

(19) DANMARK

(10) DK/EP 2441775 T3



(12)

Oversættelse af
europæisk patent

Patent- og
Varemærkestyrelsen

(51) Int.Cl.: **C 07 K 14/705 (2006.01)** **G 01 N 33/574 (2006.01)** **G 01 N 33/68 (2006.01)**

(45) Oversættelsen bekendtgjort den: **2016-10-03**

(80) Dato for Den Europæiske Patentmyndigheds
bekendtgørelse om meddelelse af patentet: **2016-08-17**

(86) Europæisk ansøgning nr.: **12150462.5**

(86) Europæisk indleveringsdag: **2008-02-26**

(87) Den europæiske ansøgnings publiceringsdag: **2012-04-18**

(30) Prioritet: **2007-02-26 US 903510 P** **2007-02-26 US 903509 P**
2008-02-25 WO PCT/GB2008/050124

(62) Stamansøgningsnr: **08709647.5**

(84) Designerede stater: **AT BE BG CH CY CZ DE DK EE ES FI FR GB GR HR HU IE IS IT LI LT LU LV MC**
MT NL NO PL PT RO SE SI SK TR

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(54) Benævnelse: **LYMFOCYTANTIGEN 75 (Ly75)**

(56) Fremdragne publikationer:
WO-A1-2004/053138
WO-A2-2005/016962

DK/EP 2441775 T3

DESCRIPTION

INTRODUCTION

[0001] The present invention relates to the identification of membrane protein associated with pancreatic cancer which has utility as a marker for pancreatic cancer metastases and which also forms a biological target against which therapeutic antibodies (or other affinity reagents) or other pharmaceutical agents can be made.

BACKGROUND OF THE INVENTION

Pancreatic Cancer

[0002] Pancreatic cancer is a very difficult cancer to detect and the prognosis for patients is usually very poor. The number of new cases and deaths per year is almost equal. Global incidence of pancreatic cancer is approximately 230,000 cases (about 2% of all cancer cases), with about 225,000 deaths (3.4% of cancer deaths) per year. It is much more prevalent in the developed world. In the USA, there are about 34,000 new cases per year, with about 32,000 deaths. It has been estimated that approximately US\$1.5 billion is spent in the USA each year on treating pancreatic cancer.

Pancreatic Cancer Diagnosis

[0003] Pancreatic cancer is very difficult to detect and very few pancreatic cancers are found early. Patients usually have no symptoms until the cancer has spread to other organs. There are currently no blood tests or easily available screening tests that can accurately detect early cancers of the pancreas. An endoscopic ultrasound followed by a biopsy is the best way to diagnose pancreatic cancer. Other detection methods include CT, CT-guided needle biopsy, PET, ultrasonography and MRI. Blood levels of CA 19-9 and carcinoembryonic antigen (CEA) may be elevated but by the time blood levels are high enough to be detected, the cancer is no longer in its early stages.

Pancreatic Cancer Staging

[0004] Pancreatic cancer has four stages, stage I to stage IV according to the American Joint Committee on Cancer (AJCC) TNM system. Pancreatic cancer is also divided into resectable, locally advanced (unresectable) and metastatic cancer. For patients with advanced cancers, the overall survival rate is <1% at 5 years with most patients dying within 1 year.

Pancreatic Cancer Treatment

[0005] Surgery is the only method of curing pancreatic cancer. About 10% of pancreatic cancers are contained entirely within the pancreas at the time of diagnosis and attempts to remove the entire cancer by surgery may be successful in some of these patients. The 5-year survival for those undergoing surgery with the intent of completely removing the cancer is about 20%. Potentially curative surgery, usually by pancreaticoduodenectomy (Whipple procedure), is used when it may be possible to remove all of the cancer. Palliative surgery may be performed if the tumour is too widespread to be completely removed. Removing only part of the cancer does not allow patients to live longer. Pancreatic cancer surgery is difficult to perform with a high likelihood of complications.

[0006] External beam radiation therapy combined with chemotherapy can be given before or after surgery and can also be given to patients whose tumours are too widespread to be removed by surgery. The main chemotherapeutic agents which are used are gemcitabine and 5-fluorouracil. Targeted therapy using drugs such as erlotinib and cetuximab may be of benefit to patients with advanced pancreatic cancer.

Therapeutic Challenges

[0007] The major challenges in treatment of pancreatic cancer is to improve early detection rates, to find new non-invasive markers that can be used to follow disease progression and identify relapse, and to find improved and less toxic therapies, especially for more advanced disease where 5 year survival is still poor. There is a great need to identify targets which are more specific to the cancer cells, e.g. ones which are expressed on the surface of the tumour cells so that they can be attacked by promising new approaches like immunotherapeutics and targeted toxins.

[0008] WO2004/053138 describes DEC 205 splice variants associated with Hodgkin's disease. These splice variants have been identified on Reed-Sternberg cells and thus suggested to have application in therapy and investigation of this disease.

[0009] WO2005/016962 describes compositions containing novel proteins and methods of using those compositions for the diagnosis and treatment of immune related diseases.

SUMMARY OF THE INVENTION

[0010] The present invention provides methods and compositions for diagnosis and therapy of pancreatic cancer, also described in the present disclosure are methods for colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer patients' stratification, for monitoring the effectiveness of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer treatment, and for drug development for treatment of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.

[0011] We have used mass spectrometry to identify peptides generated by gel electrophoresis or tagging with iTRAQ reagents and tryptic digest of membrane proteins extracted from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer tissue samples. Peptide sequences were compared to existing protein and cDNA databases and the corresponding gene sequences identified. The protein of the invention has not been previously reported to originate from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer cell membranes and represents a protein of new diagnostic and therapeutic value.

[0012] According to one aspect of the present invention there is provided an antibody which is capable of specific binding to Lymphocyte antigen 75 for use in treatment of pancreatic cancer wherein said antibody contains or is conjugated to a therapeutic moiety.

[0013] In one embodiment, the antibody for use is an isolated monoclonal antibody, or an antigen-binding portion thereof, or an antibody fragment.

[0014] In a further embodiment the isolated monoclonal antibody for use is:

1. (A) a full-length antibody of an IgG1, IgG2, IgG3, or IgG4 isotype;
2. (B) selected from the group consisting of a whole antibody, an antibody fragment, a chimeric antibody, a human antibody, a humanised antibody, a single chain antibody, an immunoconjugate, a defucosylated antibody, and a bispecific antibody;
3. (C) the fragment a domain antibody;
4. (D) a monoclonal antibody capable of specific binding to Lymphocyte antigen 75 and which has cytotoxicity against Lymphocyte antigen 75 expressing cells in the presence of a human complement; or
5. (E) a monoclonal antibody capable of specific binding to Lymphocyte antigen 75 and which has cytotoxicity against Lymphocyte antigen 75 expressing cells in the presence of human immune effector cells.

[0015] In a further embodiment the therapeutic moiety is a cytotoxic moiety or a radioactive isotope or wherein the affinity reagent is an antibody drug conjugate.

[0016] In a further embodiment the antibody is a fully human or humanised antibody that is capable of inducing antibody-directed cell-mediated cytotoxicity (ADCC)

[0017] In a further aspect there is provided an antibody drug conjugate which is capable of specific binding to Lymphocyte antigen 75 and which contains or is conjugated to a therapeutic moiety, wherein the antibody is an isolated monoclonal antibody, or an antigen-binding portion thereof, or an antibody fragment.

[0018] In one embodiment the therapeutic moiety is a cytotoxic moiety or a radioactive isotope.

[0019] In a further embodiment the antibody is:

1. (A) a full-length antibody of an IgG1, IgG2, IgG3, or IgG4 isotype;
2. (B) selected from the group consisting of a whole antibody, an antibody fragment, a chimeric antibody, a human antibody, a humanised antibody, a single chain antibody, an immunoconjugate, a defucosylated antibody, and a bispecific antibody
3. (C) the antibody fragment is a domain antibody;
4. (D) a monoclonal antibody which has cytotoxicity against Lymphocyte antigen 75 expressing cells in the presence of a human complement; or
5. (E) a monoclonal antibody which has cytotoxicity against Lymphocyte antigen 75 expressing cells in the presence of human immune effector cells.

[0020] In a further embodiment the antibody is a fully human or humanised antibody that is capable of inducing antibody-directed cell-mediated cytotoxicity (ADCC).

[0021] The disclosure provides a method of detecting, diagnosing and/or screening for or monitoring the progression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or of monitoring the effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy in a subject which comprises detecting the presence or level of OGTA076, or one or more fragments thereof, or the presence or level of nucleic acid encoding OGTA076 or the presence or level of the activity of OGTA076 or which comprises detecting a change in the level thereof in said subject.

[0022] Also described is a method of detecting, diagnosing and/or screening for colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in a candidate subject which comprises detecting the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 in said candidate subject, in which either (a) the presence of an elevated level of OGTA076 or said one or more fragments thereof or an elevated level of nucleic acid encoding OGTA076 or the presence of an elevated level of OGTA076 activity in the candidate subject as compared with the level in a healthy subject or (b) the presence of a detectable level of OGTA076 or said one or more fragments thereof or a detectable level of nucleic acid encoding OGTA076 or the presence of a detectable level of OGTA076 activity in the candidate subject as compared with a corresponding undetectable level in a healthy subject indicates the presence of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in said subject.

[0023] Further described is a method of monitoring the progression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in a subject or of monitoring the effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy which comprises detecting the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 in said candidate subject at a first time point and at a later time point, the presence of an elevated or lowered level of OGTA076 or said one or more fragments thereof or an elevated or lowered level of nucleic acid encoding OGTA076 or the presence of an elevated or lowered level of OGTA076 activity in the subject at the later time point as compared with the level in the subject at said first time point, indicating the progression or regression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or indicating the effect or non-effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy in said subject.

[0024] The presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the

presence of the activity of OGTA076 may, for example, be detected by analysis of a biological sample obtained from said subject.

[0025] The method described may typically include the step of obtaining a biological sample for analysis from said subject. However, in one or more examples the method does not include the step of obtaining the biological sample.

[0026] The biological sample used can be from any source such as a serum sample or a tissue sample e.g. colorectal, kidney, liver, lung, lymphoid, ovarian or pancreatic tissue. For instance, when looking for evidence of metastatic colorectal cancer, kidney cancer, liver cancer, lung cancer, ovarian cancer or pancreatic cancer, one would look at major sites of colorectal cancer, kidney cancer, liver cancer, lung cancer, ovarian cancer or pancreatic cancer metastasis, e.g. the liver, the peritoneal cavity, the pelvis, the retroperitoneum and the lungs for colorectal cancer; the bones, the lungs and the liver for kidney cancer; the lungs and bones for liver cancer; the brain, the liver, the bones and adrenal glands for lung cancer; the abdomen for ovarian cancer and the liver for pancreatic cancer.

[0027] Alternatively the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 may be detected by analysis *in situ*.

[0028] In certain examples, methods of diagnosis described herein may be at least partly, or wholly, performed *in vitro*.

[0029] Suitably the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 is detected quantitatively.

[0030] For example, quantitatively detecting may comprise:

1. (a) contacting a biological sample with an affinity reagent that is specific for OGTA076, said affinity reagent optionally being conjugated to a detectable label; and
2. (b) detecting whether binding has occurred between the affinity reagent and at least one species in the sample, said detection being performed either directly or indirectly.

[0031] Alternatively the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 may be detected quantitatively by means involving use of an imaging technology.

[0032] In another example, the method of the disclosure involves use of immunohistochemistry on tissue sections in order to determine the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076, and thereby to localise colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer cells.

[0033] In one example the presence of OGTA076 or one or more epitope-containing fragments thereof is detected, for example using an affinity reagent capable of specific binding to OGTA076 or one or more fragments thereof, such as an antibody.

[0034] In another example the activity of OGTA076 is detected.

[0035] According to one aspect of the invention there is provided a method of detecting, diagnosing and/or screening for or monitoring the progression of pancreatic cancer or of monitoring the effect of an anti-pancreatic cancer drug or therapy in a subject which comprises detecting the presence or level of antibodies capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof or which comprises detecting a change in the level thereof in said subject.

[0036] According to another aspect of the invention there is also provided a method of detecting, diagnosing and/or screening for pancreatic cancer in a subject which comprises detecting the presence of antibodies capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof in said subject, in which (a) the presence of an elevated level of antibodies capable of immunospecific binding to OGTA076 or said one or more epitope-containing fragments thereof in said subject as compared with the level in a healthy subject or (b) the presence of a detectable level of antibodies capable of immunospecific binding to OGTA076 or said one or more epitope-containing fragments thereof in said subject as compared with a corresponding undetectable level in a healthy subject indicates the presence of pancreatic cancer in said subject.

[0037] One particular a method of detecting, diagnosing and/or screening for pancreatic cancer comprises:

1. (a) bringing into contact with a biological sample to be tested OGTA076, or one or more epitope-containing fragments thereof; and
2. (b) detecting the presence of antibodies in the subject capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof

[0038] Also described is a method of monitoring the progression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or of monitoring the effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy in a subject which comprises detecting the presence of antibodies capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof in said subject at a first time point and at a later time point, the presence of an elevated or lowered level of antibodies capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof in said subject at the later time point as compared with the level in said subject at said first time point, indicating the progression or regression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or the effect or non-effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy in said subject.

[0039] The presence of antibodies capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof is typically detected by analysis of a biological sample obtained from said subject (exemplary biological samples are mentioned above, e.g. the sample is a sample of colorectal, kidney, liver, lung, lymphoid, ovarian or pancreatic tissue, or else a sample of blood or saliva).

[0040] The method may include the step of obtaining said biological sample for analysis from said subject.

[0041] The antibodies that may be detected include IgA, IgM and IgG antibodies.

[0042] In any of the above methods, the level that may be detected in the candidate subject who has colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer is 2 or more fold higher than the level in the healthy subject.

[0043] Also disclosed is an agent capable of specific binding to OGTA076, or a fragment thereof, or a hybridising agent capable of hybridizing to nucleic acid encoding OGTA076 or an agent capable of detecting the activity of OGTA076 for use in screening for, detecting and/or diagnosing disease, such as cancer, and especially colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.

[0044] Also described is OGTA076, or a fragment thereof for use in screening for, detecting and/or diagnosing disease such as cancer, and especially colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.

