleotide sequence and 86% for amino acid sequence in regions translated into proteins.

(4) Gene expression analysis in human pancreatic cancer cell lines

[0069] Expression of the genes obtained by the above method in human normal tissues (i.e., mammary gland, brain, bone marrow, lung, esophagus, pancreas, and testis) and 4 types of pancreatic cancer cell lines (i.e., Capan-2, MIAPaCa-2, PANC-1, and BxPC-3) was examined by RT-PCR (reverse transcription-PCR). A reverse transcription reaction was performed as follows. Specifically, total RNA was extracted from each tissue (50 mg to 100 mg) and each cell line (5-10 x 10⁶ cells) using the TRIZOL reagent (Invitrogen) in accordance with the attached protocols. cDNA was synthesized using the total RNA and the Superscript First-Strand Synthesis System for RT-PCR (Invitrogen) in accordance with the attached protocols. PCR was performed as follows using primers specific to the obtained genes (represented by SEQ ID NOs: 33 and 34). Specifically, PCR was performed by preparing a reaction solution to bring a total amount thereof to 25 μ l with the addition of reagents and an included buffer (i.e., 0.25 μ l of a sample prepared by reverse transcription reaction, the above primers (2 μ M each), dNTPs (0.2 mM each), 0.65 U of ExTaq polymerase (Takara Shuzo, Co., Ltd.)), the resulting solution was subjected to a cycle of 94°C for 30 seconds, 60°C for 30 seconds, and 72°C for 30 seconds using a Thermal Cycler (BIO RAD), and this cycle was repeated 30 times. The gene-specific primers mentioned above were used to amplify the region between the nucleotide No. 698 and the nucleotide No. 1124 in the nucleotide sequence represented by SEQ ID NO: 1 (the human CAPRIN-1 gene). For comparison, GAPDH-specific primers (represented by SEQ ID NOs: 35 and 36) were used at the same time. As a result of inspection of human CAPRIN-1 gene expression, expression thereof was observed only in the testis in the case of healthy canine tissues, although expression was observed in the pancreatic cancer cells. The results demonstrate that CAPRIN-1 expression was not observed in normal tissues other than those of the testis, while CAPRIN-1 expression was observed in the pancreatic cancer cells.

(5) CAPRIN-1 expression in normal mouse and canine tissues

[0070] Mice (Balb/c, female) and dogs (beagle dogs, female) were exsanguinated under ether anesthesia and ketamine/isoflurane anesthesia. After laparotomy, organs (stomach, liver, eyeball, thymus gland, muscle, bone marrow, uterus, small intestine, esophagus, heart, kidney, salivary gland, large intestine, mammary gland, brain, lung, skin, adrenal gland, ovary, pancreas, spleen, and bladder) were each transferred to a 10 cm dish containing PBS. Each organ was cut open in PBS and then fixed by perfusion overnight with 0.1 M phosphate buffer (pH 7.4) containing 4% paraformaldehyde (PFA). The perfusate was discarded, the tissue surface of each organ was rinsed with PBS, and a PBS solution containing 10% sucrose was then introduced into a 50 ml centrifugal tube. Each tissue was then introduced into each tube, followed by shaking using a rotor at 4°C for 2 hours. Each solution was substituted with a PBS solution containing 20% sucrose and then allowed to stand at 4°C until tissues precipitated. Each solution was substituted with a PBS solution containing 30% sucrose and then allowed to stand at 4°C until tissues precipitated. Each tissue was removed and a necessary portion was excised with a surgical scalpel. Subsequently, the OCT compound (Tissue Tek) was applied and spread over each tissue surface, and the tissues were then placed on Cryomold. Cryomold was placed on dry ice for rapid freezing. Tissues were sliced into pieces of 10 to 20 μ m long using a cryostat (LEICA), the sliced tissue pieces were then air-dried on glass slides for 30 minutes using a hair dryer, and glass slides onto which sliced tissue pieces had been applied were thus prepared. Subsequently, each glass slide was introduced into a staining bottle filled with PBS-T (saline containing 0.05% Tween 20), a procedure involving exchanging PBS-T with fresh PBS-T was performed every 5 minutes, and this procedure was repeated 3 times. Excess water around each specimen was removed using Kimwipes and each section was then encircled using DAKOPEN (DAKO). As blocking solutions, a MOM mouse Ig blocking reagent (VECTASTAIN) was applied onto mouse tissue and a PBS-T solution containing a 10% fetal calf serum was applied onto canine tissue. The resultants were allowed to stand in a moist chamber at room temperature for 1 hour. Subsequently, a solution prepared with the blocking solution to a 10 μ g/ml anti-CAPRIN-1 monoclonal antibody (monoclonal antibody #8) having the heavy chain variable region of SEQ ID NO: 55 and the light chain variable region of SEQ ID NO: 56, which reacts with the cancer cell surfaces prepared in Example 3, was applied onto each slide glass and then allowed to stand within a moist chamber at 4°C overnight. After 3 instances of 10-minutes-washing with PBS-T, a MOM biotin-labeled anti-IgG antibody (VECTASTAIN) diluted 250-fold with the blocking solution was applied onto each glass slide and then allowed to stand within a moist chamber at room temperature for 1 hour. After 3 instances of 10-minutes-washing with PBS-T, an avidin-biotin ABC reagent (VECTASTAIN) was applied and then allowed to stand

within a moist chamber at room temperature for 5 minutes. After 3times of 10-minutes-washing with PBS-T, a DAB staining solution (10 mg of DAB + 10 μ l of 30% H₂O₂, and 50 ml of 0.05M Tris-HCl, pH 7.6) was applied, and the glass slides were then allowed to stand within a moist chamber at room temperature for 30 minutes. Glass slides were rinsed with distilled water and a hematoxylin reagent (DAKO) was then applied. After being allowed to stand at room temperature for 1 minute, the glass slides were rinsed with distilled water. The glass slides were put in 70%, 80%, 90%, 95%, and 100% ethanol solutions in such order for 1 minute each and then allowed to stand in xylene overnight. The glass slides were removed, coverslipped with Glycergel Mounting Medium (DAKO), and then observed. As a result, CAPRIN-1 expression was observed to a slight degree within cells in all of salivary gland, kidney, colon, and stomach tissues, but no CAPRIN-1 expression was observed on cell surfaces. Also, absolutely no CAPRIN-1 expression was observed in tissues from other organs.

[Example 2] Preparation of canine and human CAPRIN-1 proteins

(1) Preparation of recombinant protein

[0071] A recombinant protein was prepared by the following method based on the gene of SEQ ID NO: 5 obtained in Example 1. PCR was performed by preparing a reaction solution to bring a total amount thereof to 50 μ l with the addition of reagents and an included buffer (i.e., 1 μ l of a vector prepared from the phagemid solution obtained in Example 1 and then subjected to sequence analysis, 2 types of primers containing NdeI and KpnI restriction enzyme cleavage sequences (0.4 μ M each; SEQ ID NOs: 37 and 38), 0.2 mM dNTPs, and 1.25 U PrimeSTAR HS polymerase (Takara Shuzo, Co., Ltd.)), the resulting reaction solution was subjected to a cycle of 98°C for 10 seconds and 68°C for 1.5 minutes using a Thermal Cycler (BIO RAD), and this cycle was repeated 30 times. The above 2 types of primers were used to amplify the region encoding the full-length amino acid sequence of SEQ ID NO: 6 (canine CAPRIN-1). After PCR, the amplified DNA was subjected to 1% agarose gel electrophoresis, and a DNA fragment of about 1.4 kbp was then purified from the gel using a QIAquick Gel Extraction Kit (QIAGEN).

[0072] The purified DNA fragment was ligated to a pCR-Blunt cloning vector (Invitrogen). The vector was transformed into *Escherichia coli*, the plasmid was collected, and the amplified gene fragment was confirmed to match the target sequence via sequencing. The plasmid that matched the target sequence was treated with NdeI and KpnI restriction enzymes, the resultant was purified using a QIAquick Gel Extraction Kit, and the target gene sequence was inserted into a pET30b expression vector (Novagen) for *Escherichia coli* treated with NdeI and KpnI restriction enzymes. With the use of the resulting vector, a His tag-fused recombinant protein can be produced. The plasmid was transformed into *Escherichia coli* BL21 (DE3) for expression, and the target protein was induced to express in *Escherichia coli* with the aid of 1 mM IPTG.

[0073] Separately, the recombinant protein of a canine homolog gene was prepared by the following method based on the gene of SEQ ID NO: 7. PCR was performed by preparing a reaction solution to bring a total amount thereof to 50 μ l with the addition of reagents and an included buffer (i.e., 1 μ l of cDNA, the expression of which was confirmed via RT-PCR, selected from among the various tissues and cellular cDNAs prepared in Example 1, 2 types of primers containing NdeI and KpnI restriction enzyme cleavage sequences (0.4 μ M each; SEQ ID NOs: 39 and 40), 0.2 mM dNTPs, and 1.25 U PrimeSTAR HS polymerase (Takara Shuzo, Co., Ltd.)), the resulting reaction solution was subjected to a cycle of 98°C for 10 seconds and 68°C for 2.5 minutes using a Thermal Cycler (BIO RAD), and this cycle was repeated 30 times. The above 2 types of primers were used to amplify the region encoding the full-length amino acid sequence of SEQ ID NO: 8. After PCR, the amplified DNA was subjected to 1% agarose gel electrophoresis, and a DNA fragment of about 2.2 kbp was then purified from the gel using a QIAquick Gel Extraction Kit (QIAGEN).

[0074] The purified DNA fragment was ligated to a pCR-Blunt cloning vector (Invitrogen). The vector was transformed into *Escherichia coli*, the plasmid was collected, and the amplified gene fragment was confirmed to match the target sequence via sequencing. The plasmid that matched the target sequence was treated with NdeI and KpnI restriction enzymes, the resultant was purified using a QIAquick Gel Extraction Kit, and the target gene sequence was inserted into a pET30b expression vector (Novagen) for *Escherichia coli* treated with NdeI and KpnI restriction enzymes. With the use of the resulting vector, a His tag-fused recombinant protein can be produced. The plasmid was transformed into *Escherichia coli* BL21 (DE3) for expression, and the target protein was induced to express in *Escherichia coli* with the aid of 1 mM IPTG.

[0075] Separately, the recombinant protein of a human homolog gene was prepared by the following method based on the gene of SEQ ID NO: 1. PCR was performed by preparing a reaction solution to bring a total amount thereof to 50 μ l with the addition of reagents and an included buffer (i.e., 1 μ l of cDNA, the expression of which was confirmed via RT-PCR, selected from among the various tissue and cellular cDNAs prepared in Example 1, 2 types of primers containing SacI and XhoI restriction enzyme cleavage sequences (0.4 μ M each; SEQ ID NOs: 41 and 42), 0.2 mM dNTPs, and 1.25 U PrimeSTAR HS polymerase (Takara Shuzo, Co., Ltd.)), the resulting reaction solution was subjected to a cycle of 98°C for 10 seconds and 68°C for 2.5 minutes using a Thermal Cycler (BIO RAD), and this cycle was repeated 30

times. The above 2 types of primers were used to amplify the region encoding the full-length amino acid sequence of SEQ ID NO: 2. After PCR, the amplified DNA was subjected to 1% agarose gel electrophoresis, and a DNA fragment of about 2.1 kbp was then purified from the gel using a QIAquick Gel Extraction Kit (QIAGEN).

5 [0076] The purified DNA fragment was ligated to a pCR-Blunt cloning vector (Invitrogen). The vector was transformed into *Escherichia coli*, the plasmid was collected, and the amplified gene fragment was confirmed to match the target sequence via sequencing. The plasmid that matched the target sequence was treated with *SacI* and *XhoI* restriction enzymes, the resultant was purified using a QIAquick Gel Extraction Kit, and the target gene sequence was inserted into a pET30a expression vector (Novagen) for *Escherichia coli* treated with *SacI* and *XhoI* restriction enzymes. With the use of the resulting vector, a His tag-fused recombinant protein can be produced. The plasmid was transformed into 10 *Escherichia coli* BL21 (DE3) for expression, and the target protein was induced to express in *Escherichia coli* with the aid of 1 mM IPTG.

(2) Purification of recombinant protein

15 [0077] The above-obtained recombinant *Escherichia coli* strain expressing SEQ ID NO: 1, 5, or 7 was cultured at 37°C in LB medium containing 30 µg/ml kanamycin until the absorbance at 600 nm reached around 0.7. Thereafter, isopropyl-β-D-1-thiogalactopyranoside was added to a final concentration of 1 mM, and culture was conducted at 37°C for 4 hours. Subsequently, the culture was centrifuged at 4800 rpm for 10 minutes to collect cells. The cell pellet was suspended in phosphate buffered saline and then centrifuged at 4800 rpm for 10 minutes to wash the cells.

20 [0078] The cells were suspended in phosphate buffered saline and then subjected to ultrasonication on ice. The ultrasonicated *Escherichia coli* solution was centrifuged at 6000 rpm for 20 minutes, the resulting supernatant was designated as a soluble fraction, and the resulting precipitate was designated as an insoluble fraction.