[0045] Further described is an affinity reagent capable of specific binding to OGTA076 or a fragment thereof, for example an affinity reagent which contains or is conjugated to a detectable label or contains or is conjugated to a therapeutic moiety such as a cytotoxic moiety. The affinity reagent may, for example, be an antibody.

[0046] Also described is a hybridizing agent capable of hybridizing to nucleic acid encoding OGTA076, for example, a hybridizing agent which contains or is conjugated to a detectable label. One example of a hybridizing agent is an inhibitory RNA (RNAi). Other examples include anti-sense oligonucleotides and ribozymes.

[0047] The disclosure also describes a kit containing OGTA076 and/or one or more fragments thereof or containing one or more aforementioned affinity reagents and/or hybridizing agents or containing one or more agents capable of detecting the activity of OGTA076 together with instructions for their use in an aforementioned method. The kit may further contain reagents capable of detecting and reporting the binding of said affinity reagents and/or hybridizing agents to their binding partners.

[0048] Also disclosed is a pharmaceutical composition comprising a therapeutically effective amount of an affinity reagent capable of specific binding to OGTA076 or a fragment thereof.

[0049] Further disclosed is a pharmaceutically acceptable diluent or carrier and a pharmaceutical composition comprising one or more affinity reagents or hybridizing reagents as aforesaid and a pharmaceutically acceptable diluent or carrier.

[0050] In one example the cancer to be detected, prevented or treated is colorectal cancer.

[0051] In another example the cancer to be detected, prevented or treated is kidney cancer.

[0052] In another example the cancer to be detected, prevented or treated is liver cancer.

[0053] In another example the cancer to be detected, prevented or treated is lung cancer.

[0054] In another example the cancer to be detected, prevented or treated is lymphoid leukaemia (particularly chronic lymphocytic leukaemia).

[0055] In another example the cancer to be detected, prevented or treated is ovarian cancer

[0056] In another embodiment the cancer to be detected, or treated is pancreatic cancer.

BRIEF DESCRIPTION OF THE FIGURES

[0057]

Figure 1 shows the amino acid sequences of the three isoforms of the protein of the disclosure. The tryptic peptides detected experimentally by mass spectrometry are highlighted - mass match peptides are shown in bold, tandem peptides are underlined.

Figure 2 shows the Protein Index for the protein of the disclosure.

DETAILED DESCRIPTION OF THE INVENTION

[0058] The invention described in detail below provides methods and compositions for clinical screening, diagnosis and prognosis of pancreatic cancer. Also described are methods and compositions for clinical screening, diagnosis and prognosis of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia) or ovarian cancer in a mammalian subject for identifying patients most likely to respond to a particular therapeutic treatment, for monitoring the results of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer therapy, for drug screening and drug development. The invention also encompasses the administration of therapeutic compositions to a mammalian subject to treat pancreatic cancer. The disclosure also describes the administration of therapeutic compositions to a mammalian subject to treat or prevent colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer. The mammalian subject may be a non-human mammal, for example a human, such as a human adult, i.e. a human subject at least 21 (for example at least 35, at least 50, at least 60, at least 70, or at least 80) years old. The methods and compositions of the present invention are specially suited for screening, diagnosis and prognosis of a living subject, but may also be used for postmortem diagnosis in a subject, for example, to identify family members at risk of developing the same disease.

OGTA076

[0059] As described herein, one-dimensional electrophoresis or isobaric tags for relative and absolute quantification (iTRAQ) or other appropriate methods are used to analyze colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer tissue samples from a subject, preferably a living subject, in order to measure the expression of the protein for screening or diagnosis of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer, to determine the prognosis of a colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer patient, to monitor the effectiveness of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or

pancreatic cancer therapy, or for drug development.

[0060] As used herein, the term "Protein of the disclosure", or "OGTA076", refers to the protein illustrated in Figure 1 in all its isoforms, in particular in its three different isoforms detected experimentally by 1D gel electrophoresis and iTRAQ analysis of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer tissue samples (OGTA076a [SEQ ID No: 1]; OGTA076b [SEQ ID No: 2] and OGTA076c [SEQ ID No: 3]). Protein derivatives of these sequences may also be useful for the same purposes as described herein.

[0061] This protein has been identified in membrane protein extracts of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer tissue samples from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer patients, through the methods and apparatus of the Preferred Technologies (1D gel electrophoresis or iTRAQ together with tryptic digest of membrane protein extracts). Peptide sequences were compared to the SWISS-PROT and trEMBL databases (held by the Swiss Institute of Bioinformatics (SIB) and the European Bioinformatics Institute (EBI) which are available at www.expasy.com), and the following entry: 060449, Lymphocyte antigen 75, was identified.

[0062] According to SWISS-PROT, Lymphocyte antigen 75 is expressed in spleen, thymus, colon and peripheral blood lymphocytes. It has been detected in myeloid and B lymphoid cell lines. Isoforms OGTA076b and OGTA076c are expressed in malignant Hodgkin's lymphoma cells called Hodgkin's and Reed-Sternberg (HRS) cells. Lymphocyte antigen 75 acts as an endocytic receptor to direct captured antigens from the extracellular space to a specialized antigen-processing compartment. It causes reduced proliferation of B-lymphocytes.

[0063] The protein of the disclosure is useful as are fragments particularly epitope containing fragments e.g. antigenic or immunogenic fragments thereof and derivatives thereof. Epitope containing fragments including antigenic or immunogenic fragments will typically be of length 12 amino acids or more e.g. 20 amino acids or more e.g. 50 or 100 amino acids or more. Fragments may be 95% or more of the length of the full protein e.g. 90% or more e.g. 75% or 50% or 25% or 10% or more of the length of the full protein.

[0064] Epitope containing fragments including antigenic or immunogenic fragments will be capable of eliciting a relevant immune response in a patient, for example when presented to the immune system in an appropriate form such as in a formulation containing one or more adjuvants. DNA encoding the protein of the disclosure is also useful as are fragments thereof e.g. DNA encoding fragments of the protein of the disclosure such as immunogenic fragments thereof. Fragments of nucleic acid (e.g. DNA) encoding the protein of the disclosure may be 95% or more of the length of the full coding region e.g. 90% or more e.g. 75% or 50% or 25% or 10% or more of the length of the full coding region. Fragments of nucleic acid (e.g. DNA) may be 36 nucleotides or more e.g. 60 nucleotides or more e.g. 150 or 300 nucleotides or more in length.

[0065] Derivatives of the protein of the disclosure include variants on the sequence in which one or more (e.g. 1-20 such as 15 amino acids, or up to 20% such as up to 10% or 5% or 1% by number of amino acids based on the total length of the protein) deletions, insertions or substitutions have been made. Substitutions may typically be conservative substitutions. Derivatives will typically have essentially the same biological function as the protein from which they are derived. Derivatives will typically be comparably antigenic or immunogenic to the protein from which they are derived. Derivatives will typically have either the ligand-binding activity, or the active receptor-complex forming ability, or preferably both, of the protein from which they are derived.

[0066] Tables 1a-1f below illustrates the different occurrences of OGTA076 as detected by iTRAQ and mass spectrometry of membrane protein extracts of colorectal, kidney, liver, lung and ovarian tissue samples from colorectal cancer, kidney cancer, liver cancer, non-small cell lung cancer, small cell lung cancer and ovarian cancer patients respectively. The first column provides the samples batch number, the second column gives the iTRAQ experiment number for those samples and the last column provides a list of the sequences observed by mass spectrometry and the corresponding SEQ ID Nos.

[0067] Table 2a-2c below illustrates the different occurrences of OGTA076 as detected by 1D gel electrophoresis and mass spectrometry of membrane protein extracts of colorectal, lymphoid and pancreatic tissue samples from colorectal cancer, chronic lymphocytic leukaemia and pancreatic cancer patients respectively. The first column provides the molecular weight, the second column gives information on the subfractionation protocol used, if any (see Example 1 below), and the last column provides a list of the sequences observed by mass spectrometry and the corresponding SEQ ID Nos.

Table 1a - Colorectal cancer iTRAQ

Samples batch no.	Experiment no.	Tryptics identified [SEQ ID No]
Samples 1	Experiment 1	EGIAK [14], IIPK [28], LALK [35], LNPK [40]

Samples batch no.	Experiment no.	Tryptics identified [SEQ ID No]
Samples 1	Experiment 2	EGIAK [14], LALK [35]
Samples 1	Experiment 3	EGIAK [14], IIPK [28], LALK [35]
Samples 1	Experiment 4	LNPK [40]

Table 1b - Kidney cancer iTRAQ

Sample no.	Experiment no.	Tryptics identified [SEQ ID No]
Samples 1	Experiment 1	EGIAK [14], LALK [35], LNPK [40], YLNNLYK [66]
Samples 1	Experiment 2	LALK [35]

Table 1c - Liver cancer iTRAQ

Sample no.	Experiment no.	Tryptics identified [SEQ ID No]
Samples 1	Experiment 1	YLNNLYK [66]

Table 1d - Non-small cell lung cancer iTRAQ

Sample no.	Experiment no.	Tryptics identified [SEQ ID No]
Samples 1	Experiment 1	YLNNLYK [66]
Samples 1	Experiment 2	LALK [35]
Samples 2	Experiment 1	IIPK [28], LALK [35], YLNNLYK [66]
Samples 2	Experiment 2	YLNNLYK [66]

Table 1e- Small cell lung cancer iTRAQ

Sample no.	Experiment no.	Tryptics identified [SEQ ID No]
Samples 1	Experiment 1	EGIAK [14]
Samples 1	Experiment 2	EGIAK [14]

Table 1f - Ovarian cancer iTRAQ

Sample no.	Experiment no.	Tryptics identified [SEQ ID No]
Samples 1	Experiment 1	LALK [35]
Samples 1	Experiment 2	LALK [35], YLNNLYK [66]
Samples 1	Experiment 3	LALK [35]

Table 2a - Colorectal cancer 1D gel

MW (Da)	Subfractionation	Tryptics identified [SEQ ID No]
167535		CLGLDITK [7], DGAICYKPTK [9], DVDSCEYNWATVGGRR [11], EEVWIGLK [12], ENNNITMR [16], EVKPVDSVK [17], GWHFYDDR [21], HDHSATIVSIK [22], HFVSLCQE [23], HGETCYK [24], IPENFFEEESR [29], KYFWTGGLR [34], LNDASSDK [39], LPFICEK [41], MCPDDEGWKR [42], MFSCDSSAMLWK [43], NNSLMWFDK [45], QTLQNASETVK [47], RHGETCYK [49], TLTWHSAK [56], TPDWYNPDR [57]

Table 2b - Chronic lymphocytic leukaemia 1D gel

MW (Da)	Subfractionation	Tryptics identified [SEQ ID No]
198008		AANDPFTIVHGNTGK [4], CSMLIASNETWKK [8], DGAICYKPTK [9], ELTYSNFHPLLVSQR [15], FEQEYLNNDLMK [18], GWHFYDDR [21], HDHSATIVSIK [22], HMATTQDEVHTK [25], IPENFFEEESR [29], ISEWPIDDHFTYSR [30], KYFWTGGLR [34], LFHLHSQK [36], NWEEAER [46], RGWHFYDDR [48], SHILSIR [53], SPDQGSWQWSDR [55], TPLSYTHWR [58], TPVSTIIMPNEFQQDYDIR [59], VFHRPWR [61], VQCSEQWIPFQNK [63], WVSQHR [64], YFWTGGLR [65]

MW (Da)	Subfractionation	Tryptics identified [SEQ ID No]
223568		AANDPFTIVHGNTGK [4], DGAICYKPTK [9], GWHFYDDR [21], HDHSATIVSIK [22], HFVSLCQK [23], HMATTQDEVHTK [25], IPENFFEEESR [29], KVECEHGFGFGR [33], LFHLHSQK [36], LHNEDIK [38], NWEEEAER [46], SDQALHSFSEAK [52], SHILSIR [53], SNFHPLLVSGR [54], TPLSYTHWR [58], VFHRPWR [61], VQCSEQWIPFQNK [63], WVSQHR [64], YLNNLYK [66]
238644		CEHHSLYGAAR [6], CLGLDITK [7], IEMVVDYK [27], IPENFFEEESR [29], KGNCEVSSVEGTLCK [31], KVECEHGFGFGR [33], LFHLHSQK [36], LHNEDIK [38], RGWHFYDDR [48], RLHFSR [50], RNWEEEAER [51], SDQALHSFSEAK [52], SHILSIR [53], TPLSYTHWR [58], VCEHGFGFGR [60], YFWTGLR [65], YLNNLYK [66]

Table 2c - Pancreatic cancer 1D gel

MW (Da)	Subfractionation	Tryptics identified [SEQ ID No]
161314		DGAICYKPTK [9], EFiYLRPFACDTK [13], FPVTGEECLYMSAK [19], GWHFYDDR [21], HFVSLCQK [23], IPENFFEEESR [29], ISEWPIDDHFTYSR [30], KRNWEEEAER [32], NNSLMWFDK [45], RGWHFYDDR [48], RHGETCYK [49], SNFHPLLVSGR [54], SPDLQGSWQWSDR [55], TLTWHSAK [56], TPDWYNPDR [57], TPLSYTHWR [58], VQCSEQWIPFQNK [63], WVSQHR [64]
168430		DGAICYKPTK [9], EEWIGLK [12], EFiYLRPFACDTK [13], FEQEYLNDLMK [18], FPVTGEECLYMSAK [19], GWHFYDDR [21], HFVSLCQK [23], IANISGDGQK [26], IPENFFEEESR [29], ISEWPIDDHFTYSR [30], KRNWEEEAER [32], KYFWTGLR [34], MSGPLGPEEASPK [44], NNSLMWFDK [45], RGWHFYDDR [48], RHGETCYK [49], SDQALHSFSEAK [52], SHILSIR [53], SPDLQGSWQWSDR [55], TPDWYNPDR [57], TPLSYTHWR [58], TPVSTIIMPNEFQQDYDIR [59], VCEHGFGFGR [60], VFHRPWR [61], VIEEAVYFHQH [62], VQCSEQWIPFQNK [63], WVSQHR [64], YFWTGLR [65]
176135		DGAICYKPTK [9], EEWIGLK [12], FEQEYLNDLMK [18], FPVTGEECLYMSAK [19], GNCEVSSVEGTLCK [20], GWHFYDDR [21], HMATTQDEVHTK [25], IANISGDGQK [26], IPENFFEEESR [29], ISEWPIDDHFTYSR [30], LHLAGFSSVR [37], NNSLMWFDK [45], RNWEEEAER [51], SDQALHSFSEAK [52], SPDLQGSWQWSDR [55], TPDWYNPDR [57], TPLSYTHWR [58], VCEHGFGFGR [60], VFHRPWR [61], VQCSEQWIPFQNK [63], WVSQHR [64], YFWTGLR [65]
207419		AFSSDLISIHSLADVEVVVTK [5], CLGLDITK [7], DVDSCEYNWATVGGR [11], EEWIGLK [12], FEQEYLNDLMK [18], FPVTGEECLYMSAK [19], HMATTQDEVHTK [25], IPENFFEEESR [29], RGWHFYDDR [48], RHGETCYK [49], SHILSIR [53], TLTWHSAK [56]

MW (Da)	Subfractionation	Tryptics identified [SEQ ID No]
261976		AFSSDLISIHSLADVEVVTK [5], DGAICYKPTK [9], DGHGTAISNASDVWK [10], DWDSCGEYNWATVGGR [11], EEWVIGLK [12], EFIYLRPACDTK [13], FEQEYLNDLMK [18], FPVTGEECLYMSAK [19], GWHFYDDR [21], HMATTQDEVHTK [25], IPENFEEESR [29], ISEWPIDDHFTYSR [30], LHLAGFSSVR [37], RGWHFYDDR [48], SPDQGSWQWSDR [55], TPLSYTHWR [58], TPVSTIIMPNEFQQDYDIR [59], VFHRPWR [61], VIEEAVYFHQH [62], YFWTGLR [65]

[0068] For OGTA076, the detected level obtained upon analyzing tissue from subjects having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer relative to the detected level obtained upon analyzing tissue from subjects free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer will depend upon the particular analytical protocol and detection technique that is used. Accordingly, the present disclosure contemplates that each laboratory will establish a reference range in subjects free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer according to the analytical protocol and detection technique in use, as is conventional in the diagnostic art. Preferably, at least one control positive tissue sample from a subject known to have colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or at least one control negative tissue sample from a subject known to be free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer (and more preferably both positive and negative control samples) are included in each batch of test samples analysed.

[0069] OGTA076 can be used for detection, prognosis, diagnosis, or monitoring of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or for drug development. In one example, tissue from a subject (e.g. a subject suspected of having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer) is analysed by 1D gel electrophoresis or iTRAQ for detection of OGTA076. An increased abundance of OGTA076 in the tissue from the subject relative to tissue from a subject or subjects free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer (e.g. a control sample) or a previously determined reference range indicates the presence of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.

[0070] In relation to fragments, epitope containing fragments, immunogenic fragments or antigenic fragments of OGTA076:

- for colorectal cancer applications, these may comprise the sequences identified as tryptic sequences in the 3rd column of Table 1a or the 3rd column of Table 2a;
- for kidney cancer applications, these may comprise the sequences identified as tryptic sequences in the 3rd column of Table 1b;
- for liver cancer applications, these may comprise the sequences identified as tryptic sequences in the 3rd column of Table 1c;
- for non-small cell lung cancer applications, these may comprise the sequences identified as tryptic sequences in the 3rd column of Table 1d;
- for small cell lung cancer applications, these may comprise the sequences identified as tryptic sequences in the 3rd column of Table 1e;
- for ovarian cancer applications, these may comprise the sequences identified as tryptic sequences in the 3rd column of Table 1f;
- for chronic lymphocytic leukaemia applications, these may comprise the sequences identified as tryptic sequences in the 3rd column of Table 2b;
- for pancreatic cancer applications, these may comprise the sequences identified as tryptic sequences in the 3rd column of Table 2c.

[0071] As used herein, OGTA076 is "isolated" when it is present in a preparation that is substantially free of contaminating proteins, i.e. a preparation in which less than 10% (for example less than 5%, such as less than 1%) of the total protein present is contaminating protein(s). A contaminating protein is a protein having a significantly different amino acid sequence from that of isolated OGTA076, as determined by mass spectral analysis. As used herein, a "significantly different" sequence is one that permits the contaminating protein to be resolved from OGTA076 by mass spectral analysis, performed according to the Reference Protocols.

[0072] OGTA076 can be assayed by any method known to those skilled in the art, including but not limited to, the Preferred Technologies described herein, kinase assays, enzyme assays, binding assays and other functional assays, immunoassays, and western blotting. In one example, OGTA076 is separated on a 1-D gel by virtue of its MW and visualized by staining the gel. In one example, OGTA076 is stained with a fluorescent dye and imaged with a fluorescence scanner. Sypro Red (Molecular Probes, Inc., Eugene, Oregon) is a suitable dye for this purpose. A preferred fluorescent dye is disclosed in U.S. Application No. 09/412,168, filed on October 5, 1999,. In another example, OGTA076 is analysed using isobaric tags for relative and absolute quantification (iTRAQ).

[0073] Alternatively, OGTA076 can be detected in an immunoassay. In one example, an immunoassay is performed by contacting a sample from a subject to be tested with an anti-OGTA076 antibody (or other affinity reagent) under conditions such that binding (e.g. immunospecific binding) can occur if OGTA076 is present, and detecting or measuring the amount of any binding (e.g. immunospecific binding) by the agent. OGTA076 binding agents can be produced by the methods and techniques taught herein.

[0074] OGTA076 may be detected by virtue of the detection of a fragment thereof e.g. an epitope containing (e.g. an immunogenic or antigenic) fragment thereof. Fragments may have a length of at least 10, more typically at least 20 amino acids e.g. at least 50 or 100 amino acids e.g. at least 200 or 500 amino acids e.g. at least 1000 or 1500 amino acids.

[0075] In one or more examples OGTA076 is differentially expressed in patients/subjects with colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Differentially expressed is intended to refer to a higher or lower level of expression in a relevant sample in comparison to cancer free/healthy sample.

[0076] In one example, binding of an affinity reagent (e.g. an antibody) in tissue sections can be used to detect aberrant OGTA076 localization or an aberrant level of OGTA076. In a specific example, an antibody (or other affinity reagent) to OGTA076 can be used to assay a patient tissue (e.g. a colorectal, kidney, liver, lung, lymphoid, ovarian or pancreatic tissue) for the level of OGTA076 where an aberrant level of OGTA076 is indicative of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. As used herein, an "aberrant level" means a level that is increased compared with the level in a subject free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer or a reference level.

[0077] Any suitable immunoassay can be used, including, without limitation, competitive and non-competitive assay systems using techniques such as western blots, radioimmunoassays, ELISA (enzyme linked immunosorbent assay), "sandwich" immunoassays, immunoprecipitation assays, precipitin reactions, gel diffusion precipitin reactions, immunodiffusion assays, agglutination assays, complement-fixation assays, immunoradiometric assays, fluorescent immunoassays and protein A immunoassays.

[0078] For example, OGTA076 can be detected in a fluid sample (e.g. blood, urine, or saliva) by means of a two-step sandwich assay. In the first step, a capture reagent (e.g. an anti-OGTA076 antibody or other affinity reagent) is used to capture OGTA076. The capture reagent can optionally be immobilized on a solid phase. In the second step, a directly or indirectly labeled detection reagent is used to detect the captured OGTA076. In one example, the detection reagent is a lectin. Any lectin can be used for this purpose that, for example binds to OGTA076 rather than to other isoforms that have the same core protein as OGTA076 or to other proteins that share the antigenic determinant recognized by the antibody. In a preferred example, the chosen lectin binds OGTA076 with at least 2-fold greater affinity, more preferably at least 5-fold greater affinity, still more preferably at least 10-fold greater affinity, than to said other isoforms that have the same core protein as OGTA076 or to said other proteins that share the antigenic determinant recognized by the affinity reagent. Based on the present description, a lectin that is suitable for detecting OGTA076 can readily be identified by methods well known in the art, for instance upon testing one or more lectins enumerated in Table I on pages 158-159 of Sumar et al., Lectins as Indicators of Disease-Associated Glycoforms, In: Gabius H-J & Gabius S

(eds.), 1993, Lectins and Glycobiology, at pp. 158-. In an alternative example, the detection reagent is an antibody (or other affinity reagent), e.g. an antibody that specifically (e.g. immunospecifically) detects other post-translational modifications, such as an antibody that immunospecifically binds to phosphorylated amino acids. Examples of such antibodies include those that bind to phosphotyrosine (BD Transduction Laboratories, catalog nos.: P11230-050/P11230-150; P11120; P38820; P39020), those that bind to phosphoserine (Zymed Laboratories Inc., South San Francisco, CA, catalog no. 61-8100) and those that bind to phosphothreonine (Zymed Laboratories Inc., South San Francisco, CA, catalogue nos. 71-8200, 13-9200).

[0079] If desired, a gene encoding OGTA076, a related gene, or related nucleic acid sequences or subsequences, including complementary sequences, can also be used in hybridization assays. A nucleotide encoding OGTA076, or subsequences thereof comprising at least 8 nucleotides, for example at least 12 nucleotides, such as at least 15 nucleotides can be used as a hybridization probe. Hybridization assays can be used for detection, prognosis, diagnosis, or monitoring of conditions, disorders, or disease states, associated with aberrant expression of the gene encoding OGTA076, or for differential diagnosis of subjects with signs or symptoms suggestive of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. In particular, such a hybridization assay can be carried out by a method comprising contacting a subject's sample containing nucleic acid with a nucleic acid probe capable of hybridizing to a DNA or RNA that encodes OGTA076, under conditions such that hybridization can occur, and detecting or measuring any resulting hybridization.

[0080] Hence nucleic acid encoding OGTA076 (e.g. DNA or more suitably RNA) may be detected, for example, using a hybridizing agent capable of hybridizing to nucleic acid encoding OGTA076.

[0081] One such exemplary method comprises:

1. (a) contacting one or more oligonucleotide probes comprising 10 or more consecutive nucleotides complementary to a nucleotide sequence encoding OGTA076, with an RNA obtained from a biological sample from the subject or with cDNA copied from the RNA, wherein said contacting occurs under conditions that permit hybridization of the probe to the nucleotide sequence if present;
2. (b) detecting hybridization, if any, between the probe and the nucleotide sequence; and
3. (c) comparing the hybridization, if any, detected in step (b) with the hybridization detected in a control sample, or with a previously determined reference range.

[0082] The disclosure also provides diagnostic kits, comprising an anti-OGTA076 antibody (or other affinity reagent). In addition, such a kit may optionally comprise one or more of the following: (1) instructions for using the anti-OGTA076 affinity reagent for diagnosis, prognosis, therapeutic monitoring or any combination of these applications; (2) a labeled binding partner to the affinity reagent; (3) a solid phase (such as a reagent strip) upon which the anti-OGTA076 affinity reagent is immobilized; and (4) a label or insert indicating regulatory approval for diagnostic, prognostic or therapeutic use or any combination thereof. If no labeled binding partner to the affinity reagent is provided, the anti-OGTA076 affinity reagent itself can be labeled with a detectable marker, e.g. a chemiluminescent, enzymatic, fluorescent, or radioactive moiety.

[0083] The invention also provides a kit comprising a nucleic acid probe capable of hybridizing to nucleic acid, suitably RNA, encoding OGTA076. In a specific example, a kit comprises in one or more containers a pair of primers (e.g. each in the size range of 6-30 nucleotides, more preferably 10-30 nucleotides and still more preferably 10-20 nucleotides) that under appropriate reaction conditions can prime amplification of at least a portion of a nucleic acid encoding OGTA076, such as by polymerase chain reaction (see, e.g. Innis et al., 1990, PCR Protocols, Academic Press, Inc., San Diego, CA), ligase chain reaction (see EP 320,308) use of Q β replicase, cyclic probe reaction, or other methods known in the art.

[0084] A kit can optionally further comprise a predetermined amount of OGTA076 or a nucleic acid encoding OGTA076, e.g. for use as a standard or control.

Use in Clinical Studies

[0085] The diagnostic methods and compositions of the present invention can assist in monitoring a clinical study, e.g. to evaluate drugs for therapy of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. In one example, candidate molecules are tested for their ability to restore OGTA076 levels in a subject having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia

(particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer to levels found in subjects free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer or, in a treated subject, to preserve OGTA076 levels at or near non-colorectal cancer, non-kidney cancer, non-liver cancer, non-lung cancer, non-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), non-ovarian cancer or non-pancreatic cancer values.

[0086] In another example, the methods and compositions of the present invention are used to screen candidates for a clinical study to identify individuals having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer; such individuals can then be excluded from the study or can be placed in a separate cohort for treatment or analysis.

Production of Protein of the Disclosure and Corresponding Nucleic Acid

[0087] A DNA of the present disclosure can be obtained by isolation as a cDNA fragment from cDNA libraries using as starter materials commercial mRNAs and determining and identifying the nucleotide sequences thereof. That is, specifically, clones are randomly isolated from cDNA libraries, which are prepared according to Ohara et al's method (DNA Research Vol.4, 53-59 (1997)). Next, through hybridization, duplicated clones (which appear repeatedly) are removed and then in vitro transcription and translation are carried out. Nucleotide sequences of both termini of clones, for which products of 50 kDa or more are confirmed, are determined.

Furthermore, databases of known genes are searched for homology using the thus obtained terminal nucleotide sequences as queries. The entire nucleotide sequence of a clone revealed to be novel as a result is determined. In addition to the above screening method, the 5' and 3' terminal sequences of cDNA are related to a human genome sequence. Then an unknown long-chain gene is confirmed in a region between the sequences, and the full-length of the cDNA is analyzed. In this way, an unknown gene that is unable to be obtained by a conventional cloning method that depends on known genes can be systematically cloned.

[0088] Moreover, all of the regions of a human-derived gene containing a DNA of the present disclosure can also be prepared using a PCR method such as RACE while paying sufficient attention to prevent artificial errors from taking place in short fragments or obtained sequences. As described above, clones having DNA of the present disclosure can be obtained.