25 [0079] The soluble fraction was added to a nickel chelate column (carrier: Chelating Sepharose™ Fast Flow (GE Healthcare), column capacity: 5 ml, 50 mM hydrochloric acid buffer (pH 8.0) as equilibrating buffer) prepared in accordance with a conventional method. The non-adsorbed fraction was washed with 10 column volumes of 50 mM hydrochloric acid buffer (pH 8.0) and 20 mM phosphate buffer (pH 8.0) containing 20 mM imidazole. Immediately thereafter, 6 beds were eluted with 20 mM phosphate buffer (pH 8.0) containing 100 mM imidazole. After the elution of the protein of interest had been confirmed by Coomassie staining, an elution fraction of 20 mM phosphate buffer (pH 8.0) containing 100 mM imidazole was added to a strong anion exchange column (carrier: Q Sepharose™ Fast Flow (GE Healthcare), column 30 volume: 5 ml, and 20 mM phosphate buffer (pH 8.0) as equilibrating buffer). The non-adsorbed fraction was washed with 10 column volumes of 20 mM phosphate buffer (pH 7.0) and 20 mM phosphate buffer (pH 7.0) containing 200 mM sodium chloride. Immediately thereafter, 5 beds were eluted using 20 mM phosphate buffer (pH 7.0) containing 400 mM sodium chloride. Thus, purified fractions of proteins having the amino acid sequences represented by SEQ ID NO: 2, 6, and 8 were obtained, and these purified fractions were hereafter used as materials for administration tests.

35 [0080] Each of the purified preparations obtained by the above method (200 µl each) was dispensed into 1 ml of a reaction buffer (20 mM Tris-HCl, 50 mM NaCl, 2 mM CaCl₂, pH 7.4), and 2 µl of enterokinase (Novagen) was then added. The preparation was allowed to stand at room temperature overnight for reaction, a His tag was cleaved, and purification was then performed in accordance with the protocols attached to the Enterokinase Cleavage Capture Kit (Novagen). Subsequently, 1.2 ml of the purified preparation obtained by the above method was substituted with physiological phosphate buffer (Nissui Pharmaceutical Co., Ltd.) using ultrafiltration NANOSEP 10K OMEGA (PALL). Sterilized 40 filtration was performed using 0.22 µm HT Tuffryn Acrodisc (PALL), and the resultants were used for the following experiments.

[Example 3] Preparation of antibody against CAPRIN-1

(1) Preparation of polyclonal antibody against CAPRIN-1-derived peptide

45 [0081] In order to obtain an antibody binding to CAPRIN-1, a CAPRIN-1-derived peptide (Arg-Asn-Leu-Glu-Lys-Lys-Lys-Gly-Lys-Leu-Asp-Asp-Tyr-Gln; SEQ ID NO: 43) was synthesized. The peptide as an antigen (1 mg) was mixed with the equivalent volume of an incomplete Freund's adjuvant (IFA) solution, and the mixture was subcutaneously administered to a rabbit 4 times every 2 weeks. Thereafter, blood was collected, and an antiserum containing a polyclonal antibody was obtained. Further, the antiserum was purified using a protein G carrier (GE Healthcare Bio-Sciences), and a polyclonal antibody against the CAPRIN-1-derived peptide was then obtained. Subsequently, the reactivity of the resulting polyclonal antibody to CAPRIN-1 on the cancer cell surface was examined using breast cancer cells. Specifically, 55 10⁶ cells of the human breast cancer cell line MDA-MB-231 V were subjected to centrifugation in a 1.5 ml microcentrifugal tube, a PBS solution supplemented with 0.1% fetal bovine serum (FBS) containing the polyclonal antibody was added thereto, and the resultant was then allowed to stand on ice for 1 hour. After washing with PBS, an FITC-labeled goat anti-mouse IgG antibody (Invitrogen) diluted 500-fold with PBS containing 0.1% FBS was added to the solution, and the

solution was then allowed to stand on ice for 1 hour. After washing with PBS, fluorescence intensity was measured using a FACS Calibur (Becton, Dickinson and Company). Separately, a control was prepared in accordance with a procedure similar to the above, except that PBS containing 0.1% FBS was added instead of the polyclonal antibody. As a result, fluorescence intensity in cells treated with the polyclonal antibody was found to be stronger than that in control cells, and the obtained polyclonal antibody was thus found to bind to the breast cancer cell surface.

(2) Preparation of monoclonal antibody against CAPRIN-1 protein

[0082] The antigenic protein (human CAPRIN-1) (100 μ g) represented by SEQ ID NO: 2 prepared in Example 2 was mixed with the equivalent amount of a MPL+TDM adjuvant (Sigma), and the mixture was used as an antigen solution per mouse. The antigen solution was administered intraperitoneally to a 6-week-old Balb/c mouse (Japan SLC Inc.) and further administered 3 times every week. The spleen was removed 3 days after the final immunization, ground in between two sterilized glass slides, washed with PBS (-) (Nissui), and then centrifuged at 1500 rpm for 10 minutes to remove supernatants. This procedure was repeated 3 times to obtain spleen cells. The thus obtained spleen cells were mixed with the mouse myeloma SP2/0 cells (purchased from ATCC) at a ratio of 10:1. The PEG solution prepared by mixing 200 μ l of RPMI1640 medium containing 10% FBS heated to 37°C and 800 μ l of PEG1500 (Boehringer) was added to the cells. The solution was allowed to stand for 5 minutes for cell fusion. Centrifugation was performed at 1700 rpm for 5 minutes to remove supernatants, the cells were suspended in 150 ml of RPMI1640 medium (HAT selective medium) containing 15% FBS supplemented with 2% equivalent of HAT solution (Gibco), and the suspension was then seeded onto fifteen 96-well plates (Nunc) at 100 μ l/well. Cells were cultured for 7 days at 37°C in the presence of 5% CO₂. Thus, hybridomas resulting from fusion of spleen cells with myeloma cells were obtained.

[0083] Hybridomas were selected using, as an indicator, the binding affinity of the antibody produced by the prepared hybridomas for the CAPRIN-1 protein. The CAPRIN-1 protein solution (1 μ g/ml) prepared in Example 2 was added to a 96-well plate at 100 μ l/well, and the resultant was allowed to stand at 4°C for 18 hours. Each well was washed 3 times with PBS-T, a 0.5% bovine serum albumin (BSA) solution (Sigma) was added at 400 μ l/well, and the plate was then allowed to stand at room temperature for 3 hours. The solution was removed and each well was washed 3 times with 400 μ l of PBS-T. Thereafter, each culture supernatant of the hybridomas obtained above was added at 100 μ l/well, and the resultant was then allowed to stand at room temperature for 2 hours. Each well was washed 3 times with PBS-T, an HRP-labeled anti-mouse IgG (H+L) antibody (Invitrogen) diluted 5000-fold with PBS was added thereto at 100 μ l/well, and the resultant was allowed to stand at room temperature for 1 hour. After each well was washed 3 times with PBS-T, a TMB substrate solution (Thermo) was added at 100 μ l per/well, and the resultant was allowed to stand for 15 to 30 minutes, so as to allow the color to develop. Thereafter, 1N sulfuric acid was added at 100 μ l/well to terminate the reaction. The absorbance was measured at 450 nm and 595 nm using a spectrophotometer. As a result, a plurality of hybridomas producing antibodies exhibiting high absorbance values were selected.

[0084] The thus selected hybridomas were added to a 96-well plate at 0.5 hybridomas per well and then cultured. After 1 week, hybridomas forming single colonies in wells were observed. Cells in these wells were further cultured, and hybridomas were selected using, as an indicator, the binding affinity of the antibody produced by the cloned hybridomas for the CAPRIN-1 protein. The CAPRIN-1 protein solution (1 μ g/ml) prepared in Example 2 was added to a 96-well plate at 100 μ l/well, and the resultant was allowed to stand at 4°C for 18 hours. Each well was washed 3 times with PBS-T, a 0.5% BSA solution was added at 400 μ l/well, and the plate was then allowed to stand at room temperature for 3 hours. The solution was removed and each well was washed 3 times with 400 μ l of PBS-T. Thereafter, each culture supernatant of the hybridomas obtained above was added at 100 μ l/well, and the resultant was then allowed to stand at room temperature for 2 hours. Each well was washed 3 times with PBS-T, an HRP-labeled anti-mouse IgG (H+L) antibody (Invitrogen) diluted 5000-fold with PBS was added thereto at 100 μ l/well, and the resultant was allowed to stand at room temperature for 1 hour. After each well was washed 3 times with PBS-T, a TMB substrate solution (Thermo) was added at 100 μ l per/well and then allowed to stand for 15 to 30 minutes, so as to allow the color to develop. Thereafter, 1N sulfuric acid was added at 100 μ l/well to terminate the reaction, and the absorbance was measured at 450 nm and 595 nm using a spectrophotometer. As a result, a plurality of hybridomas producing monoclonal antibodies exhibiting reactivity to the CAPRIN-1 protein were selected, the culture supernatant of hybridomas was purified using a protein G carrier, and 150 monoclonal antibodies binding to the CAPRIN-1 protein were obtained.

[0085] Subsequently, monoclonal antibodies exhibiting reactivity to the surfaces of cancer cells expressing CAPRIN-1 were selected from among these monoclonal antibodies using breast cancer cells. Specifically, 10⁶ cells of the human breast cancer cell line MDA-MB-231 V were subjected to centrifugation in a 1.5 ml microcentrifugal tube, 100 μ l of the culture supernatant of the hybridomas was added thereto, and the resultant was then allowed to stand on ice for 1 hour. After washing with PBS, an FITC-labeled goat anti-mouse IgG antibody (Invitrogen) diluted 500-fold with PBS containing 0.1% FBS was added to the solution, and the solution was then allowed to stand on ice for 1 hour. After washing with PBS, fluorescence intensity was measured using a FACS Calibur (Becton, Dickinson and Company). Separately, a control was prepared in accordance with a procedure similar to the above, except that a medium was added instead of

the antibody. As a result, 10 monoclonal antibodies exhibiting stronger fluorescence intensity than that of the control; i.e., 10 monoclonal antibodies exhibiting reactivity to the surfaces of breast cancer cells (#1 to #10), were selected. The heavy chain variable regions and the light chain variable regions of these monoclonal antibodies are shown in SEQ ID NOs: 44 to 60. The above monoclonal antibody #1 comprises the heavy chain variable region of SEQ ID NO: 44 and the light chain variable region of SEQ ID NO: 45, the monoclonal antibody #2 comprises the heavy chain variable region of SEQ ID NO: 44 and the light chain variable region of SEQ ID NO: 46, the monoclonal antibody #3 comprises the heavy chain variable region of SEQ ID NO: 44 and the light chain variable region of SEQ ID NO: 47, the monoclonal antibody #4 comprises the heavy chain variable region of SEQ ID NO: 44 and the light chain variable region of SEQ ID NO: 48, the monoclonal antibody #5 comprises the heavy chain variable region of SEQ ID NO: 49 and the light chain variable region of SEQ ID NO: 50, the monoclonal antibody #6 comprises the heavy chain variable region of SEQ ID NO: 51 and the light chain variable region of SEQ ID NO: 52, the monoclonal antibody #7 comprises the heavy chain variable region of SEQ ID NO: 53 and the light chain variable region of SEQ ID NO: 54, the monoclonal antibody #8 comprises the heavy chain variable region of SEQ ID NO: 55 and the light chain variable region of SEQ ID NO: 56, the monoclonal antibody #9 comprises the heavy chain variable region of SEQ ID NO: 57 and the light chain variable region of SEQ ID NO: 58, and the monoclonal antibody #10 comprises the heavy chain variable region of SEQ ID NO: 59 and the light chain variable region of SEQ ID NO: 60.

(3) Identification of peptides in CAPRIN-1 protein to which antibodies against CAPRIN-1 reacting with breast cancer cell surface bind

[0086] With the use of monoclonal antibodies #1 to #10 against CAPRIN-1 reacting with the surfaces of breast cancer cells obtained above, partial sequences in the CAPRIN-1 protein recognized by these monoclonal antibodies were identified.

[0087] To 100 μ l of a recombinant CAPRIN-1 protein solution adjusted to a concentration of 1 μ g/ μ l with PBS, first of all, DTT (Fluka) was added to result in a final concentration of 10 mM therein, and a reaction was allowed to proceed at 95°C for 5 minutes, so as to reduce disulfide bonds within the CAPRIN-1 protein. Subsequently, iodoacetamide (final concentration: 20 mM; Wako Pure Chemical Industries, Ltd.) was added, and thiol groups were subjected to alkylation at 37°C for 30 minutes under shaded conditions. The monoclonal antibodies #1 to #10 against CAPRIN-1 (50 μ g each) were added to 40 μ g of the reduced-alkylated CAPRIN-1 protein, the volume of the mixture was adjusted to 1 ml with 20 mM phosphate buffer (pH7.0), and the reaction was allowed to proceed at 4°C overnight with stirring and mixing.

[0088] Subsequently, trypsin (Promega) was added to a final concentration of 0.2 μ g. After the reaction was allowed to proceed at 37°C for 1 hour, 2 hours, 4 hours, and then 12 hours, the resultants were mixed with protein A-glass beads (GE), which had been subjected to blocking with PBS containing 1% BSA (Sigma) and washing with PBS in advance, in 1 mM calcium carbonate and NP-40 buffer (20 mM phosphate buffer (pH 7.4), 5 mM EDTA, 150 mM NaCl, and 1% NP-40), and the reaction was allowed to further proceed for 30 minutes.

[0089] The reaction solutions were each washed with 25 mM ammonium carbonate buffer (pH 8.0), antigen-antibody complexes were then eluted using 100 μ l of 0.1 % formic acid, and the eluates were subjected to LC-MS analysis using Q-TOF Premier (Waters-MicroMass) in accordance with the protocols attached to the instrument.

[0090] As a result, the polypeptide of SEQ ID NO: 61 was identified as a partial sequence of CAPRIN-1, which was recognized by all of the monoclonal antibodies #1 to #10 against CAPRIN-1. Further, the peptide of SEQ ID NO: 62 was identified as a partial sequence in the polypeptide of SEQ ID NO: 61 above, which was recognized by the monoclonal antibodies #1 to #4, #5 to #7, and #9. In addition, the monoclonal antibodies #1 to #4 were found to recognize the peptide of SEQ ID NO: 63, which was a partial peptide sequence thereof.

[Example 4] Diagnosis of pancreatic cancer using CAPRIN-1 polypeptide 1) Diagnosis of canine pancreatic cancer

[0091] As a result of pathological diagnosis using the removed tumor tissue samples, blood samples were collected from afflicted dogs confirmed to have malignant pancreatic ductal carcinoma, and sera were separated. With the use of the canine CAPRIN-1 protein (SEQ ID NO: 8) prepared in Example 2 and the anti-canine IgG antibody, the titer of the serum IgG antibody specifically reacting with the canine CAPRIN-1 protein was measured by an ELISA method.

[0092] The prepared canine CAPRIN-1 protein was immobilized by adding a recombinant protein solution diluted to 5 μ g/ml with phosphate buffered saline to a 96-well immobilizer amino plate (Nunc) at 100 μ l/well and then allowing the plate to stand at 4°C overnight. Blocking was performed by adding a 50 mM sodium bicarbonate buffer solution (pH 8.4) (hereafter, referred to as a "blocking solution") containing 0.5% BSA (bovine serum albumin, Sigma Aldrich Japan) at 100 μ l/well, followed by shaking at room temperature for 1 hour. Serum diluted 1000-fold with the blocking solution was added at 100 μ l/well and the mixture was then subjected to a reaction via shaking at room temperature for 3 hours. The reaction product was washed 3 times with phosphate buffered saline containing 0.05% Tween 20 (Wako Pure Chemical Industries, Ltd.; this solution is referred to as "PBS-T" herein), an HRP-modified canine IgG antibody (Goat anti-Dog

IgG-h+I HRP conjugated: BETHYL Laboratories) diluted 3000-fold with the blocking solution was added at 100 μ l/well, and the mixture was subjected to a reaction via shaking at room temperature for 1 hour. After the reaction product was washed 3 times with PBS-T, HRP substrate TMB (1-Step Turbo TMB (tetramethylbenzidine), PIERCE) was added at 100 μ l/well, and an enzyme-substrate reaction was then conducted at room temperature for 30 minutes. Thereafter, a 0.5 M sulfuric acid solution (Sigma Aldrich Japan) was added at 100 μ l/well to terminate the reaction, and the absorbance at 450 nm was measured using a microplate reader. As controls, a specimen onto which no recombinant protein prepared had been immobilized and a specimen to which the serum of a cancer-bearing dog would not be allowed to react were subjected to the treatment and comparison in the same manner as described above.

[0093] As a result, the titer of the antibody against a canine CAPRIN-1 protein of the sera derived from cancer-carrying dogs was found to be higher than that of the controls.

(2) Diagnosis of canine pancreatic cancer using human CAPRIN-1 protein

[0094] With the use of the human CAPRIN-1 protein (SEQ ID NO: 2) prepared in Example 2, the IgG antibody titer of the canine serum reacting with the human CAPRIN-1 protein was measured in the same manner as described above. When serum samples obtained from healthy dogs were subjected to the same measurement, the absorbance at 450 nm was not substantially observed as described above. The serum samples obtained from pancreatic cancer patient dogs of (1) exhibited a higher titer of the antibody against the human CAPRIN-1 protein than that of the control.

(3) Diagnosis of human pancreatic cancer

[0095] With the use of the human CAPRIN-1 protein (SEQ ID NO: 2) prepared in Example 2 and the anti-human IgG antibody, the IgG antibody titer of the serum samples obtained from healthy individuals reacting with the polypeptide was measured. The human CAPRIN-1 protein was immobilized by adding a recombinant protein solution diluted to 100 μ g/ml with phosphate buffered saline to a 96-well immobilizer amino plate (Nunc) at 100 μ l/well and then allowing the plate to stand at 4°C overnight. Blocking was performed in the following manner. That is, 4 g of Block Ace powder (DS PHARMA BIOMEDICAL Co., Ltd.) was dissolved in 100 ml of purified water, the solution was diluted 4-fold with purified water (hereafter, referred to as a "blocking solution"), the blocking solution was added at 100 μ l/well, and the mixture was subjected to shaking at room temperature for 1 hour. Serum diluted 1000-fold with the blocking solution was added at 100 μ l/well and then subjected to a reaction via shaking at room temperature for 3 hours. After washing the resultant 3 times with phosphate buffered saline containing 0.05% Tween 20 (Wako Pure Chemical Industries, Ltd.; this solution is referred to as "PBS-T" herein), an HRP-modified anti-human IgG antibody (HRP-Goat Anti-Human IgG (H+L) Conjugate: Zymed Laboratories) diluted 10000-fold with the blocking solution was added at 100 μ l/well and then subjected to a reaction via shaking at room temperature for 1 hour. After the reaction product was washed 3 times with PBS-T, HRP substrate TMB (1-Step Turbo TMB (tetramethylbenzidine), PIERCE) was added at 100 μ l/well, and an enzyme-substrate reaction was then performed at room temperature for 30 minutes. Thereafter, a 0.5 M sulfuric acid solution (Sigma Aldrich Japan) was added at 100 μ l/well to terminate the reaction, and the absorbance at 450 nm was then measured using a microplate reader. An ovalbumin antigen adjusted to 50 μ g/ml with phosphate buffered saline was immobilized and then used as a positive control. As a result, the absorbance at 450 nm was found to be high in the case of the ovalbumin antigen, although no absorbance (0) was detected in the case of the human CAPRIN-1 protein.

[0096] Further, the serum samples obtained from patients with pancreatic ductal carcinoma were subjected to measurement of the titer of the serum IgG antibody specifically reacting with the human CAPRIN-1 protein (the amino acid sequence of SEQ ID NO: 2) in the same manner as described above. As a result, the absorbance at 450 nm was found to be lower than the lowest detection limit in the case of healthy subjects, although it was found to be high in the case of patients with pancreatic cancer. With the use of the canine CAPRIN-1 protein (SEQ ID NO: 8) prepared in Example 2 and the anti-human IgG antibody, the titer of the human serum IgG antibody specifically reacting with the canine CAPRIN-1 protein was measured in the same manner as described above. As a result, pancreatic cancer patients were found to exhibit higher titers than healthy individuals.

[0097] Thus, it was demonstrated that human pancreatic cancer could be detected by the method of the present invention.

[Example 5] Diagnosis of pancreatic cancer using antibody against CAPRIN-1

(1) Diagnosis of pancreatic cancer by measuring CAPRIN-1 protein

[0098] With the use of the polyclonal antibody against the CAPRIN-1-derived peptide (SEQ ID NO: 43) obtained in Example 3 (1) in combination with each monoclonal antibody against the CAPRIN-1 protein obtained in Example 3 (2), Sandwich ELISA was carried out in order detect the CAPRIN-1 protein (cancer-bearing individual-derived serum) reacted

positive upon cancer diagnosis using the CAPRIN-1 protein in Example 4 (1)-(3). The polyclonal antibody was used as a primary antibody and each monoclonal antibody was used as a secondary antibody. The amount of the proteins specifically reacting with each of the above antibodies in the sera was measured.

5 [0099] The primary antibody was immobilized by adding a polyclonal antibody solution diluted to 5 $\mu\text{g/ml}$ with phosphate buffered saline to a 96-well immobilizer amino plate (Nunc) at 100 $\mu\text{l/well}$ and shaking the plate at room temperature for 2 hours. Blocking was performed by adding a 50 mM sodium bicarbonate buffer solution (pH 8.4) (hereafter, referred to as a "blocking solution") containing 0.5% BSA (bovine serum albumin, Sigma Aldrich Japan) at 100 $\mu\text{l/well}$, followed by shaking at room temperature for 1 hour. Thereafter, the serum samples obtained from cancer-bearing individuals 10 diluted with a blocking solution were added at 100 $\mu\text{l/well}$ and then subjected to the reaction via shaking at room temperature for 3 hours. The dilution rate at this time was adjusted with 10-fold dilution series (i.e., 10-1000-fold dilutions). The reaction product was washed 3 times with phosphate buffered saline containing 0.05% Tween 20 (Wako Pure Chemical Industries, Ltd.; this solution is referred to as "PBS-T" herein), each monoclonal antibody as a secondary antibody diluted to a concentration of 1 $\mu\text{g/ml}$ with the blocking solution was added at 100 $\mu\text{l/well}$, and the resultant was then subjected to shaking at room temperature for 1 hour for reaction. The reaction product was washed 3 times with 15 PBS-T, an HRP-labeled anti-mouse IgG (H+L) antibody (Invitrogen) as a tertiary antibody diluted 5000-fold with the blocking solution was added at 100 $\mu\text{l/well}$, and the resultant was then allowed to stand at room temperature for 1 hour. After each well was washed 3 times with PBS-T, a TMB substrate solution (Thermo) was added at 100 μl per/well, and the resultant was allowed to stand for 15 to 30 minutes, so as to allow the color to develop. Thereafter, 1N sulfuric acid was added at 100 $\mu\text{l/well}$ to terminate the reaction. The absorbance was measured at 450 nm using a spectrophotometer.

20 [0100] When the monoclonal antibodies #1 to #10 reacting with the surfaces of cancer cells were used as secondary antibodies, as a result, high absorbance values were detected in all the dogs with pancreatic ductal carcinoma, although no absorbance was detected in healthy dogs. When monoclonal antibodies that react with the CAPRIN-1 proteins but do not react with the surfaces of cancer cells were used as secondary antibodies, polypeptide values were detected in all specimens. However, all the absorbance values were lower than the detection limit, which were lower than the results 25 for combinations of antibodies reacting with the surfaces of cancer cells.

[0101] Therefore, cancer can also be diagnosed or examined by this technique that comprises detection of the CAPRIN-1 proteins using antibodies against CAPRIN-1.

30 (2) Diagnosis or examination of cancer by measuring antigenic polypeptide on pancreatic cancer tissue by immunohistochemical staining

[0102] Immunohistochemical staining was performed using an array (BIOMAX) having 101 paraffin-embedded human pancreatic cancer tissue specimens. The array of human pancreatic cancer tissues was treated at 60°C for 3 hours, the resultant was put in a staining bottle filled with xylene, xylene was replaced with fresh xylene every 5 minutes, and this 35 procedure was repeated 3 times. Subsequently, a similar procedure was carried out using ethanol and PBS-T instead of xylene. The array of human pancreatic cancer tissues was put in a staining bottle filled with 10 mM citrate buffer (pH 6.0) containing 0.05% Tween 20, treated at 125°C for 5 minutes, and then allowed to stand at room temperature for 40 minutes or longer. Excess water around each specimen was removed using Kimwipes, each section was encircled with DAKOPEN (DAKO), and an appropriate amount of Peroxidase Block (DAKO) was then added dropwise onto the array. The array was allowed to stand at room temperature for 5 minutes, the array was put in a staining bottle filled with PBS-T, and PBS-T was replaced with fresh PBS-T every 5 minutes. This procedure was performed 3 times. As a blocking 40 solution, a PBS-T solution containing 10% FBS was applied onto the array, and the array was then allowed to stand in a moist chamber at room temperature for 1 hour. Subsequently, the monoclonal antibodies #1 to #10 prepared in Example 3 adjusted to 10 $\mu\text{g/ml}$ with a PBS-T solution containing 5% FBS were applied onto the array, and the array was allowed to stand in a moist chamber at 4°C overnight. After the array was washed with PBS-T for 10 minutes 3 times, an 45 appropriate amount of Peroxidase Labeled Polymer Conjugated (DAKO) was added dropwise onto the array, and the array was allowed to stand in a moist chamber at room temperature for 30 minutes. After the array was washed with PBS-T for 10 minutes 3 times, a DAB color-developing solution (DAKO) was applied onto the array, and the array was then allowed to stand at room temperature for about 10 minutes. After the color-developing solution was discarded, the 50 array was washed with PBS-T for 10 minutes 3 times, rinsed with distilled water, successively put in 70%, 80%, 90%, 95%, and 100% ethanol solutions for 1 minute each, and then allowed to stand in xylene overnight. The glass slides were removed, coverslipped with Glycergel Mounting Medium (DAKO), and then observed. As a result, CAPRIN-1 expression was observed in pancreatic cancer cell membranes and in pancreatic cancer cells in the pancreatic cancer tissue samples with the use of any antibodies. When immunohistochemical staining was carried out with the use of 55 Antibody #8, for example, strong CAPRIN-1 expression was observed in 54 specimens among the total pancreatic cancer tissue specimens (101 specimens) (i.e., 54%).