[0089] In another means for cloning DNA of the present disclosure, a synthetic DNA primer having an appropriate nucleotide sequence of a portion of a polypeptide of the present disclosure is produced, followed by amplification by the PCR method using an appropriate library. Alternatively, selection can be carried out by hybridization of the DNA of the present disclosure with a DNA that has been incorporated into an appropriate vector and labeled with a DNA fragment or a synthetic DNA encoding some or all of the regions of the polypeptide of the present disclosure. Hybridization can be carried out by, for example, the method described in Current Protocols in Molecular Biology (edited by Frederick M. Ausubel et al., 1987). DNA of the present disclosure may be any DNA, as long as they contain nucleotide sequences encoding the polypeptides of the present disclosure as described above. Such a DNA may be a cDNA identified and isolated from cDNA libraries or the like that are derived from colorectal, kidney, liver, lung, lymphoid, ovarian or pancreatic tissue. Such a DNA may also be a synthetic DNA or the like. Vectors for use in library construction may be any of bacteriophages, plasmids, cosmids, phagemids, or the like. Furthermore, by the use of a total RNA fraction or a mRNA fraction prepared from the above cells and/or tissues, amplification can be carried out by a direct reverse transcription coupled polymerase chain reaction (hereinafter abbreviated as "RT-PCR method").

[0090] DNA encoding the above polypeptide consisting of an amino acid sequence that is substantially identical to the amino acid sequence of OGTA076 or DNA encoding the above polypeptide consisting of an amino acid sequence derived from the amino acid sequence of OGTA076 by deletion, substitution, or addition of one or more amino acids composing a portion of the amino acid sequence can be easily produced by an appropriate combination of, for example, a site-directed mutagenesis method, a gene homologous recombination method, a primer elongation method, and the PCR method known by persons skilled in the art. In addition, at this time, a possible method for causing a polypeptide to have substantially equivalent biological activity is substitution of homologous amino acids (e.g. polar and nonpolar amino acids, hydrophobic and hydrophilic amino acids, positively-charged and negatively charged amino acids, and aromatic amino acids) among amino acids composing the polypeptide. Furthermore, to maintain substantially equivalent biological activity, amino acids within functional domains contained in the polypeptide of the present disclosure are preferably conserved.

[0091] Furthermore, examples of DNA of the present disclosure include DNA comprising a nucleotide sequence that encodes the amino acid sequence of OGTA076 and DNA hybridizing under stringent conditions to the DNA and encoding a polypeptide (protein) having biological activity (function) equivalent to the function of the polypeptide consisting of the amino acid sequence of

OGTA076. Under such conditions, an example of such DNA capable of hybridizing to DNA comprising the nucleotide sequence that encodes the amino acid sequence of OGTA076 is DNA comprising a nucleotide sequence that has a degree of overall mean homology with the entire nucleotide sequence of the DNA, such as approximately 80% or more, preferably approximately 90% or more, and more preferably approximately 95% or more. Hybridization can be carried out according to a method known in the art such as a method described in Current Protocols in Molecular Biology (edited by Frederick M. Ausubel et al., 1987) or a method according thereto. Here, "stringent conditions" are, for example, conditions of approximately "1*SSC, 0.1% SDS, and 37°C, more stringent conditions of approximately "0.5*SSC, 0.1% SDS, and 42°C, or even more stringent conditions of approximately "0.2*SSC, 0.1% SDS, and 65°C. With more stringent hybridization conditions, the isolation of a DNA having high homology with a probe sequence can be expected. The above combinations of SSC, SDS, and temperature conditions are given for illustrative purposes. Stringency similar to the above can be achieved by persons skilled in the art using an appropriate combination of the above factors or other factors (for example, probe concentration, probe length, and reaction time for hybridization) for determination of hybridization stringency.

[0092] A cloned DNA of the present disclosure can be directly used or used, if desired, after digestion with a restriction enzyme or addition of a linker, depending on purposes. The DNA may have ATG as a translation initiation codon at the 5' terminal side and have TAA, TGA, or TAG as a translation termination codon at the 3' terminal side. These translation initiation and translation termination codons can also be added using an appropriate synthetic DNA adapter.

[0093] Where it is provided for use with the methods of the disclosure OGTA076 may, for example be provided in isolated form. In one example the OGTA076 polypeptide has been purified to at least to some extent. OGTA076 polypeptide may be provided in substantially pure form, that is to say free, to a substantial extent, from other proteins. OGTA076 polypeptide can also be produced using recombinant methods, synthetically produced or produced by a combination of these methods. OGTA076 can be easily prepared by any method known by persons skilled in the art, which involves producing an expression vector containing a DNA of the present disclosure or a gene containing a DNA of the present disclosure, culturing a transformant transformed using the expression vector, generating and accumulating a polypeptide of the present disclosure or a recombinant protein containing the polypeptide, and then collecting the resultant.

[0094] Recombinant OGTA076 polypeptide may be prepared by processes well known in the art from genetically engineered host cells comprising expression systems. Accordingly, the present disclosure also relates to expression systems which comprise an OGTA076 polypeptide or nucleic acid, to host cells which are genetically engineered with such expression systems and to the production of OGTA076 polypeptide by recombinant techniques. For recombinant OGTA076 polypeptide production, host cells can be genetically engineered to incorporate expression systems or portions thereof for nucleic acids. Such incorporation can be performed using methods well known in the art, such as, calcium phosphate transfection, DEAE-dextran mediated transfection, transvection, microinjection, cationic lipid-mediated transfection, electroporation, transduction, scrape loading, ballistic introduction or infection (see e.g. Davis et al., *Basic Methods in Molecular Biology*, 1986 and Sambrook et al., *Molecular Cloning: A Laboratory Manual*, 2nd Ed., Cold Spring Harbour Laboratory Press, Cold Spring Harbour, NY, 1989).

[0095] As host cells, for example, bacteria of the genus *Escherichia*, *Streptococci*, *Staphylococci*, *Streptomyces*, bacteria of the genus *Bacillus*, yeast, *Aspergillus* cells, insect cells, insects, and animal cells are used. Specific examples of bacteria of the genus *Escherichia*, which are used herein, include *Escherichia coli* K12 and DH1 (Proc. Natl. Acad. Sci. U.S.A., Vol. 60, 160 (1968)), JM103 (Nucleic Acids Research, Vol. 9, 309 (1981)), JA221 (Journal of Molecular Biology, Vol. 120, 517 (1978)), and HB101 (Journal of Molecular Biology, Vol. 41, 459 (1969)). As bacteria of the genus *Bacillus*, for example, *Bacillus subtilis* M114 (Gene, Vol. 24, 255 (1983)) and 207-21 (Journal of Biochemistry, Vol. 95, 87 (1984)) are used. As yeast, for example, *Saccharomyces cerevisiae* AH22, AH22R-, NA87-11A, DKD-5D, and 20B-12, *Schizosaccharomyces pombe* NCYC1913 and NCYC2036, and *Pichia pastoris* are used. As insect cells, for example, *Drosophila* S2 and *Spodoptera* Sf9 cells are used. As animal cells, for example, COS-7 and Vero monkey cells, CHO Chinese hamster cells (hereinafter abbreviated as CHO cells), dhfr-gene-deficient CHO cells, mouse L cells, mouse AtT-20 cells, mouse myeloma cells, rat GH3 cells, human FL cells, COS, HeLa, C127,3T3, HEK 293, BHK and Bowes melanoma cells are used.

[0096] Cell-free translation systems can also be employed to produce recombinant polypeptides (e.g. rabbit reticulocyte lysate, wheat germ lysate, SP6/T7 in vitro T&T and RTS 100 E. Coli HY transcription and translation kits from Roche Diagnostics Ltd., Lewes, UK and the TNT Quick coupled Transcription/Translation System from Promega UK, Southampton, UK).

[0097] The expression vector can be produced according to a method known in the art. For example, the vector can be produced by (1) excising a DNA fragment containing a DNA of the present disclosure or a gene containing a DNA of the present disclosure and (2) ligating the DNA fragment downstream of the promoter in an appropriate expression vector. A wide variety of expression systems can be used, such as and without limitation, chromosomal, episomal and virus-derived systems, e.g. plasmids derived from *Escherichia coli* (e.g. pBR322, pBR325, pUC18, and pUC118), plasmids derived from *Bacillus subtilis* (e.g. pUB110,

pTP5, and pC194), from bacteriophage, from transposons, from yeast episomes (e.g. pSH19 and pSH15), from insertion elements, from yeast chromosomal elements, from viruses such as baculoviruses, papova viruses such as SV40, vaccinia viruses, adenoviruses, fowl pox viruses, pseudorabies viruses and retroviruses, and vectors derived from combinations thereof, such as those derived from plasmid and bacteriophage (such as $[\lambda]$ phage) genetic elements, such as cosmids and phagemids. The expression systems may contain control regions that regulate as well as engender expression. Promoters to be used in the present disclosure may be any promoters as long as they are appropriate for hosts to be used for gene expression. For example, when a host is Escherichia coli, a trp promoter, a lac promoter, a recA promoter, a pL promoter, an lpp promoter, and the like are preferred. When a host is Bacillus subtilis, an SPO1 promoter, an SPO2 promoter, a penP promoter, and the like are preferred. When a host is yeast, a PHO5 promoter, a PGK promoter, a GAP promoter, an ADH promoter, and the like are preferred. When an animal cell is used as a host, examples of promoters for use in this case include an SRα promoter, an SV40 promoter, an LTR promoter, a CMV promoter, and an HSV-TK promoter. Generally, any system or vector that is able to maintain, propagate or express a nucleic acid to produce a polypeptide in a host may be used.

[0098] The appropriate nucleic acid sequence may be inserted into an expression system by any variety of well known and routine techniques, such as those set forth in Sambrook et al., *supra*. Appropriate secretion signals may be incorporated into the OGTA076 polypeptide to allow secretion of the translated protein into the lumen of the endoplasmic reticulum, the periplasmic space or the extracellular environment. These signals may be endogenous to the OGTA076 polypeptide or they may be heterologous signals. Transformation of the host cells can be carried out according to methods known in the art. For example, the following documents can be referred to: Proc. Natl. Acad. Sci. U.S.A., Vol. 69, 2110 (1972); Gene, Vol. 17, 107 (1982); Molecular & General Genetics, Vol. 168, 111 (1979); Methods in Enzymology, Vol. 194, 182-187 (1991); Proc. Natl. Acad. Sci. U.S.A.), Vol. 75, 1929 (1978); Cell Technology, separate volume 8, New Cell Technology, Experimental Protocol. 263-267 (1995) (issued by Shujunsha); and Virology, Vol. 52, 456 (1973). The thus obtained transformant transformed with an expression vector containing a DNA of the present disclosure or a gene containing a DNA of the present disclosure can be cultured according to a method known in the art. For example, when hosts are bacteria of the genus Escherichia, the bacteria are generally cultured at approximately 15°C to 43°C for approximately 3 to 24 hours. If necessary, aeration or agitation can also be added. When hosts are bacteria of the genus Bacillus, the bacteria are generally cultured at approximately 30°C to 40°C for approximately 6 to 24 hours. If necessary, aeration or agitation can also be added. When transformants whose hosts are yeast are cultured, culture is generally carried out at approximately 20°C to 35°C for approximately 24 to 72 hours using media with pH adjusted to be approximately 5 to 8. If necessary, aeration or agitation can also be added. When transformants whose hosts are animal cells are cultured, the cells are generally cultured at approximately 30°C to 40°C for approximately 15 to 60 hours using media with the pH adjusted to be approximately 6 to 8. If necessary, aeration or agitation can also be added.

[0099] If an OGTA076 polypeptide is to be expressed for use in cell-based screening assays, it is preferred that the polypeptide be produced at the cell surface. In this event, the cells may be harvested prior to use in the screening assay. If the OGTA076 polypeptide is secreted into the medium, the medium can be recovered in order to isolate said polypeptide. If produced intracellularly, the cells must first be lysed before the OGTA076 polypeptide is recovered.

[0100] OGTA076 polypeptide can be recovered and purified from recombinant cell cultures or from other biological sources by well known methods including, ammonium sulphate or ethanol precipitation, acid extraction, anion or cation exchange chromatography, phosphocellulose chromatography, affinity chromatography, hydrophobic interaction chromatography, hydroxylapatite chromatography, molecular sieving chromatography, centrifugation methods, electrophoresis methods and lectin chromatography. In one example, a combination of these methods is used. In another example, high performance liquid chromatography is used. In a further example, an antibody which specifically binds to an OGTA076 polypeptide can be used to deplete a sample comprising an OGTA076 polypeptide of said polypeptide or to purify said polypeptide.

[0101] To separate and purify a polypeptide or a protein of the present disclosure from the culture products, for example, after culture, microbial bodies or cells are collected by a known method, they are suspended in an appropriate buffer, the microbial bodies or the cells are disrupted by, for example, ultrasonic waves, lysozymes, and/or freeze-thawing, the resultant is then subjected to centrifugation or filtration, and then a crude extract of the protein can be obtained. The buffer may also contain a protein denaturation agent such as urea or guanidine hydrochloride or a surfactant such as Triton X-100(TM). When the protein is secreted in a culture solution, microbial bodies or cells and a supernatant are separated by a known method after the completion of culture and then the supernatant is collected. The protein contained in the thus obtained culture supernatant or the extract can be purified by an appropriate combination of known separation and purification methods. The thus obtained polypeptide (protein) of the present disclosure can be converted into a salt by a known method or a method according thereto. Conversely, when the polypeptide (protein) of the present disclosure is obtained in the form of a salt, it can be converted into a free protein or peptide or another salt by a known method or a method according thereto. Moreover, an appropriate protein modification enzyme such as trypsin or chymotrypsin is caused to act on a protein produced by a recombinant before or after

purification, so that modification can be arbitrarily added or a polypeptide can be partially removed. The presence of a polypeptide (protein) of the present disclosure or a salt thereof can be measured by various binding assays, enzyme immunoassays using specific antibodies, and the like.

[0102] Techniques well known in the art may be used for refolding to regenerate native or active conformations of the OGTA076 polypeptide when the polypeptide has been denatured during isolation and or purification. In the context of the present disclosure, OGTA076 polypeptide can be obtained from a biological sample from any source, such as and without limitation, a blood sample or tissue sample, e.g. a colorectal, kidney, liver, lung, lymphoid, ovarian or pancreatic tissue sample.