[0103] Similarly, immunohistochemical staining was carried out using an array (BIOMAX) of paraffin-embedded normal human tissues including normal human pancreatic tissues. Excess water around each specimen was removed using

Kimwipes, each section was encircled with DAKOPEN (DAKO), and an appropriate amount of Peroxidase Block (DAKO) was then added dropwise onto the array. The array was allowed to stand at room temperature for 5 minutes, the array was put in a staining bottle filled with PBS-T, and PBS-T was replaced with fresh PBS-T every 5 minutes. This procedure was performed 3 times. As a blocking solution, a PBS-T solution containing 10% FBS was applied onto the array, and the array was then allowed to stand in a moist chamber at room temperature for 1 hour. Subsequently, the monoclonal antibodies #1 to #10 prepared in Example 3 adjusted to 10 $\mu\text{g/ml}$ in a PBS-T solution containing 5% FBS were applied onto the array, and the array was then allowed to stand in a moist chamber at 4°C overnight. After the array was washed with PBS-T for 10 minutes 3 times, an appropriate amount of Peroxidase Labeled Polymer Conjugated (DAKO) was added dropwise onto the array, and the array was allowed to stand in a moist chamber at room temperature for 30 minutes. After the array was washed with PBS-T for 10 minutes 3 times, a DAB color-developing solution (DAKO) was applied onto the array, and the array was then allowed to stand at room temperature for about 10 minutes. After the color-developing solution was discarded, the array was washed with PBS-T for 10 minutes 3 times, rinsed with distilled water, successively put in 70%, 80%, 90%, 95%, and 100% ethanol solutions for 1 minute each, and then allowed to stand in xylene overnight. The glass slides were removed, coverslipped with Glycergel Mounting Medium (DAKO), and then observed. As a result, none of pancreas-derived normal tissue samples were stained and no CAPRN-1 expression was observed, no matter what antibody was used.

Industrial Applicability

[0104] The present invention is industrially useful for diagnosis or detection of pancreatic cancer.

[0105] All publications, patents, and patent applications cited herein are incorporated herein by reference in their entirety.

Free Text of Sequence Listing

[0106] SEQ ID NOs: 31 to 42: primers

EP 2 741 085 A1

SEQUENCE LISTING

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<150> JP2011-171364

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75	caaatt	tttaa	ttttt	gaatg	attta	atttt	ccctg	tact	atata	aaactg	tottg	aaaac					2474
80	taga	acatat	tctctt	ctca	gaaaa	agtgt	ttttc	caact	gaaa	attatt	tttc	caggtcc					2534
85	taaa	acctgc	taaat	gtttt	tagga	agtac	ttact	gaaac	attttt	gtaa	gacatt	ttttg					2594
90	gaat	gagatt	gaacatt	tat	ataa	atttat	tattc	ctctt	tcatt	ttttt	gaaac	atgcc					2654
95	tatt	atattt	taggg	ccaga	cacc	tttaa	tggcc	ggata	agcc	atagtt	aacatt	taga					2714

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gaaccattta gaagtgatag aactaatgga atttgcaatg ccttttggac ctctattagt 2774
gatataaata tcaagttatt tctgactttt aaacaaaact cccaaattcc taacttattg 2834
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atgtaagctc catgagagca ggtaccttgt ctgtcttctc tgctgtatct attccaacg 3494
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Gly Ala Ala Ala Pro Ala Ser Gln His Pro Ala Thr Gly Thr Gly Ala
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45

Val Gln Thr Glu Ala Met Lys Gln Ile Leu Gly Val Ile Asp Lys Lys
50 55 60

50

Leu Arg Asn Leu Glu Lys Lys Lys Gly Lys Leu Asp Asp Tyr Gln Glu
65 70 75 80

55

Arg Met Asn Lys Gly Glu Arg Leu Asn Gln Asp Gln Leu Asp Ala Val
85 90 95

Ser Lys Tyr Gln Glu Val Thr Asn Asn Leu Glu Phe Ala Lys Glu Leu
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Gln Arg Ser Phe Met Ala Leu Ser Gln Asp Ile Gln Lys Thr Ile Lys

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15	Asp	Asp	Glu	Val	Arg	Thr	Asp	Leu	Lys	Gln	Gly	Leu	Asn	Gly	Val	Pro
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20	Ile	Leu	Ser	Glu	Glu	Glu	Leu	Ser	Leu	Leu	Asp	Glu	Phe	Tyr	Lys	Leu
				180					185					190		
25	Val	Asp	Pro	Glu	Arg	Asp	Met	Ser	Leu	Arg	Leu	Asn	Glu	Gln	Tyr	Glu
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35	Val	Cys	Gly	Thr	Thr	Tyr	Lys	Val	Leu	Lys	Glu	Ile	Val	Glu	Arg	Val
		225				230					235					240
40	Phe	Gln	Ser	Asn	Tyr	Phe	Asp	Ser	Thr	His	Asn	His	Gln	Asn	Gly	Leu
				245						250					255	
45	Cys	Glu	Glu	Glu	Glu	Ala	Ala	Ser	Ala	Pro	Ala	Val	Glu	Asp	Gln	Val
				260					265					270		
50	Pro	Glu	Ala	Glu	Pro	Glu	Pro	Ala	Glu	Glu	Tyr	Thr	Glu	Gln	Ser	Glu
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65	Val	Glu	Val	Val	Asn	Ser	Leu	Gln	Gln	Gln	Pro	Gln	Ala	Ala	Ser	Pro
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70	Ser	Val	Pro	Glu	Pro	His	Ser	Leu	Thr	Pro	Val	Ala	Gln	Ala	Asp	Pro
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75	Leu	Val	Arg	Arg	Gln	Arg	Val	Gln	Asp	Leu	Met	Ala	Gln	Met	Gln	Gly
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Leu Asp Pro Ala Ile Val Ser Ala Gln Pro Met Asn Pro Thr Gln Asn
 385 390 395 400
 5 Met Asp Met Pro Gln Leu Val Cys Pro Pro Val His Ser Glu Ser Arg
 405 410 415
 10 Leu Ala Gln Pro Asn Gln Val Pro Val Gln Pro Glu Ala Thr Gln Val
 420 425 430
 15 Pro Leu Val Ser Ser Thr Ser Glu Gly Tyr Thr Ala Ser Gln Pro Leu
 435 440 445
 Tyr Gln Pro Ser His Ala Thr Glu Gln Arg Pro Gln Lys Glu Pro Ile
 450 455 460
 20 Asp Gln Ile Gln Ala Thr Ile Ser Leu Asn Thr Asp Gln Thr Thr Ala
 465 470 475 480
 25 Ser Ser Ser Leu Pro Ala Ala Ser Gln Pro Gln Val Phe Gln Ala Gly
 485 490 495
 Thr Ser Lys Pro Leu His Ser Ser Gly Ile Asn Val Asn Ala Ala Pro
 500 505 510
 30 Phe Gln Ser Met Gln Thr Val Phe Asn Met Asn Ala Pro Val Pro Pro
 515 520 525
 35 Val Asn Glu Pro Glu Thr Leu Lys Gln Gln Asn Gln Tyr Gln Ala Ser
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 Tyr Asn Gln Ser Phe Ser Ser Gln Pro His Gln Val Glu Gln Thr Glu
 545 550 555 560
 40 Leu Gln Gln Glu Gln Leu Gln Thr Val Val Gly Thr Tyr His Gly Ser
 565 570 575
 45 Pro Asp Gln Ser His Gln Val Thr Gly Asn His Gln Gln Pro Pro Gln
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 50 Gln Asn Thr Gly Phe Pro Arg Ser Asn Gln Pro Tyr Tyr Asn Ser Arg
 595 600 605
 Gly Val Ser Arg Gly Gly Ser Arg Gly Ala Arg Gly Leu Met Asn Gly
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 55 Tyr Arg Gly Pro Ala Asn Gly Phe Arg Gly Gly Tyr Asp Gly Tyr Arg
 625 630 635 640

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Pro Ser Phe Ser Asn Thr Pro Asn Ser Gly Tyr Thr Gln Ser Gln Phe
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5 Ser Ala Pro Arg Asp Tyr Ser Gly Tyr Gln Arg Asp Gly Tyr Gln Gln
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Gly Asn Ile Leu Trp Trp
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Gln Asp Ile Gln Lys Thr Ile Lys Lys Thr Ala Arg Arg Glu Gln Leu
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atg aga gag gaa gcg gaa caa aaa cgt tta aaa act gta ctt gag ctc 153
Met Arg Glu Glu Ala Glu Gln Lys Arg Leu Lys Thr Val Leu Glu Leu
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Gln Tyr Val Leu Asp Lys Leu Gly Asp Asp Glu Val Arg Thr Asp Leu
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40 aag caa ggt ttg aat gga gtg cca ata ttg tct gaa gaa gaa ttg tcg 249
Lys Gln Gly Leu Asn Gly Val Pro Ile Leu Ser Glu Glu Glu Leu Ser
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ttg ttg gat gaa ttc tac aaa tta gca gac cct gaa cgg gac atg agc 297
Leu Leu Asp Glu Phe Tyr Lys Leu Ala Asp Pro Glu Arg Asp Met Ser
70 75 80

45 ttg agg ttg aat gag cag tat gaa cat gct tcc att cac ctg tgg gac 345
Leu Arg Leu Asn Glu Gln Tyr Glu His Ala Ser Ile His Leu Trp Asp
85 90 95 100

50 ttg ctg gaa gga aag gaa aag tct gta tgt gga aca acc tat aaa gca 393
Leu Leu Glu Gly Lys Glu Lys Ser Val Cys Gly Thr Thr Tyr Lys Ala
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cta aag gaa att gtt gag cgt gtt ttc cag tca aat tac ttt gac agc 441
Leu Lys Glu Ile Val Glu Arg Val Phe Gln Ser Asn Tyr Phe Asp Ser
120 125 130

55 act cac aac cac cag aat ggg cta tgt gag gaa gaa gag gca gcc tca 489

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	Ala	Pro	Thr	Val	Glu	Asp	Gln	Val	Ala	Glu	Ala	Glu	Pro	Glu	Pro	Ala	
		150					155					160					
10	gaa	gaa	tac	act	gaa	caa	agt	gaa	ggt	gaa	tca	aca	gag	tat	gta	aat	585
	Glu	Glu	Tyr	Thr	Glu	Gln	Ser	Glu	Val	Glu	Ser	Thr	Glu	Tyr	Val	Asn	
	165					170					175				180		
15	aga	caa	ttt	atg	gca	gaa	aca	cag	ttc	agc	agt	ggt	gaa	aag	gag	cag	633
	Arg	Gln	Phe	Met	Ala	Glu	Thr	Gln	Phe	Ser	Ser	Gly	Glu	Lys	Glu	Gln	
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	Val	Asp	Glu	Trp	Thr	Val	Glu	Thr	Val	Glu	Val	Val	Asn	Ser	Leu	Gln	
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25	cag	caa	cct	cag	gct	gcg	tct	cct	tca	gta	cca	gag	ccc	cac	tct	ttg	729
	Gln	Gln	Pro	Gln	Ala	Ala	Ser	Pro	Ser	Val	Pro	Glu	Pro	His	Ser	Leu	
			215					220					225				
30	act	ccg	gtg	gct	cag	gca	gat	ccc	ctt	gtg	aga	aga	cag	cga	gtc	cag	777
	Thr	Pro	Val	Ala	Gln	Ala	Asp	Pro	Leu	Val	Arg	Arg	Gln	Arg	Val	Gln	
		230					235					240					
35	gac	ctt	atg	gcg	cag	atg	cag	ggg	ccc	tat	aat	ttc	ata	cag	gat	tca	825
	Asp	Leu	Met	Ala	Gln	Met	Gln	Gly	Pro	Tyr	Asn	Phe	Ile	Gln	Asp	Ser	
	245					250					255				260		
40	atg	ctg	gat	ttt	gaa	aac	cag	aca	ctc	gat	cct	gcc	att	gta	tct	gca	873
	Met	Leu	Asp	Phe	Glu	Asn	Gln	Thr	Leu	Asp	Pro	Ala	Ile	Val	Ser	Ala	
					265					270					275		
45	cag	cct	atg	aat	ccg	aca	caa	aac	atg	gac	atg	ccc	cag	ctg	gtt	tgc	921
	Gln	Pro	Met	Asn	Pro	Thr	Gln	Asn	Met	Asp	Met	Pro	Gln	Leu	Val	Cys	
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50	cct	cca	ggt	cat	tct	gaa	tct	aga	ctt	gct	caa	cct	aat	caa	gtt	cct	969
	Pro	Pro	Val	His	Ser	Glu	Ser	Arg	Leu	Ala	Gln	Pro	Asn	Gln	Val	Pro	
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	Val	Gln	Pro	Glu	Ala	Thr	Gln	Val	Pro	Leu	Val	Ser	Ser	Thr	Ser	Glu	
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60	ggg	tat	aca	gca	tct	caa	ccc	ttg	tac	cag	cct	tct	cat	gct	aca	gag	1065
	Gly	Tyr	Thr	Ala	Ser	Gln	Pro	Leu	Tyr	Gln	Pro	Ser	His	Ala	Thr	Glu	
	325					330					335				340		
65	caa	cga	cca	caa	aag	gaa	cca	att	gac	cag	att	cag	gca	aca	atc	tct	1113
	Gln	Arg	Pro	Gln	Lys	Glu	Pro	Ile	Asp	Gln	Ile	Gln	Ala	Thr	Ile	Ser	
					345					350					355		
70	tta	aat	aca	gac	cag	act	aca	gcg	tca	tca	tcc	ctt	ccg	gct	gct	tct	1161
	Leu	Asn	Thr	Asp	Gln	Thr	Thr	Ala	Ser	Ser	Ser	Leu	Pro	Ala	Ala	Ser	
				360					365					370			
75	cag	cct	cag	gta	ttc	cag	gct	ggg	aca	agc	aaa	cca	tta	cat	agc	agt	1209
	Gln	Pro	Gln	Val	Phe	Gln	Ala	Gly	Thr	Ser	Lys	Pro	Leu	His	Ser	Ser	
			375				380						385				
80	gga	atc	aat	gta	aat	gca	gct	cca	ttc	caa	tcc	atg	caa	acg	gtg	ttc	1257
	Gly	Ile	Asn	Val	Asn	Ala	Ala	Pro	Phe	Gln	Ser	Met	Gln	Thr	Val	Phe	