[0103] OGTA076 polypeptide may be in the form of a "mature protein" or may be part of a larger protein such as a fusion protein. It is often advantageous to include an additional amino acid sequence which contains secretory or leader sequences, a pre-, pro- or prepro-protein sequence, or a sequence which aids in purification such as an affinity tag, for example, but without limitation, multiple histidine residues, a FLAG tag, HA tag or myc tag.

[0104] An additional sequence that may provide stability during recombinant production may also be used. Such sequences may be optionally removed as required by incorporating a cleavable sequence as an additional sequence or part thereof. Thus, an OGTA076 polypeptide may be fused to other moieties including other polypeptides or proteins (for example, glutathione S-transferase and protein A). Such a fusion protein can be cleaved using an appropriate protease, and then separated into each protein. Such additional sequences and affinity tags are well known in the art. In addition to the above, features known in the art, such as an enhancer, a splicing signal, a polyA addition signal, a selection marker, and an SV40 replication origin can be added to an expression vector, if desired.

Production of Affinity Reagents to OGTA076

[0105] According to those in the art, there are three main types of immunoaffinity reagent - monoclonal antibodies, phage display antibodies and smaller antibody-derived molecules such as Affibodies, Domain Antibodies (dAbs), Nanobodies or UniBodies. In general in applications according to the present invention where the use of antibodies is stated, other affinity reagents (e.g. Affibodies, domain antibodies, Nanobodies or UniBodies) may be employed. Such substances may be said to be capable of immunospecific binding to OGTA076. Where appropriate the term "affinity agent" shall be construed to embrace immunoaffinity reagents and other substances capable of specific binding to OGTA076 including but not limited to ligands, lectins, streptavidins, antibody mimetics and synthetic binding agents.

Production of Antibodies to OGTA076

[0106] According to the invention OGTA076, an OGTA076 analog, an OGTA076-related protein or a fragment or derivative of any of the foregoing may be used as an immunogen to generate antibodies which immunospecifically bind such an immunogen. Such immunogens can be isolated by any convenient means, including the methods described above. The term "antibody" as used herein refers to a peptide or polypeptide derived from, modeled after or substantially encoded by an immunoglobulin gene or immunoglobulin genes, or fragments thereof, capable of specifically binding an antigen or epitope. See, e.g. Fundamental Immunology, 3rd Edition, W.E. Paul, ed., Raven Press, N.Y. (1993); Wilson (1994) J. Immunol. Methods 175:267-273; Yarmush (1992) J. Biochem. Biophys. Methods 25:85-97. The term antibody includes antigen-binding portions, i.e., "antigen binding sites" (e.g. fragments, subsequences, complementarity determining regions (CDRs)) that retain capacity to bind antigen, including (i) a Fab fragment, a monovalent fragment consisting of the VL, VH, CL and CH1 domains; (ii) a F(ab')₂ fragment, a bivalent fragment comprising two Fab fragments linked by a disulfide bridge at the hinge region; (iii) a Fd fragment consisting of the VH and CH1 domains; (iv) a Fv fragment consisting of the VL and VH domains of a single arm of an antibody, (v) a dAb fragment (Ward et al., (1989) Nature 341:544-546), which consists of a VH domain; and (vi) an isolated complementarity determining region (CDR). Single chain antibodies are also included by reference in the term "antibody." Antibodies of the invention include, but are not limited to polyclonal, monoclonal, bispecific, humanized or chimeric antibodies, single chain antibodies, Fab fragments and F(ab')₂ fragments, fragments produced by a Fab expression library, anti-idiotypic (anti-Id) antibodies, and epitope-binding fragments of any of the above. The immunoglobulin molecules of the invention can be of any class (e.g. IgG, IgE, IgM, IgD and IgA) or subclass of immunoglobulin molecule.

[0107] The term "specifically binds" (or "immunospecifically binds") is not intended to indicate that an antibody binds exclusively to its intended target. Rather, an antibody "specifically binds" if its affinity for its intended target is typically about 5-fold greater when compared to its affinity for a non-target molecule. Suitably there is no significant cross-reaction or cross-binding with

undesired substances, especially naturally occurring proteins or tissues of a healthy person or animal. In one embodiment the affinity of the antibody will be at least about 5 fold, for example 10 fold, such as 25-fold, particularly 50-fold, and especially 100-fold or more, greater for a target molecule than its affinity for a non-target molecule. In some embodiments, specific binding between an antibody or other binding agent and an antigen means a binding affinity of at least 10^6 M^{-1} . Antibodies bind with affinities of at least about 10^7 M^{-1} , for example between about 10^8 M^{-1} to about 10^9 M^{-1} , about 10^9 M^{-1} to about 10^{10} M^{-1} , or about 10^{10} M^{-1} to about 10^{11} M^{-1} may be particularly suitable for use in the invention.

[0108] Affinity is calculated as $K_d = k_{off}/k_{on}$ (k_{off} is the dissociation rate constant, k_{on} is the association rate constant and K_d is the equilibrium constant. Affinity can be determined at equilibrium by measuring the fraction bound (r) of labeled ligand at various concentrations (c). The data are graphed using the Scatchard equation:

(c). The data are graphed using the Scatchard equation: $r/c = K(n-r)$.

where

r = moles of bound ligand/mole of receptor at equilibrium;

c = free ligand concentration at equilibrium;

K = equilibrium association constant; and

n = number of ligand binding sites per receptor molecule

By graphical analysis, r/c is plotted on the Y-axis versus r on the X-axis thus producing a Scatchard plot. The affinity is the negative slope of the line. k_{off} can be determined by competing bound labeled ligand with unlabeled excess ligand (see, e.g. U.S. Pat No. 6,316,409). The affinity of a targeting agent for its target molecule is preferably at least about 1×10^{-6} moles/liter, is more preferably at least about 1×10^{-7} moles/liter, is even more preferably at least about 1×10^{-8} moles/liter, is yet even more preferably at least about 1×10^{-9} moles/liter, and is most preferably at least about 1×10^{-10} moles/liter. Antibody affinity measurement by Scatchard analysis is well known in the art. See, e.g. van Erp et al., *J. Immunoassay* 12: 425-43, 1991; Nelson and Griswold, *Comput. Methods Programs Biomed.* 27: 65-8, 1988.

[0109] In one embodiment, antibodies that recognize gene products of genes encoding OGTA076 are publicly available. In another embodiment, methods known to those skilled in the art are used to produce antibodies that recognize OGTA076, an OGTA076 analog, an OGTA076-related polypeptide, or a fragment or derivative of any of the foregoing. One skilled in the art will recognize that many procedures are available for the production of antibodies, for example, as described in *Antibodies, A Laboratory Manual*, Ed Harlow and David Lane, Cold Spring Harbor Laboratory (1988), Cold Spring Harbor, N.Y. One skilled in the art will also appreciate that binding fragments or Fab fragments which mimic antibodies can also be prepared from genetic information by various procedures (*Antibody Engineering: A Practical Approach* (Borrebaek, C., ed.), 1995, Oxford University Press, Oxford; *J. Immunol.* 149, 3914-3920 (1992)).

[0110] In one example, antibodies to a specific domain of OGTA076 are produced. In a specific example, hydrophilic fragments of OGTA076 are used as immunogens for antibody production.

[0111] In the production of antibodies, screening for the desired antibody can be accomplished by techniques known in the art, e.g. ELISA (enzyme-linked immunosorbent assay). For example, to select antibodies which recognize a specific domain of OGTA076, one may assay generated hybridomas for a product which binds to an OGTA076 fragment containing such domain. For selection of an antibody that specifically binds a first OGTA076 homolog but which does not specifically bind to (or binds less avidly to) a second OGTA076 homolog, one can select on the basis of positive binding to the first OGTA076 homolog and a lack of binding to (or reduced binding to) the second OGTA076 homolog. Similarly, for selection of an antibody that specifically binds OGTA076 but which does not specifically bind to (or binds less avidly to) a different isoform of the same protein (such as a different glycoform having the same core peptide as OGTA076), one can select on the basis of positive binding to OGTA076 and a lack of binding to (or reduced binding to) the different isoform (e.g. a different glycoform). Thus, the present invention provides an antibody (preferably a monoclonal antibody) that binds with greater affinity (for example at least 2-fold, such as at least 5-fold, particularly at least 10-fold greater affinity) to OGTA076 than to a different isoform or isoforms (e.g. glycoforms) of OGTA076.

[0112] Polyclonal antibodies which may be used in the methods of the invention are heterogeneous populations of antibody molecules derived from the sera of immunized animals. Unfractionated immune serum can also be used. Various procedures known in the art may be used for the production of polyclonal antibodies to OGTA076, a fragment of OGTA076, an OGTA076-related polypeptide, or a fragment of an OGTA076-related polypeptide. For example, one way is to purify polypeptides of interest or to synthesize the polypeptides of interest using, e.g. solid phase peptide synthesis methods well known in the art. See, e.g.

Guide to Protein Purification, Murray P. Deutcher, ed., Meth. Enzymol. Vol 182 (1990); Solid Phase Peptide Synthesis, Greg B. Fields ed., Meth. Enzymol. Vol 289 (1997); Kiso et al., Chem. Pharm. Bull. (Tokyo) 38: 1192-99, 1990; Mostafavi et al., Biomed. Pept. Proteins Nucleic Acids 1: 255-60, 1995; Fujiwara et al., Chem. Pharm. Bull. (Tokyo) 44: 1326-31, 1996. The selected polypeptides may then be used to immunize by injection various host animals, including but not limited to rabbits, mice, rats, etc., to generate polyclonal or monoclonal antibodies. The Preferred Technology described herein in Example 1 provides isolated OGTA076 suitable for such immunization. If OGTA076 is purified by gel electrophoresis, OGTA076 can be used for immunization with or without prior extraction from the polyacrylamide gel. Various adjuvants (i.e. immunostimulants) may be used to enhance the immunological response, depending on the host species, including, but not limited to, complete or incomplete Freund's adjuvant, a mineral gel such as aluminum hydroxide, surface active substance such as lysolecithin, pluronic polyol, a polyanion, a peptide, an oil emulsion, keyhole limpet hemocyanin, dinitrophenol, and an adjuvant such as BCG (bacille Calmette-Guerin) or corynebacterium parvum. Additional adjuvants are also well known in the art.

[0113] For preparation of monoclonal antibodies (mAbs) directed toward OGTA076, a fragment of OGTA076, an OGTA076-related polypeptide, or a fragment of an OGTA076-related polypeptide, any technique which provides for the production of antibody molecules by continuous cell lines in culture may be used. For example, the hybridoma technique originally developed by Kohler and Milstein (1975, *Nature* 256:495-497), as well as the trioma technique, the human B-cell hybridoma technique (Kozbor et al., 1983, *Immunology Today* 4:72), and the EBV-hybridoma technique to produce human monoclonal antibodies (Cole et al., 1985, in *Monoclonal Antibodies and Cancer Therapy*, Alan R. Liss, Inc., pp. 77-96). Such antibodies may be of any immunoglobulin class including IgG, IgM, IgE, IgA, IgD and any subclass thereof. The hybridoma producing the mAbs of the invention may be cultivated *in vitro* or *in vivo*. In an additional example, monoclonal antibodies can be produced in germ-free animals utilizing known technology (PCT/US90/02545).

[0114] The monoclonal antibodies include but are not limited to human monoclonal antibodies and chimeric monoclonal antibodies (e.g. human-mouse chimeras). A chimeric antibody is a molecule in which different portions are derived from different animal species, such as those having a human immunoglobulin constant region and a variable region derived from a murine mAb. (See, e.g. Cabilly et al., U.S. Patent No. 4,816,567; and Boss et al., U.S. Patent No. 4,816,397.) Humanized antibodies are antibody molecules from non-human species having one or more complementarity determining regions (CDRs) from the non-human species and a framework region from a human immunoglobulin molecule. (See, e.g. Queen, U.S. Patent No. 5,585,089.)

[0115] Chimeric and humanized monoclonal antibodies can be produced by recombinant DNA techniques known in the art, for example using methods described in PCT Publication No. WO 87/02671; European Patent Application 184,187; European Patent Application 171,496; European Patent Application 173,494; PCT Publication No. WO 86/01533; U.S. Patent No. 4,816,567; European Patent Application 125,023; Better et al., 1988, *Science* 240:1041-1043; Liu et al., 1987, *Proc. Natl. Acad. Sci. USA* 84:3439-3443; Liu et al., 1987, *J. Immunol.* 139:3521-3526; Sun et al., 1987, *Proc. Natl. Acad. Sci. USA* 84:214-218; Nishimura et al., 1987, *Canc. Res.* 47:999-1005; Wood et al., 1985, *Nature* 314:446-449; and Shaw et al., 1988, *J. Natl. Cancer Inst.* 80:1553-1559; Morrison, 1985, *Science* 229:1202-1207; Oi et al., 1986, *Bio/Techniques* 4:214; U.S. Patent 5,225,539; Jones et al., 1986, *Nature* 321:552-525; Verhoeyan et al. (1988) *Science* 239:1534; and Beidler et al., 1988, *J. Immunol.* 141:4053-4060.

[0116] Completely human antibodies are particularly desirable for therapeutic treatment of human subjects. Such antibodies can be produced using transgenic mice which are incapable of expressing endogenous immunoglobulin heavy and light chain genes, but which can express human heavy and light chain genes. The transgenic mice are immunized in the normal fashion with a selected antigen, e.g. all or a portion of OGTA076. Monoclonal antibodies directed against the antigen can be obtained using conventional hybridoma technology. The human immunoglobulin transgenes harbored by the transgenic mice rearrange during B cell differentiation, and subsequently undergo class switching and somatic mutation. Thus, using such a technique, it is possible to produce therapeutically useful IgG, IgA, IgM and IgE antibodies. For an overview of this technology for producing human antibodies, see Lonberg and Huszar (1995, *Int. Rev. Immunol.* 13:65-93). For a detailed discussion of this technology for producing human antibodies and human monoclonal antibodies and protocols for producing such antibodies, see, e.g. U.S. Patent 5,625,126; U.S. Patent 5,633,425; U.S. Patent 5,569,825; U.S. Patent 5,661,016; and U.S. Patent 5,545,806. In addition, companies such as Abgenix, Inc. (Freemont, CA) and Genpharm (San Jose, CA) can be engaged to provide human antibodies directed against a selected antigen using technology similar to that described above.