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	390		395		400														
	aat atg aat gcc cca gtt cct cct gtt aat gaa cca gaa act ttg aaa																		1305
5	Asn Met Asn Ala Pro Val Pro Pro Val Asn Glu Pro Glu Thr Leu Lys																		
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	caa caa aat cag tac cag gcc agt tat aac cag agc ttt tct agt cag																		1353
	Gln Gln Asn Gln Tyr Gln Ala Ser Tyr Asn Gln Ser Phe Ser Ser Gln																		
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10	cct cac caa gta gaa caa aca gag gga tgc cgc aaa tga acactcagca																		1402
	Pro His Gln Val Glu Gln Thr Glu Gly Cys Arg Lys																		
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	agtgaattaa tctgattcac aggattatgt ttaaagcga aaaacacact ggccagtgta																		1462
15	ccataatag ttaccagaag agttattatc tatttggttct ccctttcagg aaacttattg																		1522
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	Val Leu Glu Leu Gln Tyr Val Leu Asp Lys Leu Gly Asp Asp Glu Val																		
35			35					40					45						
	Arg Thr Asp Leu Lys Gln Gly Leu Asn Gly Val Pro Ile Leu Ser Glu																		
			50				55					60							
40																			
	Glu Glu Leu Ser Leu Leu Asp Glu Phe Tyr Lys Leu Ala Asp Pro Glu																		
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	Arg Asp Met Ser Leu Arg Leu Asn Glu Gln Tyr Glu His Ala Ser Ile																		
			85						90										95
45																			
	His Leu Trp Asp Leu Leu Glu Gly Lys Glu Lys Ser Val Cys Gly Thr																		
			100					105											110
50																			
	Thr Tyr Lys Ala Leu Lys Glu Ile Val Glu Arg Val Phe Gln Ser Asn																		
			115				120												125
55																			
	Tyr Phe Asp Ser Thr His Asn His Gln Asn Gly Leu Cys Glu Glu Glu																		
			130				135						140						

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Glu Ala Ala Ser Ala Pro Thr Val Glu Asp Gln Val Ala Glu Ala Glu
 145 150 155 160
 5 Pro Glu Pro Ala Glu Glu Tyr Thr Glu Gln Ser Glu Val Glu Ser Thr
 165 170 175
 Glu Tyr Val Asn Arg Gln Phe Met Ala Glu Thr Gln Phe Ser Ser Gly
 10 180 185 190
 Glu Lys Glu Gln Val Asp Glu Trp Thr Val Glu Thr Val Glu Val Val
 15 195 200 205
 Asn Ser Leu Gln Gln Gln Pro Gln Ala Ala Ser Pro Ser Val Pro Glu
 210 215 220
 20 Pro His Ser Leu Thr Pro Val Ala Gln Ala Asp Pro Leu Val Arg Arg
 225 230 235 240
 Gln Arg Val Gln Asp Leu Met Ala Gln Met Gln Gly Pro Tyr Asn Phe
 25 245 250 255
 Ile Gln Asp Ser Met Leu Asp Phe Glu Asn Gln Thr Leu Asp Pro Ala
 260 265 270
 30 Ile Val Ser Ala Gln Pro Met Asn Pro Thr Gln Asn Met Asp Met Pro
 275 280 285
 Gln Leu Val Cys Pro Pro Val His Ser Glu Ser Arg Leu Ala Gln Pro
 35 290 295 300
 Asn Gln Val Pro Val Gln Pro Glu Ala Thr Gln Val Pro Leu Val Ser
 305 310 315 320
 40 Ser Thr Ser Glu Gly Tyr Thr Ala Ser Gln Pro Leu Tyr Gln Pro Ser
 325 330 335
 His Ala Thr Glu Gln Arg Pro Gln Lys Glu Pro Ile Asp Gln Ile Gln
 45 340 345 350
 Ala Thr Ile Ser Leu Asn Thr Asp Gln Thr Thr Ala Ser Ser Ser Leu
 50 355 360 365
 Pro Ala Ala Ser Gln Pro Gln Val Phe Gln Ala Gly Thr Ser Lys Pro
 370 375 380
 55 Leu His Ser Ser Gly Ile Asn Val Asn Ala Ala Pro Phe Gln Ser Met
 385 390 395 400

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Gln Thr Val Phe Asn Met Asn Ala Pro Val Pro Pro Val Asn Glu Pro
405 410 415

5 Glu Thr Leu Lys Gln Gln Asn Gln Tyr Gln Ala Ser Tyr Asn Gln Ser
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Pro Pro Pro Pro Ser Gly Ser Ser Gly Ser Glu Ala Ala Ala Ala Ala
20 25 30

30 ggg gcg gcg ggg gcg gcg ggg gcc ggg gcg gct gcg ccc gcc tcc cag 144
Gly Ala Ala Gly Ala Ala Gly Ala Gly Ala Ala Ala Pro Ala Ser Gln
35 40 45

35 cac ccc gcg acc ggc acc ggc gct gtc cag acc gag gcc atg aag cag 192
His Pro Ala Thr Gly Thr Gly Ala Val Gln Thr Glu Ala Met Lys Gln
50 55 60

40 atc ctc ggg gtg atc gac aag aaa ctc cgg aac ctg gag aag aaa aag 240
Ile Leu Gly Val Ile Asp Lys Lys Leu Arg Asn Leu Glu Lys Lys Lys
65 70 75 80

45 ggc aag ctt gat gat tac cag gaa cga atg aac aaa ggg gaa agg ctt 288
Gly Lys Leu Asp Asp Tyr Gln Glu Arg Met Asn Lys Gly Glu Arg Leu
85 90 95

50 aat caa gat cag ctg gat gcc gta tct aag tac cag gaa gtc aca aat 336
Asn Gln Asp Gln Leu Asp Ala Val Ser Lys Tyr Gln Glu Val Thr Asn
100 105 110

55 aac ttg gag ttt gca aaa gaa tta cag agg agt ttc atg gca tta agt 384
Asn Leu Glu Phe Ala Lys Glu Leu Gln Arg Ser Phe Met Ala Leu Ser
115 120 125

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Gln Asp Ile Gln Lys Thr Ile Lys Lys Thr Ala Arg Arg Glu Gln Leu
130 135 140

65 atg aga gag gaa gcg gaa caa aaa cgt tta aaa act gta ctt gag ctc 480
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145 150 155 160

70 cag tat gtt ttg gac aaa ttg gga gat gat gaa gtg aga act gac ctc 528
Gln Tyr Val Leu Asp Lys Leu Gly Asp Asp Glu Val Arg Thr Asp Leu

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	180	185	190	
10	ttg ttg gat gaa ttc tac aaa tta gca gac cct gaa cgg gac atg agc Leu Leu Asp Glu Phe Tyr Lys Leu Ala Asp Pro Glu Arg Asp Met Ser			624
	195	200	205	
15	ttg agg ttg aat gag cag tat gaa cat gct tcc att cac ctg tgg gac Leu Arg Leu Asn Glu Gln Tyr Glu His Ala Ser Ile His Leu Trp Asp			672
	210	215	220	
20	ttg ctg gaa gga aag gaa aag tct gta tgt gga aca acc tat aaa gca Leu Leu Glu Gly Lys Glu Lys Ser Val Cys Gly Thr Thr Tyr Lys Ala			720
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25	cta aag gaa att gtt gag cgt gtt ttc cag tca aat tac ttt gac agc Leu Lys Glu Ile Val Glu Arg Val Phe Gln Ser Asn Tyr Phe Asp Ser			768
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	260	265	270	
35	gca cct aca gtt gaa gac cag gta gct gaa gct gag cct gag cca gca Ala Pro Thr Val Glu Asp Gln Val Ala Glu Ala Glu Pro Glu Pro Ala			864
	275	280	285	
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50	gta gat gag tgg acg gtc gaa aca gtg gag gtg gtg aat tca ctc cag Val Asp Glu Trp Thr Val Glu Thr Val Glu Val Val Asn Ser Leu Gln			1008
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55	cag caa cct cag gct gcg tct cct tca gta cca gag ccc cac tct ttg Gln Gln Pro Gln Ala Ala Ser Pro Ser Val Pro Glu Pro His Ser Leu			1056
	340	345	350	
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	355	360	365	
65	gac ctt atg gcg cag atg cag ggg ccc tat aat ttc ata cag gat tca Asp Leu Met Ala Gln Met Gln Gly Pro Tyr Asn Phe Ile Gln Asp Ser			1152
	370	375	380	
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	385	390	395	400
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	405	410	415	
80	cct cca gtt cat tct gaa tct aga ctt gct caa cct aat caa gtt cct Pro Pro Val His Ser Glu Ser Arg Leu Ala Gln Pro Asn Gln Val Pro			1296
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Gly Lys Leu Asp Asp Tyr Gln Glu Arg Met Asn Lys Gly Glu Arg Leu
85 90 95

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	Asn	Met	Asn	Ala	Pro	Val	Pro	Pro	Val	Asn	Glu	Pro	Glu	Thr	Leu	Lys	
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	Pro	His	Gln	Val	Glu	Gln	Thr	Asp	Leu	Gln	Gln	Glu	Gln	Leu	Gln	Thr	
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	Gly	Asn	His	Gln	Gln	Pro	Pro	Gln	Gln	Asn	Thr	Gly	Phe	Pro	Arg	Ser	
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5 ggt gct aga ggc tta atg aat gga tac agg ggc cct gcc aat gga ttc 1920
Gly Ala Arg Gly Leu Met Asn Gly Tyr Arg Gly Pro Ala Asn Gly Phe
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55 His Pro Ala Thr Gly Thr Gly Ala Val Gln Thr Glu Ala Met Lys Gln
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Ile Leu Gly Val Ile Asp Lys Lys Leu Arg Asn Leu Glu Lys Lys Lys
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Gly Lys Leu Asp Asp Tyr Gln Glu Arg Met Asn Lys Gly Glu Arg Leu
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Asn Gln Asp Gln Leu Asp Ala Val Ser Lys Tyr Gln Glu Val Thr Asn
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5 Asn Leu Glu Phe Ala Lys Glu Leu Gln Arg Ser Phe Met Ala Leu Ser
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Gln Asp Ile Gln Lys Thr Ile Lys Lys Thr Ala Arg Arg Glu Gln Leu
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Lys Gln Gly Leu Asn Gly Val Pro Ile Leu Ser Glu Glu Glu Leu Ser
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25 Leu Arg Leu Asn Glu Gln Tyr Glu His Ala Ser Ile His Leu Trp Asp
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50 Val Asp Glu Trp Thr Val Glu Thr Val Glu Val Val Asn Ser Leu Gln
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	Gln Pro Pro Gln Gln Asn Thr Gly Phe Pro Arg Ser Asn Gln Pro Tyr	
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	Asp Gly Tyr Gln Gln Asn Phe Lys Arg Gly Ser Gly Gln Ser Gly Pro	
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 45 Ala Pro Arg Asp Tyr Ser Gly Tyr Gln Arg Asp Gly Tyr Gln Gln Asn
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	Val Asn Arg Gln Phe Met Ala Glu Ala Gln Phe Ser Gly Glu Lys Glu	
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	Gln	Asp	Leu	Met	Ala	Gln	Met	Gln	Gly	Pro	Tyr	Asn	Phe	Ile	Gln	Asp	
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	Ser	Leu	Asn	Thr	Asp	Gln	Thr	Thr	Ala	Ser	Ser	Ser	Leu	Pro	Ala	Ala	
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	Ser	Gln	Pro	Gln	Val	Phe	Gln	Ala	Gly	Thr	Ser	Lys	Pro	Leu	His	Ser	
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	Phe	Asn	Met	Asn	Ala	Pro	Val	Pro	Pro	Val	Asn	Glu	Pro	Glu	Thr	Leu	
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	Asn Leu Glu Lys Lys Lys Gly Lys Leu Asp Asp Tyr Gln Glu Arg Met	
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	Asn Lys Gly Glu Arg Leu Asn Gln Asp Gln Leu Asp Ala Val Ser Lys	
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	Tyr Gln Glu Val Thr Asn Asn Leu Glu Phe Ala Lys Glu Leu Gln Arg	
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	225 230 235 240	
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	245 250 255	
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70	tcc	ctt	cct	gct	gct	tct	cag	cct	caa	gtg	ttc	cag	gct	ggg	aca	agt	1666
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	Lys	Pro	Leu	His	Ser	Ser	Gly	Ile	Asn	Val	Asn	Ala	Ala	Pro	Phe	Gln	
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15	Arg	Arg	Gln	Arg	Val	Gln	Asp	Leu	Met	Ala	Gln	Met	Gln	Gly	Pro	Tyr
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	Asp Gln Val Ala Glu Ala Glu Pro Glu Pro Ala Glu Glu Tyr Thr Glu	
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	caa agt gag gtt gaa tca aca gag tat gtc aat agg cag ttc atg gca	1035
	Gln Ser Glu Val Glu Ser Thr Glu Tyr Val Asn Arg Gln Phe Met Ala	
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	Glu Thr Gln Phe Ser Ser Gly Glu Lys Glu Gln Val Asp Glu Trp Thr	
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	Ala	Ser	Pro	Ser	Val	Pro	Glu	Pro	His	Ser	Leu	Thr	Pro	Val	Ala	Gln	
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	Ser	Asp	Pro	Leu	Val	Arg	Arg	Gln	Arg	Val	Gln	Asp	Leu	Met	Ala	Gln	
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	Met	Gln	Gly	Pro	Tyr	Asn	Phe	Ile	Gln	Thr	Leu	Asp	Pro	Ala	Ile	Val	
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	Ser	Ala	Gln	Pro	Met	Asn	Pro	Thr	Gln	Asn	Met	Asp	Met	Pro	Gln	Leu	
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	Thr	Glu	Gln	Arg	Pro	Gln	Lys	Glu	Pro	Met	Asp	Gln	Ile	Gln	Ala	Thr	
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	Val	Phe	Asn	Met	Asn	Ala	Pro	Val	Pro	Pro	Ala	Asn	Glu	Pro	Glu	Thr	
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	Leu	Lys	Gln	Gln	Ser	Gln	Tyr	Gln	Ala	Thr	Tyr	Asn	Gln	Ser	Phe	Ser	
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	Ser	Gln	Pro	His	Gln	Val	Glu	Gln	Thr	Glu	Leu	Gln	Gln	Asp	Gln	Leu	
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	Gln	Thr	Val	Val	Gly	Thr	Tyr	His	Gly	Ser	Gln	Asp	Gln	Pro	His	Gln	
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	Arg Pro Asn Arg Gly Met Pro Gln Met Asn Thr Gln Gln Val Asn			
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 45 35 40 45
 Thr Glu Ala Met Lys Gln Ile Leu Gly Val Ile Asp Lys Lys Leu Arg
 50 50 55 60
 Asn Leu Glu Lys Lys Lys Gly Lys Leu Asp Asp Tyr Gln Glu Arg Met
 55 65 70 75 80
 Asn Lys Gly Glu Arg Leu Asn Gln Asp Gln Leu Asp Ala Val Ser Lys
 85 90 95
 Tyr Gln Glu Val Thr Asn Asn Leu Glu Phe Ala Lys Glu Leu Gln Arg
 100 105 110

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Ser Phe Met Ala Leu Ser Gln Asp Ile Gln Lys Thr Ile Lys Lys Thr
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 5 Ala Arg Arg Glu Gln Leu Met Arg Glu Glu Ala Glu Gln Lys Arg Leu
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 20 Pro Glu Arg Asp Met Ser Leu Arg Leu Asn Glu Gln Tyr Glu His Ala
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 225 230 235 240
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 50 Pro Glu Pro His Ser Leu Thr Pro Val Ala Gln Ser Asp Pro Leu Val
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 55 Arg Arg Gln Arg Val Gln Asp Leu Met Ala Gln Met Gln Gly Pro Tyr
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 Asn Pro Thr Gln Asn Met Asp Met Pro Gln Leu Val Cys Pro Gln Val
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 His Ser Glu Ser Arg Leu Ala Gln Ser Asn Gln Val Pro Val Gln Pro
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Glu Ala Thr Gln Val Pro Leu Val Ser Ser Thr Ser Glu Gly Tyr Thr
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 Ala Ser Gln Pro Leu Tyr Gln Pro Ser His Ala Thr Glu Gln Arg Pro
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Asp Gln Thr Thr Ala Ser Ser Ser Leu Pro Ala Ala Ser Gln Pro Gln
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 Val Phe Gln Ala Gly Thr Ser Lys Pro Leu His Ser Ser Gly Ile Asn
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Ala Pro Val Pro Pro Ala Asn Glu Pro Glu Thr Leu Lys Gln Gln Ser
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Thr Tyr His Gly Ser Gln Asp Gln Pro His Gln Val Pro Gly Asn His
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 610 615 620

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 Tyr Asp Gly Tyr Arg Pro Ser Phe Ser Asn Thr Pro Asn Ser Gly Tyr

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	Ala	Arg	Arg	Glu	Gln	Leu	Met	Arg	Glu	Glu	Ala	Glu	Gln	Lys	Arg	Leu	
		130					135					140					
10	aaa	act	gta	ctt	gag	tta	cag	tat	gta	ttg	gat	aag	ctg	gga	gat	gat	658
	Lys	Thr	Val	Leu	Glu	Leu	Gln	Tyr	Val	Leu	Asp	Lys	Leu	Gly	Asp	Asp	
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15	gat	gtg	aga	aca	gat	ctg	aaa	caa	ggt	ttg	agt	gga	gtg	cca	ata	ttg	706
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	Ser	Glu	Glu	Glu	Leu	Ser	Leu	Leu	Asp	Glu	Phe	Tyr	Lys	Leu	Val	Asp	
				180					185					190			
25	cct	gag	cgt	gac	atg	agt	tta	agg	tta	aat	gag	cag	tat	gaa	cat	gcc	802
	Pro	Glu	Arg	Asp	Met	Ser	Leu	Arg	Leu	Asn	Glu	Gln	Tyr	Glu	His	Ala	
			195					200					205				
30	tca	att	cac	ttg	tgg	gat	ttg	ctg	gaa	ggg	aaa	gaa	aag	cct	gtg	tgt	850
	Ser	Ile	His	Leu	Trp	Asp	Leu	Leu	Glu	Gly	Lys	Glu	Lys	Pro	Val	Cys	
		210					215					220					
35	gga	aca	acc	tat	aaa	gct	cta	aag	gaa	att	gtt	gag	cgt	gtt	ttc	cag	898
	Gly	Thr	Thr	Tyr	Lys	Ala	Leu	Lys	Glu	Ile	Val	Glu	Arg	Val	Phe	Gln	
	225					230					235					240	
40	tca	aac	tac	ttt	gat	agc	act	cac	aat	cat	caa	aat	ggg	ttg	tgt	gag	946
	Ser	Asn	Tyr	Phe	Asp	Ser	Thr	His	Asn	His	Gln	Asn	Gly	Leu	Cys	Glu	
				245					250						255		
45	gag	gaa	gag	gcg	gct	tca	gcg	ccc	aca	gtg	gag	gac	cag	gta	gct	gaa	994
	Glu	Glu	Glu	Ala	Ala	Ser	Ala	Pro	Thr	Val	Glu	Asp	Gln	Val	Ala	Glu	
				260					265					270			
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	Ala	Glu	Pro	Glu	Pro	Ala	Glu	Glu	Tyr	Thr	Glu	Gln	Ser	Glu	Val	Glu	
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	Ser	Thr	Glu	Tyr	Val	Asn	Arg	Gln	Phe	Met	Ala	Glu	Thr	Gln	Phe	Ser	
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65	gtt	gta	aac	tca	ctc	cag	cag	caa	cct	cag	gct	gcg	tcc	cct	tca	gtc	1186
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	Pro	Glu	Pro	His	Ser	Leu	Thr	Pro	Val	Ala	Gln	Ser	Asp	Pro	Leu	Val	
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75	aga	agg	cag	cgt	gta	caa	gat	ctt	atg	gca	caa	atg	caa	ggg	ccc	tat	1282
	Arg	Arg	Gln	Arg	Val	Gln	Asp	Leu	Met	Ala	Gln	Met	Gln	Gly	Pro	Tyr	
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	Met Pro Gln Leu Val Cys Pro Gln Val His Ser Glu Ser Arg Leu Ala																			
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	Gln Ser Asn Gln Val Pro Val Gln Pro Glu Ala Thr Gln Val Pro Leu																			
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 5 Asp Val Arg Thr Asp Leu Lys Gln Gly Leu Ser Gly Val Pro Ile Leu
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 Ser Glu Glu Glu Leu Ser Leu Leu Asp Glu Phe Tyr Lys Leu Val Asp
 10 180 185 190
 Pro Glu Arg Asp Met Ser Leu Arg Leu Asn Glu Gln Tyr Glu His Ala
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 Ser Ile His Leu Trp Asp Leu Leu Glu Gly Lys Glu Lys Pro Val Cys
 20 210 215 220
 Gly Thr Thr Tyr Lys Ala Leu Lys Glu Ile Val Glu Arg Val Phe Gln
 25 225 230 235 240
 Ser Asn Tyr Phe Asp Ser Thr His Asn His Gln Asn Gly Leu Cys Glu
 30 245 250 255
 Glu Glu Glu Ala Ala Ser Ala Pro Thr Val Glu Asp Gln Val Ala Glu
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 Ala Glu Pro Glu Pro Ala Glu Glu Tyr Thr Glu Gln Ser Glu Val Glu
 40 275 280 285
 Ser Thr Glu Tyr Val Asn Arg Gln Phe Met Ala Glu Thr Gln Phe Ser
 45 290 295 300
 Ser Gly Glu Lys Glu Gln Val Asp Glu Trp Thr Val Glu Thr Val Glu
 50 305 310 315 320
 Val Val Asn Ser Leu Gln Gln Gln Pro Gln Ala Ala Ser Pro Ser Val
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 Pro Glu Pro His Ser Leu Thr Pro Val Ala Gln Ser Asp Pro Leu Val
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Met Pro Gln Leu Val Cys Pro Gln Val His Ser Glu Ser Arg Leu Ala
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5 Gln Ser Asn Gln Val Pro Val Gln Pro Glu Ala Thr Gln Val Pro Leu
 420 425 430

10 Val Ser Ser Thr Ser Glu Gly Tyr Thr Ala Ser Gln Pro Leu Tyr Gln
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Pro Ser His Ala Thr Glu Gln Arg Pro Gln Lys Glu Pro Met Asp Gln
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20 Ser Leu Pro Ala Ala Ser Gln Pro Gln Val Phe Gln Ala Gly Thr Ser
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25 Lys Pro Leu His Ser Ser Gly Ile Asn Val Asn Ala Ala Pro Phe Gln
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Ser Met Gln Thr Val Phe Asn Met Asn Ala Pro Val Pro Pro Ala Asn
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30 Glu Pro Glu Thr Leu Lys Gln Gln Ser Gln Tyr Gln Ala Thr Tyr Asn
 530 535 540