[0117] Completely human antibodies which recognize a selected epitope can be generated using a technique referred to as "guided selection". In this approach a selected non-human monoclonal antibody, e.g. a mouse antibody, is used to guide the selection of a completely human antibody recognizing the same epitope. (Jespers et al. (1994) *Bio/technology* 12:899-903).

[0118] The antibodies of the present invention can also be generated by the use of phage display technology to produce and screen libraries of polypeptides for binding to a selected target. See, e.g. Cwirla et al., *Proc. Natl. Acad. Sci. USA* 87, 6378-82, 1990; Devlin et al., *Science* 249, 404-6, 1990, Scott and Smith, *Science* 249, 386-88, 1990; and Ladner et al., U.S. Pat. No.

5,571,698. A basic concept of phage display methods is the establishment of a physical association between DNA encoding a polypeptide to be screened and the polypeptide. This physical association is provided by the phage particle, which displays a polypeptide as part of a capsid enclosing the phage genome which encodes the polypeptide. The establishment of a physical association between polypeptides and their genetic material allows simultaneous mass screening of very large numbers of phage bearing different polypeptides. Phage displaying a polypeptide with affinity to a target bind to the target and these phage are enriched by affinity screening to the target. The identity of polypeptides displayed from these phage can be determined from their respective genomes. Using these methods a polypeptide identified as having a binding affinity for a desired target can then be synthesized in bulk by conventional means. See, e.g. U.S. Patent No. 6,057,098. In particular, such phage can be utilized to display antigen binding domains expressed from a repertoire or combinatorial antibody library (e.g. human or murine). Phage expressing an antigen binding domain that binds the antigen of interest can be selected or identified with antigen, e.g. using labeled antigen or antigen bound or captured to a solid surface or bead. Phage used in these methods are typically filamentous phage including fd and M13 binding domains expressed from phage with Fab, Fv or disulfide stabilized Fv antibody domains recombinantly fused to either the phage gene III or gene VIII protein. Phage display methods that can be used to make the antibodies of the present invention include those disclosed in Brinkman et al., *J. Immunol. Methods* 182:41-50 (1995); Ames et al., *J. Immunol. Methods* 184:177-186 (1995); Kettleborough et al., *Eur. J. Immunol.* 24:952-958 (1994); Persic et al., *Gene* 187 9-18 (1997); Burton et al., *Advances in Immunology* 57:191-280 (1994); PCT Application No. PCT/GB91/01134; PCT Publications WO 90/02809; WO 91/10737; WO 92/01047; WO 92/18619; WO 93/11236; WO 95/15982; WO 95/20401; and U.S. Patent Nos. 5,698,426; 5,223,409; 5,403,484; 5,580,717; 5,427,908; 5,750,753; 5,821,047; 5,571,698; 5,427,908; 5,516,637; 5,780,225; 5,658,727; 5,733,743 and 5,969,108.

[0119] As described in the above references, after phage selection, the antibody coding regions from the phage can be isolated and used to generate whole antibodies, including human antibodies, or any other desired antigen binding fragment, and expressed in any desired host, including mammalian cells, insect cells, plant cells, yeast, and bacteria, e.g. as described in detail below. For example, techniques to recombinantly produce Fab, Fab' and F(ab')₂ fragments can also be employed using methods known in the art such as those disclosed in PCT publication WO 92/22324; Mullinax et al., *BioTechniques* 12(6):864-869 (1992); and Sawai et al., *AJRI* 34:26-34 (1995); and Better et al., *Science* 240:1041-1043 (1988).

[0120] Examples of techniques which can be used to produce single-chain Fvs and antibodies include those described in U.S. Patents 4,946,778 and 5,258,498; Huston et al., *Methods in Enzymology* 203:46-88 (1991); Shu et al., *PNAS* 90:7995-7999 (1993); and Skerra et al., *Science* 240:1038-1040 (1988).

[0121] The invention further provides for the use of bispecific antibodies, which can be made by methods known in the art. Traditional production of full length bispecific antibodies is based on the coexpression of two immunoglobulin heavy chain-light chain pairs, where the two chains have different specificities (Milstein et al., 1983, *Nature* 305:537-539). Because of the random assortment of immunoglobulin heavy and light chains, these hybridomas (quadromas) produce a potential mixture of 10 different antibody molecules, of which only one has the correct bispecific structure. Purification of the correct molecule, which is usually done by affinity chromatography steps, is rather cumbersome, and the product yields are low. Similar procedures are disclosed in WO 93/08829, published 13 May 1993, and in Traunecker et al., 1991, *EMBO J.* 10:3655-3659.

[0122] According to a different and more preferred approach, antibody variable domains with the desired binding specificities (antibody-antigen combining sites) are fused to immunoglobulin constant domain sequences. The fusion preferably is with an immunoglobulin heavy chain constant domain, comprising at least part of the hinge, CH₂, and CH₃ regions. It is preferred to have the first heavy-chain constant region (CH₁) containing the site necessary for light chain binding, present in at least one of the fusions. DNAs encoding the immunoglobulin heavy chain fusions and, if desired, the immunoglobulin light chain, are inserted into separate expression vectors, and are co-transfected into a suitable host organism. This provides for great flexibility in adjusting the mutual proportions of the three polypeptide fragments in examples when unequal ratios of the three polypeptide chains used in the construction provide the optimum yields. It is, however, possible to insert the coding sequences for two or all three polypeptide chains in one expression vector when the expression of at least two polypeptide chains in equal ratios results in high yields or when the ratios are of no particular significance.

[0123] In one example of this approach, the bispecific antibodies are composed of a hybrid immunoglobulin heavy chain with a first binding specificity in one arm, and a hybrid immunoglobulin heavy chain-light chain pair (providing a second binding specificity) in the other arm. It was found that this asymmetric structure facilitates the separation of the desired bispecific compound from unwanted immunoglobulin chain combinations, as the presence of an immunoglobulin light chain in only one half of the bispecific molecule provides for a facile way of separation. This approach is disclosed in WO 94/04690 published March 3, 1994. For further details for generating bispecific antibodies see, for example, Suresh et al., *Methods in Enzymology*, 1986, 121:210.

[0124] The invention provides functionally active fragments, derivatives or analogs of the anti-OGTA076 immunoglobulin molecules. Functionally active means that the fragment, derivative or analog is able to elicit anti-anti-idiotype antibodies (*i.e.*, tertiary antibodies) that recognize the same antigen that is recognized by the antibody from which the fragment, derivative or analog is derived. Specifically, in a preferred example the antigenicity of the idiotype of the immunoglobulin molecule may be enhanced by deletion of framework and CDR sequences that are C-terminal to the CDR sequence that specifically recognizes the antigen. To determine which CDR sequences bind the antigen, synthetic peptides containing the CDR sequences can be used in binding assays with the antigen by any binding assay method known in the art.

[0125] The present invention provides antibody fragments such as, but not limited to, F(ab')₂ fragments and Fab fragments. Antibody fragments which recognize specific epitopes may be generated by known techniques. F(ab')₂ fragments consist of the variable region, the light chain constant region and the CH1 domain of the heavy chain and are generated by pepsin digestion of the antibody molecule. Fab fragments are generated by reducing the disulfide bridges of the F(ab')₂ fragments. The invention also provides heavy chain and light chain dimers of the antibodies of the invention, or any minimal fragment thereof such as Fvs or single chain antibodies (SCAs) (*e.g.* as described in U.S. Patent 4,946,778; Bird, 1988, *Science* 242:423-42; Huston et al., 1988, *Proc. Natl. Acad. Sci. USA* 85:5879-5883; and Ward et al., 1989, *Nature* 334:544-54), or any other molecule with the same specificity as the antibody of the invention. Single chain antibodies are formed by linking the heavy and light chain fragments of the Fv region via an amino acid bridge, resulting in a single chain polypeptide. Techniques for the assembly of functional Fv fragments in *E. coli* may be used (Skerra et al., 1988, *Science* 242:1038-1041).

[0126] In other examples, the disclosure provides fusion proteins of the immunoglobulins of the invention (or functionally active fragments thereof), for example in which the immunoglobulin is fused via a covalent bond (*e.g.* a peptide bond), at either the N-terminus or the C-terminus to an amino acid sequence of another protein (or portion thereof, preferably at least 10, 20 or 50 amino acid portion of the protein) that is not the immunoglobulin. Preferably the immunoglobulin, or fragment thereof, is covalently linked to the other protein at the N-terminus of the constant domain. As stated above, such fusion proteins may facilitate purification, increase half-life *in vivo*, and enhance the delivery of an antigen across an epithelial barrier to the immune system.

[0127] The immunoglobulins of the invention include analogs and derivatives that are modified, *i.e.*, by the covalent attachment of any type of molecule as long as such covalent attachment does not impair immunospecific binding. For example, but not by way of limitation, the derivatives and analogs of the immunoglobulins include those that have been further modified, *e.g.* by glycosylation, acetylation, pegylation, phosphorylation, amidation, derivatization by known protecting/blocking groups, proteolytic cleavage, linkage to a cellular ligand or other protein, etc. Any of numerous chemical modifications may be carried out by known techniques, including, but not limited to specific chemical cleavage, acetylation, formylation, etc. Additionally, the analog or derivative may contain one or more non-classical amino acids.

[0128] The foregoing antibodies can be used in methods known in the art relating to the localization and activity of OGTA076, *e.g.* for imaging this protein, measuring levels thereof in appropriate physiological samples, in diagnostic methods, etc.

Production of Affibodies to OGTA076

[0129] Affibody molecules represent a new class of affinity proteins based on a 58-amino acid residue protein domain, derived from one of the IgG-binding domains of staphylococcal protein A. This three helix bundle domain has been used as a scaffold for the construction of combinatorial phagemid libraries, from which Affibody variants that target the desired molecules can be selected using phage display technology (Nord K, Gunnarsson E, Ringdahl J, Stahl S, Uhlen M, Nygren PA, Binding proteins selected from combinatorial libraries of an α -helical bacterial receptor domain, *Nat Biotechnol* 1997;15:772-7. Ronmark J, Gronlund H, Uhlen M, Nygren PA, Human immunoglobulin A (IgA)-specific ligands from combinatorial engineering of protein A, *Eur J Biochem* 2002;269:2647-55.). The simple, robust structure of Affibody molecules in combination with their low molecular weight (6 kDa), make them suitable for a wide variety of applications, for instance, as detection reagents (Ronmark J, Hansson M, Nguyen T, et al, Construction and characterization of Affibody-Fc chimeras produced in *Escherichia coli*, *J Immunol Methods* 2002;261:199-211) and to inhibit receptor interactions (Sandstrom K, Xu Z, Forsberg G, Nygren PA, Inhibition of the CD28-CD80 co-stimulation signal by a CD28-binding Affibody ligand developed by combinatorial protein engineering, *Protein Eng* 2003;16:691-7). Further details of Affibodies and methods of production thereof may be obtained by reference to US Patent No 5831012.

[0130] Labelled Affibodies may also be useful in imaging applications for determining abundance of Isoforms.

Production of Domain Antibodies to OGTA076

[0131] References to antibodies herein embrace references to Domain Antibodies. Domain Antibodies (dAbs) are the smallest functional binding units of antibodies, corresponding to the variable regions of either the heavy (V_H) or light (V_L) chains of human antibodies. Domain Antibodies have a molecular weight of approximately 13 kDa. Domantis has developed a series of large and highly functional libraries of fully human V_H and V_L dAbs (more than ten billion different sequences in each library), and uses these libraries to select dAbs that are specific to therapeutic targets. In contrast to many conventional antibodies, Domain Antibodies are well expressed in bacterial, yeast, and mammalian cell systems. Further details of domain antibodies and methods of production thereof may be obtained by reference to US Patent 6,291,158; 6,582,915; 6,593,081; 6,172,197; 6,696,245; US Serial No. 2004/0110941; European patent application No. 1433846 and European Patents 0368684 & 0616640; WO05/035572, WO04/101790, WO04/081026, WO04/058821, WO04/003019 and WO03/002609.

Production of Nanobodies to OGTA076

[0132] Nanobodies are antibody-derived therapeutic proteins that contain the unique structural and functional properties of naturally-occurring heavy-chain antibodies. These heavy-chain antibodies contain a single variable domain (VHH) and two constant domains (C_H2 and C_H3). Importantly, the cloned and isolated VHH domain is a perfectly stable polypeptide harbouring the full antigen-binding capacity of the original heavy-chain antibody. Nanobodies have a high homology with the VH domains of human antibodies and can be further humanised without any loss of activity. Importantly, Nanobodies have a low immunogenic potential, which has been confirmed in primate studies with Nanobody lead compounds.

[0133] Nanobodies combine the advantages of conventional antibodies with important features of small molecule drugs. Like conventional antibodies, Nanobodies show high target specificity, high affinity for their target and low inherent toxicity. However, like small molecule drugs they can inhibit enzymes and readily access receptor clefts. Furthermore, Nanobodies are extremely stable, can be administered by means other than injection (see e.g. WO 04/041867) and are easy to manufacture. Other advantages of Nanobodies include recognising uncommon or hidden epitopes as a result of their small size, binding into cavities or active sites of protein targets with high affinity and selectivity due to their unique 3-dimensional, drug format flexibility, tailoring of half-life and ease and speed of drug discovery.

[0134] Nanobodies are encoded by single genes and are efficiently produced in almost all prokaryotic and eukaryotic hosts e.g. *E. coli* (see e.g. US 6,765,087), moulds (for example *Aspergillus* or *Trichoderma*) and yeast (for example *Saccharomyces*, *Kluyveromyces*, *Hansenula* or *Pichia*) (see e.g. US 6,838,254). The production process is scalable and multi-kilogram quantities of Nanobodies have been produced. Because Nanobodies exhibit a superior stability compared with conventional antibodies, they can be formulated as a long shelf-life, ready-to-use solution.

[0135] The Nanoclone method (see e.g. WO 06/079372) is a proprietary method for generating Nanobodies against a desired target, based on automated high-throughput selection of B-cells.