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Gln Asp Gln Leu Gln Thr Val Val Gly Thr Tyr His Gly Ser Gln Asp
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40 Gln Pro His Gln Val Pro Gly Asn His Gln Gln Pro Pro Gln Gln Asn
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45 Thr Gly Phe Pro Arg Ser Ser Gln Pro Tyr Tyr Asn Ser Arg Gly Val
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50 Gly Pro Ala Asn Gly Phe Arg Gly Gly Tyr Asp Gly Tyr Arg Pro Ser
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55 Phe Ser Asn Thr Pro Asn Ser Gly Tyr Ser Gln Ser Gln Phe Thr Ala
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Pro Arg Asp Tyr Ser Gly Tyr Gln Arg Asp Gly Tyr Gln Gln Asn Phe
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Ile Leu Trp Trp
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ggc ccg ggc ggc aac gag cag gcc ccg gcg gcg gca gcg gcg gcc ccg 96
Gly Pro Gly Gly Asn Glu Gln Ala Pro Ala Ala Ala Ala Ala Pro
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cag gcg tcg ggc ggc agc atc acc tcg gtt cag acc gag gcc atg aag 144
Gln Ala Ser Gly Gly Ser Ile Thr Ser Val Gln Thr Glu Ala Met Lys
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cag atc ttg gga gtg atc gac aaa aag ctc cgc aac ctc gag aag aaa 192
Gln Ile Leu Gly Val Ile Asp Lys Lys Leu Arg Asn Leu Glu Lys Lys
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Lys Ser Lys Leu Asp Asp Tyr Gln Glu Arg Met Asn Lys Gly Glu Arg
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cta aat caa gat caa ctg gat gca gtg tca aaa tac cag gaa gtg aca 288
Leu Asn Gln Asp Gln Leu Asp Ala Val Ser Lys Tyr Gln Glu Val Thr
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aat aac ctg gaa ttc gct aaa gaa ctg cag agg agc ttt atg gca ctg 336
Asn Asn Leu Glu Phe Ala Lys Glu Leu Gln Arg Ser Phe Met Ala Leu
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agc caa gat atc cag aaa aca ata aaa aag acg gct cgc agg gag cag 384
Ser Gln Asp Ile Gln Lys Thr Ile Lys Lys Thr Ala Arg Arg Glu Gln
115 120 125

ctg atg aga gaa gag gct gag cag aag cgt tta aag act gtg cta gag 432
Leu Met Arg Glu Glu Ala Glu Gln Lys Arg Leu Lys Thr Val Leu Glu
130 135 140

ctg cag ttc att ttg gac aag ttg ggt gac gat gaa gtg cgc agt gac 480
Leu Gln Phe Ile Leu Asp Lys Leu Gly Asp Asp Glu Val Arg Ser Asp
145 150 155 160

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15	gcc ctg aag gag Ala Leu Lys Glu 225	ggt gtt gaa cgt Val Val Glu Arg 230	att ctt caa act Ile Leu Gln Thr 235	agt tac ttt gat Ser Tyr Phe Asp 240	720
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25	gca gaa gaa ttt Ala Glu Glu Phe 275	act gaa cct act Thr Glu Pro Thr 280	gaa gtt gaa tcg Glu Val Glu Ser 285	act gag tat gta Thr Glu Tyr Val 285	864
30	aac aga caa ttc Asn Arg Gln Phe 290	atg gca gag act Met Ala Glu Thr 295	cag ttc agc agt Gln Phe Ser Ser 300	agt gag aag gaa Ser Ser Ser Glu Lys Glu	912
	cag gta gat gag Gln Val Asp Glu 305	tgg aca gtt gaa Trp Thr Val Glu 310	acg gtt gag gtt Thr Val Glu Val 315	gta aat tca ctg Val Val Asn Ser Leu	960
35	cag caa caa aca Gln Gln Gln Thr 325	caa gct aca tct Gln Ala Thr Ser 330	cct cca gtt cct Pro Pro Val Pro 335	gaa cct cat aca Glu Pro His Thr 335	1008
40	ctc act act gtg Leu Thr Thr Val 340	gct caa gca gat Gln Ala Asp 345	cct ctt gtt aga Pro Leu Val Arg 350	aga aga cag aga Arg Arg Gln Arg Val	1056
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50	gca cag ccc atg Ala Gln Pro Met 385	aat cca gca cag Asn Pro Ala Gln 390	aat ttg gac atg Asn Leu Asp Met 395	ccg caa atg gtc Pro Gln Met Val 400	1200
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5	gag caa cgg cca cag aag gaa tcc att gac cag att cag gct tca atg	1392
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15	tcc cag ccg caa gtt ttc caa gct gga tct agc aaa cct ttg cat agc	1488
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	Phe Asn Met Asn Ala Pro Val Pro Pro Val Asn Glu Pro Glu Ala Leu	
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30	aag caa caa aat cag tac cag gcc agt tac aac cag agt ttc tcc aat	1632
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35	cag cca cac caa gta gaa caa tca gat ctt cag caa gaa cag ctc cag	1680
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	Thr Val Val Gly Thr Tyr His Gly Ser Pro Asp Gln Thr His Gln Val	
	565 570 575	
45	gca gga aac cac cag caa cct ccc cag cag aat act gga ttt cca cgc	1776
	Ala Gly Asn His Gln Gln Pro Pro Gln Gln Asn Thr Gly Phe Pro Arg	
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50	aac agt cag cct tat tac aac agt cgg gga gtg tct cgt ggt gga tca	1824
	Asn Ser Gln Pro Tyr Tyr Asn Ser Arg Gly Val Ser Arg Gly Gly Ser	
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55	cgt ggg act cgt gga ttg atg aat ggt tac agg gga cct gca aat gga	1872
	Arg Gly Thr Arg Gly Leu Met Asn Gly Tyr Arg Gly Pro Ala Asn Gly	
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60	ttt aga gga gga tat gat ggc tac cgt cct tca ttt tcc aac act ccg	1920
	Phe Arg Gly Gly Tyr Asp Gly Tyr Arg Pro Ser Phe Ser Asn Thr Pro	
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65	aac agt ggt tac acg cag ccc caa ttt aat gct cct cga gat tat tca	1968
	Asn Ser Gly Tyr Thr Gln Pro Gln Phe Asn Ala Pro Arg Asp Tyr Ser	
	645 650 655	
70	aac tac cag cgg gat gga tat cag cag aac ttc aaa cgt ggt tct gga	2016
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75	caa agt ggg cct cgg gga gct cct cga ggt cgt gga ggg ccc cca aga	2064
	Gln Ser Gly Pro Arg Gly Ala Pro Arg Gly Arg Gly Gly Pro Pro Arg	
	675 680 685	

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 <213> Gallus gallus

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20 Gln Ala Ser Gly Gly Ser Ile Thr Ser Val Gln Thr Glu Ala Met Lys
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Gln Ile Leu Gly Val Ile Asp Lys Lys Leu Arg Asn Leu Glu Lys Lys
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25 Lys Ser Lys Leu Asp Asp Tyr Gln Glu Arg Met Asn Lys Gly Glu Arg
 65 70 75 80

30 Leu Asn Gln Asp Gln Leu Asp Ala Val Ser Lys Tyr Gln Glu Val Thr
 85 90 95

Asn Asn Leu Glu Phe Ala Lys Glu Leu Gln Arg Ser Phe Met Ala Leu
 100 105 110

35 Ser Gln Asp Ile Gln Lys Thr Ile Lys Lys Thr Ala Arg Arg Glu Gln
 115 120 125

40 Leu Met Arg Glu Glu Ala Glu Gln Lys Arg Leu Lys Thr Val Leu Glu
 130 135 140

Leu Gln Phe Ile Leu Asp Lys Leu Gly Asp Asp Glu Val Arg Ser Asp
 145 150 155 160

45 Leu Lys Gln Gly Ser Asn Gly Val Pro Val Leu Thr Glu Glu Glu Leu
 165 170 175

50 Thr Met Leu Asp Glu Phe Tyr Lys Leu Val Tyr Pro Glu Arg Asp Met
 180 185 190

Asn Met Arg Leu Asn Glu Gln Tyr Glu Gln Ala Ser Val His Leu Trp
 195 200 205

55 Asp Leu Leu Glu Gly Lys Glu Lys Pro Val Cys Gly Thr Thr Tyr Lys

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5	Ala 225	Leu	Lys	Glu	Val	Val 230	Glu	Arg	Ile	Leu	Gln 235	Thr	Ser	Tyr	Phe	Asp 240			
		Ser	Thr	His	Asn 245	His	Gln	Asn	Gly	Leu	Cys 250	Glu	Glu	Glu	Glu	Ala 255	Ala		
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		Asn 290	Arg	Gln	Phe	Met	Ala	Glu 295	Thr	Gln	Phe	Ser	Ser 300	Ser	Glu	Lys	Glu		
20	Gln 305	Val	Asp	Glu	Trp	Thr 310	Val	Glu	Thr	Val	Glu 315	Val	Val	Asn	Ser	Leu 320			
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30	Gln	Asp	Leu	Met	Ala	Gln	Met	Gln 360	Gly	Pro	Tyr	Asn	Phe 365	Met	Gln	Asp			
35	Ser 370	Met	Leu	Glu	Phe	Glu	Asn 375	Gln	Thr	Leu	Asp	Pro 380	Ala	Ile	Val	Ser			
40	Ala 385	Gln	Pro	Met	Asn	Pro 390	Ala	Gln	Asn	Leu	Asp 395	Met	Pro	Gln	Met	Val 400			
	Cys	Pro	Pro	Val	His 405	Thr	Glu	Ser	Arg	Leu 410	Ala	Gln	Pro	Asn	Gln 415	Val			
45	Pro	Val	Gln	Pro	Glu 420	Ala	Thr	Gln	Val 425	Pro	Leu	Val	Ser	Ser 430	Thr	Ser			
50	Glu	Gly	Tyr	Thr	Ala	Ser	Gln	Pro 440	Met	Tyr	Gln	Pro	Ser 445	His	Thr	Thr			
	Glu 450	Gln	Arg	Pro	Gln	Lys	Glu 455	Ser	Ile	Asp	Gln	Ile 460	Gln	Ala	Ser	Met			
55	Ser 465	Leu	Asn	Ala	Asp	Gln 470	Thr	Pro	Ser	Ser	Ser 475	Ser	Leu	Pro	Thr	Ala 480			

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Ser Gln Pro Gln Val Phe Gln Ala Gly Ser Ser Lys Pro Leu His Ser
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 5 Ser Gly Ile Asn Val Asn Ala Ala Pro Phe Gln Ser Met Gln Thr Val
 500 505 510
 Phe Asn Met Asn Ala Pro Val Pro Pro Val Asn Glu Pro Glu Ala Leu
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 10 Lys Gln Gln Asn Gln Tyr Gln Ala Ser Tyr Asn Gln Ser Phe Ser Asn
 530 535 540
 15 Gln Pro His Gln Val Glu Gln Ser Asp Leu Gln Gln Glu Gln Leu Gln
 545 550 555 560
 Thr Val Val Gly Thr Tyr His Gly Ser Pro Asp Gln Thr His Gln Val
 565 570 575
 20 Ala Gly Asn His Gln Gln Pro Pro Gln Gln Asn Thr Gly Phe Pro Arg
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 25 Asn Ser Gln Pro Tyr Tyr Asn Ser Arg Gly Val Ser Arg Gly Gly Ser
 595 600 605
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 35 Asn Ser Gly Tyr Thr Gln Pro Gln Phe Asn Ala Pro Arg Asp Tyr Ser
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35 Pro Gly Ala Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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40 Thr Asp Tyr Asn Met Asp Trp Val Lys Gln Ser His Gly Lys Ser Leu
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45 Glu Trp Ile Gly Asp Ile Asn Pro Asn Tyr Asp Ser Thr Ser Tyr Asn
65 70 75 80

50 Gln Lys Phe Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser
85 90 95

55 Thr Ala Tyr Met Glu Leu Arg Ser Leu Thr Ser Glu Asp Thr Ala Val
100 105 110

60 Tyr Tyr Cys Ala Arg Ser Arg Ser Tyr Asp Tyr Glu Gly Phe Ala Tyr
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65 Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ala Ala Lys Thr Thr Pro
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70 Pro Ser Val Tyr
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Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu Leu Trp Ser Val
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Asn Gln Lys Asn Tyr Leu Ser Trp Tyr Gln Gln Lys Gln Arg Gln Pro
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Pro Lys Leu Leu Ile Tyr Gly Ala Ser Ile Arg Glu Ser Trp Val Pro
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Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile
 85 90 95

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Ser Asn Val His Ala Glu Asp Leu Ala Val Tyr Tyr Cys Gln His Asn
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Ser Asn Asn Arg
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 35 40 45

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Gly Lys Ser Pro Gln Leu Leu Val Tyr Asn Ala Lys Thr Leu Ala Asp
 50 55 60

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Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Gln Tyr Ser
 65 70 75 80

5 Leu Lys Ile Asn Ser Leu Gln Pro Glu Asp Phe Gly Ser Tyr Tyr Cys
 85 90 95

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15 Ile Lys Gln Ser Asp
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 35 40 45

Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Ser Val
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55 Pro Pro Lys Leu Leu Val Tyr Pro Ala Leu Leu Ile Tyr Glu Ala Ser
 35 40 45

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Ile Thr Lys Ser Cys Val Pro Asp Arg Phe Thr Arg Ser Gly Ser Gly
50 55 60

5 Thr Asn Phe Thr Leu Thr Ile Asn Phe Val His Ala Asp Asp Leu Ile
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Val Gln Val Pro Arg Arg Arg Ser Asn
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30 Asn Pro Tyr Asn Gly Gly Thr Ser Tyr Asn Gln Lys Phe Lys Gly Lys
35 40 45

Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr Met Glu Leu
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35 Leu Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Trp
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Val Ser Ser Lys
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 5 Tyr Leu Ala Ser Asn Arg Asp Thr Gly Leu Pro Asp Arg Phe Pro Gly
 35 40 45
 Arg Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile Thr Asn Val Gln Ser
 50 55 60
 10 Glu Asp Leu Glu Asp Tyr Phe Cys Leu Gln His Cys Asn Tyr Pro Asn
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 35 Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr Gln Lys Phe Lys Gly Lys
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 40 Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala Tyr Met Gln Leu
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 Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Gln Asp Tyr Ser Leu
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 Thr Ile Ser Ser Leu Glu Tyr Glu Asp Met Gly Ile Tyr Tyr Cys Leu
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Asn Leu Glu Ser Gly Ile Pro Ala Arg Phe Ser Gly Ser Gly Ser Arg
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Gly Lys Ala Leu Glu Trp Leu Gly Phe Ile Arg Asn Lys Ala Asn Gly
 35 40 45

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Tyr Thr Thr Glu Tyr Ser Ala Ser Val Lys Gly Arg Phe Thr Ile Ser
 50 55 60

Arg Asp Asn Ser Gln Ser Ile Leu Tyr Leu Gln Met Asn Thr Leu Arg
 65 70 75 80

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Ala Glu Asp Ser Ala Thr Tyr Tyr Cys Ala Arg Ala Pro Leu Leu Tyr
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25 Asn Leu Glu Ser Gly Val Pro Ala Arg Phe Ser Gly Ser Gly Ser Gly
 50 55 60

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 35 40 45

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5 Thr Val Ser Ser Lys
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 35 40 45

30 Gly Val Pro Ala Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr
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Ser Glu Val Glu Ser Thr Glu Tyr Val Asn Arg
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Claims

1. A method for detecting pancreatic cancer, comprising measuring the presence or an amount of a polypeptide having a reactivity of specifically binding to an antibody against a CAPRIN-1 protein via an antigen-antibody reaction, or the presence or an amount of a nucleic acid encoding the polypeptide, in a sample separated from a subject.
2. The method according to claim 1, wherein the polypeptide to be measured is a CAPRIN-1 protein consisting of an amino acid sequence represented by any of the even-numbered SEQ ID NOs: 2 to 30, or a polypeptide consisting of an amino acid sequence having 85-90% or higher sequence identity with the CAPRIN-1 protein.
3. The method according to claim 1 or 2, wherein the subject is a human or a dog.
4. The method according to claim 3, wherein the subject is a dog and the polypeptide to be measured comprises an amino acid sequence represented by any of the even-numbered SEQ ID NOs: 2 to 30.
5. The method according to claim 4, wherein the subject is a dog and the polypeptide to be measured comprises the amino acid sequence represented by SEQ ID NO: 6, 8, 10, 12, or 14.
6. The method according to claim 3, wherein the subject is a human and the polypeptide to be measured comprises the amino acid sequence represented by SEQ ID NO: 2 or 4.
7. The method according to any one of claims 1 to 6, wherein the presence or an amount of the polypeptide is determined by immunologically measuring an antibody induced against the polypeptide to be measured in the body of a subject, which antibody may be contained in the sample.
8. The method according to any one of claims 1 to 6, wherein the presence or an amount of a nucleic acid encoding the polypeptide is determined by measuring a nucleic acid encoding the polypeptide contained in the sample.
9. The method according to claim 8, wherein the presence or an amount of the nucleic acid in the sample is measured using a polynucleotide specifically hybridizing to a partial sequence comprising at least 15 to 19 nucleotides or at least 20 to 30 nucleotides in the nucleotide sequence of the nucleic acid or a sequence complementary thereto.
10. The method according to claim 9, wherein the subject is a dog and the polynucleotide specifically hybridizes to a partial sequence comprising at least 15 to 19 nucleotides or at least 20 to 30 nucleotides in the nucleotide sequence represented by SEQ ID NO: 5, 7, 9, 11, or 13 or a sequence complementary thereto.
11. The method according to claim 9, wherein the subject is a human and the polynucleotide specifically hybridizes to

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a partial sequence comprising at least 15 to 19 nucleotides or at least 20 to 30 nucleotides in the nucleotide sequence represented by SEQ ID NO: 1 or 3 or a sequence complementary thereto.

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12. The method according to any one of claims 1 to 6, wherein the presence or an amount of the polypeptide is determined by measuring the polypeptide contained in the sample.
13. The method according to claim 12, wherein the measurement is an immunological assay.
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14. The method according to any one of claims 1 to 14, wherein the sample is blood, serum, blood plasma, ascites fluid, pleural effusion, tissues, or cells.
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15. A reagent or kit for detecting pancreatic cancer, comprising one or more polypeptides having a reactivity of binding via an antigen-antibody reaction to an antibody induced against a CAPRIN-1 protein in the body of a subject.
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16. A reagent or kit for detecting pancreatic cancer, comprising one or more antibodies that undergo an antigen-antibody reaction with a polypeptide having a reactivity of binding, via an antigen-antibody reaction, to an antibody against a CAPRIN-1 protein and produced in the body of a subject or to an antigen-binding fragment of the antibody.
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17. The reagent or kit according to claim 15 or 16, wherein the CAPRIN-1 protein consists of an amino acid sequence represented by any of the even-numbered SEQ ID NOs: 2 to 30.
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18. The reagent or kit according to claim 16 or 17, wherein the antibody or the antigen-binding fragment thereof that undergoes an antigen-antibody reaction with the polypeptide is an antibody or an antigen-binding fragment thereof that binds to the surface of a pancreatic cancer cell.
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19. The reagent or kit according to any one of claims 16 to 18, wherein the antibody or the antigen-binding fragment thereof that undergoes an antigen-antibody reaction with the polypeptide comprises an antibody or a fragment thereof having an immunological reactivity with a polypeptide consisting of an amino acid sequence comprising at least 7 to 12 continuous amino acid residues within the region of amino acid residue Nos. 50 to 98 or amino acid residue Nos. 233 to 344 of the amino acid sequence represented by any of even-numbered SEQ IDS NO: 2 to 30 except for SEQ ID NOs: 6 and 18.
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20. The reagent or kit according to any one of claims 16 to 19, wherein the antibody or an antigen-binding fragment thereof that undergoes an antigen-antibody reaction with the polypeptide is one or more antibodies or antigen-binding fragments thereof selected from the group consisting of: an antibody binding to a polypeptide comprising the amino acid sequence represented by SEQ ID NO: 43 or an antigen-binding fragment of the antibody; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 44 and 45 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 44 and 46 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 44 and 47 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 44 and 48 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 49 and 50 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 51 and 52 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 53 and 54 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 55 and 56 or an antigen-binding fragment thereof; a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 57 and 58 or an antigen-binding fragment thereof; and a monoclonal antibody comprising the amino acid sequences represented by SEQ ID NOs: 59 and 60 or an antigen-binding fragment thereof.
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21. A reagent or kit for detecting pancreatic cancer, comprising one or more polynucleotides that specifically hybridize to a partial sequence comprising at least 15 to 19 nucleotides or at least 20 to 30 nucleotides in the nucleotide sequence represented by any of the odd-numbered SEQ ID NOs: 1 to 29 and encoding a CAPRIN-1 protein or in a sequence complementary to the nucleotide sequence.
22. A method for detecting pancreatic cancer, comprising measuring the presence or an amount of a CAPRIN-1 protein, an antibody against CAPRIN-1 protein, or a nucleic acid encoding the CAPRIN-1 protein, in a sample from a subject using at least one reagent or kit according to any one of claims 15 to 21.

23. The method for detecting pancreatic cancer according to any one of claims 1 to 14, comprising measuring the presence or an amount of a CAPRIN-1 protein, an antibody against CAPRIN-1 protein, or a nucleic acid encoding the CAPRIN-1 protein, in a sample from a subject using at least one reagent or kit according to any one of claims 15 to 21.

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INTERNATIONAL SEARCH REPORT

International application No.

PCT/JP2012/069824

A. CLASSIFICATION OF SUBJECT MATTER <i>G01N33/574(2006.01) i, C12Q1/68(2006.01) i, G01N33/53(2006.01) i, C07K14/47(2006.01) n, C07K16/18(2006.01) n, C12N15/02(2006.01) n, C12P21/08(2006.01) n</i> According to International Patent Classification (IPC) or to both national classification and IPC		
B. FIELDS SEARCHED Minimum documentation searched (classification system followed by classification symbols) <i>G01N33/574, C12Q1/68, G01N33/53, C07K14/47, C07K16/18, C12N15/02, C12P21/08</i> Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched Jitsuyo Shinan Koho 1922-1996 Jitsuyo Shinan Toroku Koho 1996-2012 Kokai Jitsuyo Shinan Koho 1971-2012 Toroku Jitsuyo Shinan Koho 1994-2012 Electronic data base consulted during the international search (name of data base and, where practicable, search terms used) MEDLINE (STN), JSTPlus/JMEDPlus/JST7580 (JDreamII)		
C. DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X/A	WO 2010/016527 A1 (Toray Industries, Inc.), 11 February 2010 (11.02.2010), claims 1 to 22 & EP 2325648 A1	15-21/4, 5, 10
A	US 2008/0107668 A1 (Immunotope), 08 May 2008 (08.05.2008), & EP 2061503 A	4, 5, 10, 15-21
<input type="checkbox"/> Further documents are listed in the continuation of Box C. <input type="checkbox"/> See patent family annex.		
* Special categories of cited documents: "A" document defining the general state of the art which is not considered to be of particular relevance "E" earlier application or patent but published on or after the international filing date "L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified) "O" document referring to an oral disclosure, use, exhibition or other means "P" document published prior to the international filing date but later than the priority date claimed "I" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention "X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art "&" document member of the same patent family		
Date of the actual completion of the international search 22 October, 2012 (22.10.12)		Date of mailing of the international search report 06 November, 2012 (06.11.12)
Name and mailing address of the ISA/ Japanese Patent Office		Authorized officer
Facsimile No.		Telephone No.

Form PCT/ISA/210 (second sheet) (July 2009)

INTERNATIONAL SEARCH REPORT

International application No.

PCT/JP2012/069824

Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. Claims Nos.: 1-3, 6-9, 11-14, 22, 23
because they relate to subject matter not required to be searched by this Authority, namely:
(See extra sheet)
2. Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying additional fees, this Authority did not invite payment of additional fees.
3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- The additional search fees were accompanied by the applicant's protest and, where applicable, the payment of a protest fee.
- The additional search fees were accompanied by the applicant's protest but the applicable protest fee was not paid within the time limit specified in the invitation.
- No protest accompanied the payment of additional search fees.

Form PCT/ISA/210 (continuation of first sheet (2)) (July 2009)

INTERNATIONAL SEARCH REPORT

International application No.
PCT/JP2012/069824

Continuation of Box No.II-1 of continuation of first sheet (2)

Claims 1-3, 6-9, 11-14, 22 and 23 include "a method for detecting pancreatic cancer" which is a method for diagnosing the human body. Thus, the inventions of these claims relate to a subject matter which this international searching authority is not required, under the provisions of PCT Article 17(2)(a)(i) and PCT Rule 39.1, to search.

REFERENCES CITED IN THE DESCRIPTION

This list of references cited by the applicant is for the reader's convenience only. It does not form part of the European patent document. Even though great care has been taken in compiling the references, errors or omissions cannot be excluded and the EPO disclaims all liability in this regard.

Patent documents cited in the description

- WO 2010016526 A [0006]
- WO 2010016527 A [0006]

Non-patent literature cited in the description

- **KARLIN ; ALTSCHUL.** *Proc. Natl. Acad. Sci. U.S.A.*, 1993, vol. 87, 2264-2268 [0025]
- **ALTSCHUL et al.** *Nucleic Acids Res.*, 1997, vol. 25, 3389-3402 [0025]
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- Molecular Cloning. **SAMBROOK et al.** Current Protocols in Molecular Biology. Cold Spring Harbor Laboratory Press, 1989 [0028]
- Short Protocols in Molecular Biology. **AUSUBEL et al.** A Compendium of Methods from Current Protocols in Molecular Biology. John Wiley & Sons, 1995 [0028]

专利名称(译)	检测胰腺癌的方法		
公开(公告)号	EP2741085A4	公开(公告)日	2015-03-04
申请号	EP2012819473	申请日	2012-08-03
[标]申请(专利权)人(译)	东丽株式会社		
申请(专利权)人(译)	TORAY INDUSTRIES , INC.		
当前申请(专利权)人(译)	TORAY INDUSTRIES , INC.		
[标]发明人	IDO TAKAYOSHI OKANO FUMIYOSHI		
发明人	IDO, TAKAYOSHI OKANO, FUMIYOSHI		
IPC分类号	G01N33/574 C12Q1/68 G01N33/53 C07K14/47 C07K16/18 C12N15/02 C12P21/08		
CPC分类号	C07K16/18 C07K14/47 C07K14/4738 C07K16/30 C07K16/303 C07K2317/34 C07K2317/56 C12Q1/6886 C12Q2600/158 G01N33/57438 G01N2333/705		
优先权	2011171364 2011-08-04 JP		
其他公开文献	EP2741085A1 EP2741085B1		
外部链接	Espacenet		

摘要(译)

本发明涉及使用新型肿瘤标志物检测胰腺癌的方法。具体地，本发明提供了一种用于检测胰腺癌的方法，包括测量通过抗原 - 抗体反应与从受试者分离的样品中的抗CAPRIN-1蛋白的抗体具有结合反应性的多肽的存在或量，并且用于检测胰腺癌的试剂或试剂盒，其包含CAPRIN-1蛋白或其片段，针对其的抗体或编码其的多核苷酸。