Production of UniBodies to OGTA076

[0136] UniBody is a new proprietary antibody technology that creates a stable, smaller antibody format with an anticipated longer therapeutic window than current small antibody formats. IgG4 antibodies are considered inert and thus do not interact with the immune system. Genmab modified fully human IgG4 antibodies by eliminating the hinge region of the antibody. Unlike the full size IgG4 antibody, the half molecule fragment is very stable and is termed a UniBody. Halving the IgG4 molecule left only one area on the UniBody that can bind to disease targets and the UniBody therefore binds univalently to only one site on target cells. This univalent binding does not stimulate cancer cells to grow like bivalent antibodies might and opens the door for treatment of some types of cancer which ordinary antibodies cannot treat.

[0137] The UniBody is about half the size of a regular IgG4 antibody. This small size can be a great benefit when treating some forms of cancer, allowing for better distribution of the molecule over larger solid tumors and potentially increasing efficacy.

[0138] Fabs typically do not have a very long half-life. UniBodies, however, were cleared at a similar rate to whole IgG4 antibodies and were able to bind as well as whole antibodies and antibody fragments in pre-clinical studies. Other antibodies

primarily work by killing the targeted cells whereas UniBodies only inhibit or silence the cells.

[0139] Further details of UniBodies may be obtained by reference to patent WO2007/059782.

Expression of Affinity Reagents

Expression of Antibodies

[0140] The antibodies of the invention can be produced by any method known in the art for the synthesis of antibodies, in particular, by chemical synthesis or by recombinant expression, and are preferably produced by recombinant expression techniques.

[0141] Recombinant expression of antibodies, or fragments, derivatives or analogs thereof, requires construction of a nucleic acid that encodes the antibody. If the nucleotide sequence of the antibody is known, a nucleic acid encoding the antibody may be assembled from chemically synthesized oligonucleotides (e.g. as described in Kutmeier et al., 1994, *BioTechniques* 17:242), which, briefly, involves the synthesis of overlapping oligonucleotides containing portions of the sequence encoding antibody, annealing and ligation of those oligonucleotides, and then amplification of the ligated oligonucleotides by PCR.

[0142] Alternatively, the nucleic acid encoding the antibody may be obtained by cloning the antibody. If a clone containing the nucleic acid encoding the particular antibody is not available, but the sequence of the antibody molecule is known, a nucleic acid encoding the antibody may be obtained from a suitable source (e.g. an antibody cDNA library, or cDNA library generated from any tissue or cells expressing the antibody) by PCR amplification using synthetic primers hybridizable to the 3' and 5' ends of the sequence or by cloning using an oligonucleotide probe specific for the particular gene sequence.

[0143] If an antibody molecule that specifically recognizes a particular antigen is not available (or a source for a cDNA library for cloning a nucleic acid encoding such an antibody), antibodies specific for a particular antigen may be generated by any method known in the art, for example, by immunizing an animal, such as a rabbit, to generate polyclonal antibodies or, more preferably, by generating monoclonal antibodies. Alternatively, a clone encoding at least the Fab portion of the antibody may be obtained by screening Fab expression libraries (e.g. as described in Huse et al., 1989, *Science* 246:1275-1281) for clones of Fab fragments that bind the specific antigen or by screening antibody libraries (See, e.g. Clarkson et al., 1991, *Nature* 352:624; Hane et al., 1997 *Proc. Natl. Acad. Sci. USA* 94:4937).

[0144] Once a nucleic acid encoding at least the variable domain of the antibody molecule is obtained, it may be introduced into a vector containing the nucleotide sequence encoding the constant region of the antibody molecule (see, e.g. PCT Publication WO 86/05807; PCT Publication WO 89/01036; and U.S. Patent No. 5,122,464). Vectors containing the complete light or heavy chain for co-expression with the nucleic acid to allow the expression of a complete antibody molecule are also available. Then, the nucleic acid encoding the antibody can be used to introduce the nucleotide substitution(s) or deletion(s) necessary to substitute (or delete) the one or more variable region cysteine residues participating in an intrachain disulfide bond with an amino acid residue that does not contain a sulphydryl group. Such modifications can be carried out by any method known in the art for the introduction of specific mutations or deletions in a nucleotide sequence, for example, but not limited to, chemical mutagenesis, *in vitro* site directed mutagenesis (Hutchinson et al., 1978, *J. Biol. Chem.* 253:6551), PCT based methods, etc.

[0145] In addition, techniques developed for the production of "chimeric antibodies" (Morrison et al., 1984, *Proc. Natl. Acad. Sci.* 81:851-855; Neuberger et al., 1984, *Nature* 312:604-608; Takeda et al., 1985, *Nature* 314:452-454) by splicing genes from a mouse antibody molecule of appropriate antigen specificity together with genes from a human antibody molecule of appropriate biological activity can be used. As described *supra*, a chimeric antibody is a molecule in which different portions are derived from different animal species, such as those having a variable region derived from a murine mAb and a human antibody constant region, e.g. humanized antibodies.

[0146] Once a nucleic acid encoding an antibody molecule of the invention has been obtained, the vector for the production of the antibody molecule may be produced by recombinant DNA technology using techniques well known in the art. Thus, methods for preparing the protein of the invention by expressing nucleic acid containing the antibody molecule sequences are described herein. Methods which are well known to those skilled in the art can be used to construct expression vectors containing an antibody molecule coding sequences and appropriate transcriptional and translational control signals. These methods include, for example, *in vitro* recombinant DNA techniques, synthetic techniques, and *in vivo* genetic recombination. See, for example, the

techniques described in Sambrook et al. (1990, Molecular Cloning, A Laboratory Manual, 2nd Ed., Cold Spring Harbor Laboratory, Cold Spring Harbor, NY) and Ausubel et al. (eds., 1998, Current Protocols in Molecular Biology, John Wiley & Sons, NY).

[0147] The expression vector is transferred to a host cell by conventional techniques and the transfected cells are then cultured by conventional techniques to produce an antibody of the invention.

[0148] The host cells used to express a recombinant antibody of the invention may be either bacterial cells such as *Escherichia coli*, or, preferably, eukaryotic cells, especially for the expression of whole recombinant antibody molecule. In particular, mammalian cells such as Chinese hamster ovary cells (CHO), in conjunction with a vector such as the major intermediate early gene promoter element from human cytomegalovirus are an effective expression system for antibodies (Foecking et al., 1986, Gene 45:101; Cockett et al., 1990, Bio/Technology 8:2).

[0149] A variety of host-expression vector systems may be utilized to express an antibody molecule of the invention. Such host-expression systems represent vehicles by which the coding sequences of interest may be produced and subsequently purified, but also represent cells which may, when transformed or transfected with the appropriate nucleotide coding sequences, express the antibody molecule of the invention *in situ*. These include but are not limited to microorganisms such as bacteria (e.g. *E. coli*, *B. subtilis*) transformed with recombinant bacteriophage DNA, plasmid DNA or cosmid DNA expression vectors containing antibody coding sequences; yeast (e.g. *Saccharomyces*, *Pichia*) transformed with recombinant yeast expression vectors containing antibody coding sequences; insect cell systems infected with recombinant virus expression vectors (e.g. baculovirus) containing the antibody coding sequences; plant cell systems infected with recombinant virus expression vectors (e.g. cauliflower mosaic virus, CaMV; tobacco mosaic virus, TMV) or transformed with recombinant plasmid expression vectors (e.g. Ti plasmid) containing antibody coding sequences; or mammalian cell systems (e.g. COS, CHO, BHK, 293, 3T3 cells) harboring recombinant expression constructs containing promoters derived from the genome of mammalian cells (e.g. metallothionein promoter) or from mammalian viruses (e.g. the adenovirus late promoter; the vaccinia virus 7.5K promoter).

[0150] In bacterial systems, a number of expression vectors may be advantageously selected depending upon the use intended for the antibody molecule being expressed. For example, when a large quantity of such a protein is to be produced, for the generation of pharmaceutical compositions comprising an antibody molecule, vectors which direct the expression of high levels of fusion protein products that are readily purified may be desirable. Such vectors include, but are not limited, to the *E. coli* expression vector pUR278 (Ruther et al., 1983, EMBO J. 2:1791), in which the antibody coding sequence may be ligated individually into the vector in frame with the *lac Z* coding region so that a fusion protein is produced; pIN vectors (Inouye & Inouye, 1985, Nucleic Acids Res. 13:3101-3109; Van Heeke & Schuster, 1989, J. Biol. Chem. 24:5503-5509); and the like. pGEX vectors may also be used to express foreign polypeptides as fusion proteins with glutathione S-transferase (GST). In general, such fusion proteins are soluble and can easily be purified from lysed cells by adsorption and binding to a matrix glutathione-agarose beads followed by elution in the presence of free glutathione. The pGEX vectors are designed to include thrombin or factor Xa protease cleavage sites so that the cloned target gene product can be released from the GST moiety.

[0151] In an insect system, *Autographa californica* nuclear polyhedrosis virus (AcNPV) is used as a vector to express foreign genes. The virus grows in *Spodoptera frugiperda* cells. The antibody coding sequence may be cloned individually into non-essential regions (for example the polyhedrin gene) of the virus and placed under control of an AcNPV promoter (for example the polyhedrin promoter). In mammalian host cells, a number of viral-based expression systems (e.g. an adenovirus expression system) may be utilized.

[0152] As discussed above, a host cell strain may be chosen which modulates the expression of the inserted sequences, or modifies and processes the gene product in the specific fashion desired. Such modifications (e.g. glycosylation) and processing (e.g. cleavage) of protein products may be important for the function of the protein.

[0153] For long-term, high-yield production of recombinant antibodies, stable expression is preferred. For example, cell lines that stably express an antibody of interest can be produced by transfecting the cells with an expression vector comprising the nucleotide sequence of the antibody and the nucleotide sequence of a selectable (e.g. neomycin or hygromycin), and selecting for expression of the selectable marker. Such engineered cell lines may be particularly useful in screening and evaluation of compounds that interact directly or indirectly with the antibody molecule.

[0154] The expression levels of the antibody molecule can be increased by vector amplification (for a review, see Bebbington and Hentschel, The use of vectors based on gene amplification for the expression of cloned genes in mammalian cells in DNA cloning, Vol.3. (Academic Press, New York, 1987)). When a marker in the vector system expressing antibody is amplifiable, increase in the level of inhibitor present in culture of host cell will increase the number of copies of the marker gene. Since the

amplified region is associated with the antibody gene, production of the antibody will also increase (Crouse et al., 1983, Mol. Cell. Biol. 3:257).

[0155] The host cell may be co-transfected with two expression vectors of the disclosure, the first vector encoding a heavy chain derived polypeptide and the second vector encoding a light chain derived polypeptide. The two vectors may contain identical selectable markers which enable equal expression of heavy and light chain polypeptides. Alternatively, a single vector may be used which encodes both heavy and light chain polypeptides. In such situations, the light chain should be placed before the heavy chain to avoid an excess of toxic free heavy chain (Proudfoot, 1986, Nature 322:52; Kohler, 1980, Proc. Natl. Acad. Sci. USA 77:2197). The coding sequences for the heavy and light chains may comprise cDNA or genomic DNA.

[0156] Once the antibody molecule of the invention has been recombinantly expressed, it may be purified by any method known in the art for purification of an antibody molecule, for example, by chromatography (e.g. ion exchange chromatography, affinity chromatography such as with protein A or specific antigen, and sizing column chromatography), centrifugation, differential solubility, or by any other standard technique for the purification of proteins.

[0157] Alternatively, any fusion protein may be readily purified by utilizing an antibody specific for the fusion protein being expressed. For example, a system described by Janknecht et al. allows for the ready purification of non-denatured fusion proteins expressed in human cell lines (Janknecht et al., 1991, Proc. Natl. Acad. Sci. USA 88:8972-897). In this system, the gene of interest is subcloned into a vaccinia recombination plasmid such that the open reading frame of the gene is translationally fused to an amino-terminal tag consisting of six histidine residues. The tag serves as a matrix binding domain for the fusion protein. Extracts from cells infected with recombinant vaccinia virus are loaded onto Ni^{2+} nitriloacetic acid-agarose columns and histidine-tagged proteins are selectively eluted with imidazole-containing buffers.

[0158] The antibodies that are generated by these methods may then be selected by first screening for affinity and specificity with the purified polypeptide of interest and, if required, comparing the results to the affinity and specificity of the antibodies with polypeptides that are desired to be excluded from binding. The screening procedure can involve immobilization of the purified polypeptides in separate wells of microtiter plates. The solution containing a potential antibody or groups of antibodies is then placed into the respective microtiter wells and incubated for about 30 min to 2 h. The microtiter wells are then washed and a labeled secondary antibody (for example, an anti-mouse antibody conjugated to alkaline phosphatase if the raised antibodies are mouse antibodies) is added to the wells and incubated for about 30 min and then washed. Substrate is added to the wells and a color reaction will appear where antibody to the immobilized polypeptide(s) is present.

[0159] The antibodies so identified may then be further analyzed for affinity and specificity in the assay design selected. In the development of immunoassays for a target protein, the purified target protein acts as a standard with which to judge the sensitivity and specificity of the immunoassay using the antibodies that have been selected. Because the binding affinity of various antibodies may differ; certain antibody pairs (e.g. in sandwich assays) may interfere with one another sterically, etc., assay performance of an antibody may be a more important measure than absolute affinity and specificity of an antibody.

[0160] Those skilled in the art will recognize that many approaches can be taken in producing antibodies or binding fragments and screening and selecting for affinity and specificity for the various polypeptides, but these approaches do not change the scope of the invention.

[0161] For therapeutic applications, antibodies (particularly monoclonal antibodies) may suitably be human or humanized animal (e.g. mouse) antibodies. Animal antibodies may be raised in animals using the human protein (e.g. OGTA076) as immunogen. Humanisation typically involves grafting CDRs identified thereby into human framework regions. Normally some subsequent retromutation to optimize the conformation of chains is required. Such processes are known to persons skilled in the art.

Expression of Affibodies

[0162] The construction of affibodies has been described elsewhere (Ronnlmark J, Gronlund H, Uhle' n, M., Nygren P.A°, Human immunoglobulin A (IgA)-specific ligands from combinatorial engineering of protein A, 2002, Eur. J. Biochem. 269, 2647-2655 .), including the construction of Affibody phage display libraries (Nord, K., Nilsson, J., Nilsson, B., Uhle' n, M. & Nygren, P.A, A combinatorial library of an a-helical bacterial receptor domain, 1995, Protein Eng. 8, 601-608. Nord, K., Gunneriusson, E., Ringdahl, J., Stahl, S., Uhle' n, M. & Nygren, P.A, Binding proteins selected from combinatorial libraries of an a-helical bacterial receptor domain, 1997, Nat. Biotechnol. 15, 772-777.)

[0163] The biosensor analyses to investigate the optimal Affibody variants using biosensor binding studies has also been described elsewhere (Ronnlmark J, Grnlund H, Uhle' n, M., Nygren P.A, Human immunoglobulin A (IgA)-specific ligands from combinatorial engineering of protein A, 2002, Eur. J. Biochem. 269, 2647-2655.).

Affinity Reagent Modifications

[0164] In one embodiment, anti-OGTA076 affinity reagents such as antibodies or fragments thereof are conjugated to a diagnostic moiety (such as a detectable label) or a therapeutic moiety. The antibodies can be used for diagnosis or to determine the efficacy of a given treatment regimen. Detection can be facilitated by coupling the antibody to a detectable substance (label). Examples of detectable substances include various enzymes, prosthetic groups, fluorescent materials, luminescent materials, bioluminescent materials, radioactive nuclides, positron emitting metals (for use in positron emission tomography), and nonradioactive paramagnetic metal ions. See generally U.S. Patent No. 4,741,900 for metal ions which can be conjugated to antibodies for use as diagnostics according to the present invention. Suitable enzymes include horseradish peroxidase, alkaline phosphatase, beta-galactosidase, or acetylcholinesterase; suitable prosthetic groups include streptavidin, avidin and biotin; suitable fluorescent materials include umbelliferon, fluorescein, fluorescein isothiocyanate, rhodamine, dichlorotriazinylamine fluorescein, dansyl chloride and phycoerythrin; suitable luminescent materials include luminol; suitable bioluminescent materials include luciferase, luciferin, and aequorin; and suitable radioactive nuclides include ^{125}I , ^{131}I , ^{111}In and ^{99}Tc . ^{68}Ga may also be employed.

[0165] Anti-OGTA076 antibodies or fragments thereof as well as other affinity reagents can be conjugated to a therapeutic agent or drug moiety to modify a given biological response. An exemplary therapeutic agent to which the affinity reagent may be conjugated is a cytotoxic moiety. The therapeutic agent or drug moiety is not to be construed as limited to classical chemical therapeutic agents. For example, the drug moiety may be a protein or polypeptide possessing a desired biological activity. Such proteins may include, for example, a toxin such as abrin, ricin A, pseudomonas exotoxin, or diphtheria toxin; a protein such as tumor necrosis factor, α -interferon, β -interferon, nerve growth factor, platelet derived growth factor, tissue plasminogen activator, a thrombotic agent or an anti-angiogenic agent, e.g. angiostatin or endostatin; or, a biological response modifier such as a lymphokine, interleukin-1 (IL-1), interleukin-2 (IL-2), interleukin-6 (IL-6), granulocyte macrophage colony stimulating factor (GM-CSF), granulocyte colony stimulating factor (G-CSF), nerve growth factor (NGF) or other growth factor.

[0166] Techniques for conjugating such therapeutic moiety to antibodies are well known, see, e.g. Arnon et al., "Monoclonal Antibodies For Immunotargeting Of Drugs In Cancer Therapy", in Monoclonal Antibodies And Cancer Therapy, Reisfeld et al. (eds.), pp. 243-56 (Alan R. Liss, Inc. 1985); Hellstrom et al., "Antibodies For Drug Delivery", in Controlled Drug Delivery (2nd Ed.), Robinson et al. (eds.), pp. 623-53 (Marcel Dekker, Inc. 1987); Thorpe, "Antibody Carriers Of Cytotoxic Agents In Cancer Therapy: A Review", in Monoclonal Antibodies '84: Biological And Clinical Applications, Pinchera et al. (eds.), pp. 475-506 (1985); "Analysis, Results, And Future Prospective Of The Therapeutic Use Of Radiolabeled Antibody In Cancer Therapy", in Monoclonal Antibodies For Cancer Detection And Therapy, Baldwin et al. (eds.), pp. 303-16 (Academic Press 1985), and Thorpe et al., "The Preparation And Cytotoxic Properties Of Antibody-Toxin Conjugates", Immunol. Rev., 62:119-58 (1982).

[0167] Alternatively, an antibody can be conjugated to a second antibody to form an antibody heteroconjugate as described by Segal in U.S. Patent No. 4,676,980.

[0168] An antibody with or without a therapeutic moiety conjugated to it can be used as a therapeutic that is administered alone or in combination with cytotoxic factor(s) and/or cytokine(s).

[0169] The invention also provides for fully human, or humanised antibodies that induce antibody-directed cell-mediated cytotoxicity (ADCC). A fully human antibody is one in which the protein sequences are encoded by naturally occurring human immunoglobulin sequences, either from isolated antibody-producing human B-lymphocytes, or from transgenic murine B-lymphocytes of mice in which the murine immunoglobulin coding chromosomal regions have been replaced by orthologous human sequences. Transgenic antibodies of the latter type include, but are not restricted to, HuMab (Medarex, Inc, CA) and Xenomouse (Abgenix Inc., CA). A humanised antibody is one in which the constant region of a non-human antibody molecule of appropriate antigen specificity, is replaced by the constant region of a human antibody, for example of the IgG subtype, with appropriate effector functions (Morrison et al., 1984, Proc. Natl. Acad. Sci. 81:851-855; Neuberger et al., 1984, Nature 312:604-608; Takeda et al., 1985, Nature 314:452-454). Appropriate effector functions include ADCC, which is a natural process by which fully-human antibodies or humanized antibodies, when bound to targets on the surface of cancer cells, switch on the cell killing properties of lymphocytes that are part of the normal immune system. These active lymphocytes, called Natural Killer (NK) cells, use a cytotoxic process to destroy living cells to which the antibodies are bound. ADCC activity may be detected and quantified by measuring

release of Europium (Eu3+) from Eu3+ labelled, living cells in the presence of an antigen-specific antibody and peripheral blood mononuclear cells extracted from an immunocompetent, living human subject. The ADCC process is described in detail in Janeway Jr. C.A. et al., *Immunobiology*, 5th ed., 2001, Garland Publishing, ISBN 0-8153-3642-X; Pier G.B. et al., *Immunology, Infection, and Immunity*, 2004, p246-5; Albanell J. et al., *Advances in Experimental Medicine and Biology*, 2003, 532:p2153-68 and Weng, W.-K. et al., *Journal of Clinical Oncology*, 2003, 21:p 3940-3947. Suitable methods for the detection and quantification of ADCC can be found in Blomberg et al., *Journal of Immunological Methods*. 1986, 86:p225-9; Blomberg et al., *Journal of Immunological Methods*. 1986, 21:92:p117-23 and Patel & Boyd, *Journal of Immunological Methods*. 1995, 184:p29-38.

[0170] ADCC typically involves activation of NK cells and is dependent on the recognition of antibody-coated cells by Fc receptors on the surface of the NK cell. The Fc receptors recognize the Fc (crystalline) portion of antibodies such as IgG, bound specifically to the surface of a target cell. The Fc receptor that triggers activation of the NK cell is called CD16 or Fc_YR_{IIIA}. Once the Fc_YR_{IIIA} receptor is bound to the IgG Fc, the NK cell releases cytokines such as IFN- γ , and cytotoxic granules containing perforin and granzymes that enter the target cell and promote cell death by triggering apoptosis.

[0171] The induction of antibody-dependent cellular cytotoxicity (ADCC) by an antibody can be enhanced by modifications that alter interactions between the antibody constant region (Fc) and various receptors that are present on the surface of cells of the immune system. Such modifications include the reduction or absence of alpha1,6-linked fucose moieties in the complex oligosaccharide chains that are normally added to the Fc of antibodies during natural or recombinant synthesis in mammalian cells. In one embodiment, non-fucosylated anti-OGTA076 affinity reagents such as antibodies or fragments thereof are produced for the purpose of enhancing their ability to induce the ADCC response.

[0172] Techniques for reducing or ablating alpha1,6-linked fucose moieties in the oligosaccharide chains of the Fc are well established. In one example, the recombinant antibody is synthesized in a cell line that is impaired in its ability to add fucose in an alpha 1,6 linkage to the innermost N-acetylglucosamine of the N-linked biantennary complex-type Fc oligosaccharides. Such cell lines include, but are not limited to, the rat hybridoma YB2/0, which expresses a reduced level of the alpha 1,6-fucosyltransferase gene, FUT8. For example, the antibody is synthesized in a cell line that is incapable of adding alpha 1,6-linked fucosyl moieties to complex oligosaccharide chains, due to the deletion of both copies of the FUT8 gene. Such cell lines include, but are not limited to, FUT8-/- CHO/DG44 cell lines. Techniques for synthesizing partially fucosylated, or non-fucosylated antibodies and affinity reagents are described in Shinkawa et al., *J. Biol. Chem.* 278:3466-34735 (2003); Yamane-Ohnuki et al., *Biotechnology and Bioengineering* 87: 614-22 (2004) and in WO00/61739 A1, WO02/31140 A1 and WO03/085107 A1. In a second example, the fucosylation of a recombinant antibody is reduced or abolished by synthesis in a cell line that has been genetically engineered to overexpress a glycoprotein-modifying glycosyl transferase at a level that maximizes the production of complex N-linked oligosaccharides carrying bisecting N-acetylglucosamine. For example, the antibody is synthesized in a Chinese Hamster Ovary cell line expressing the enzyme N-acetyl glucosamine transferase III (GnT III). Cell lines stably transfected with suitable glycoprotein-modifying glycosyl transferases, and methods of synthesizing antibodies using these cells are described in WO9954342.

[0173] A non-fucosylated antibody or affinity reagent can be used as a therapeutic that is administered alone or in combination with cytotoxic factor(s) and/or cytokine(s).

[0174] In a further modification, the amino acid sequences of the antibody Fc are altered in a way that enhances ADCC activation, without affecting ligand affinity. Examples of such modifications are described in Lazar et al., *Proceedings of the National Academy of Sciences* 2006, 103:p4005-4010; WO03074679 and WO2007039818. In these examples, substitution of amino acids in the antibody Fc, such as aspartate for serine at position 239, and isoleucine for glutamate at position 332, altered the binding affinity of an antibody for Fc receptors, leading to an increase in ADCC activation.

[0175] An antibody reagent with enhanced ADCC activation due to amino acid substitutions can be used as a therapeutic that is administered alone or in combination with cytotoxic factor(s) and/or cytokine(s).

Diagnosis of Colorectal Cancer, Kidney Cancer, Liver Cancer, Lung Cancer, Lymphoid Leukaemia (Particularly Chronic Lymphocytic Leukaemia), Ovarian Cancer or Pancreatic Cancer

[0176] In accordance with the present disclosure, test samples of colorectal, kidney, liver, lung, lymphoid, ovarian or pancreatic tissue, serum, plasma or urine obtained from a subject suspected of having or known to have colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer can be used for diagnosis or monitoring. In one example, a change in the abundance of OGTA076 in a test sample relative to a

control sample (from a subject or subjects free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer) or a previously determined reference range indicates the presence of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. In another example, the relative abundance of OGTA076 in a test sample compared to a control sample or a previously determined reference range indicates a subtype of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer (e.g. familial or sporadic colorectal cancer, fibrolamellar hepatocellular carcinoma, squamous cell lung carcinoma, T-cell chronic lymphocytic leukaemia, malignant papillary serous adenocarcinoma or endocrine tumours of the pancreas). In yet another example, the relative abundance of OGTA076 in a test sample relative to a control sample or a previously determined reference range indicates the degree or severity of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer (e.g. the likelihood for metastasis). In any of the aforesaid methods, detection of OGTA076 may optionally be combined with detection of one or more of additional biomarkers for colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Any suitable method in the art can be employed to measure the level of OGTA076, including but not limited to the Preferred Technologies described herein, kinase assays, immunoassays to detect and/or visualize the OGTA076 (e.g. Western blot, immunoprecipitation followed by sodium dodecyl sulfate polyacrylamide gel electrophoresis, immunocytochemistry, etc.). In a further example, a change in the abundance of mRNA encoding OGTA076 in a test sample relative to a control sample or a previously determined reference range indicates the presence of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Any suitable hybridization assay can be used to detect OGTA076 expression by detecting and/or visualizing mRNA encoding the OGTA076 (e.g. Northern assays, dot blots, *in situ* hybridization, etc.).

[0177] In another example of the disclosure, labeled antibodies (or other affinity reagents), derivatives and analogs thereof, which specifically bind to OGTA076 can be used for diagnostic purposes to detect, diagnose, or monitor colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Preferably, colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer is detected in an animal, more preferably in a mammal and most preferably in a human.

Screening Assays

[0178] The disclosure provides methods for identifying agents (e.g. candidate compounds or test compounds) that bind to OGTA076 or have a stimulatory or inhibitory effect on the expression or activity of OGTA076. The disclosure also provides methods of identifying agents, candidate compounds or test compounds that bind to an OGTA076-related polypeptide or an OGTA076 fusion protein or have a stimulatory or inhibitory effect on the expression or activity of an OGTA076-related polypeptide or an OGTA076 fusion protein. Examples of agents, candidate compounds or test compounds include, but are not limited to, nucleic acids (e.g. DNA and RNA), carbohydrates, lipids, proteins, peptides, peptidomimetics, small molecules and other drugs. Agents can be obtained using any of the numerous approaches in combinatorial library methods known in the art, including:

biological libraries; spatially addressable parallel solid phase or solution phase libraries; synthetic library methods requiring deconvolution; the "one-bead one-compound" library method; and synthetic library methods using affinity chromatography selection. The biological library approach is limited to peptide libraries, while the other four approaches are applicable to peptide, non-peptide oligomer or small molecule libraries of compounds (Lam, 1997, *Anticancer Drug Des.* 12:145; U.S. Patent No. 5,738,996; and U.S. Patent No. 5,807,683).

[0179] Examples of methods for the synthesis of molecular libraries can be found in the art, for example in: DeWitt et al., 1993, *Proc. Natl. Acad. Sci. USA* 90:6909; Erb et al., 1994, *Proc. Natl. Acad. Sci. USA* 91:11422; Zuckermann et al., 1994, *J. Med. Chem.* 37:2678; Cho et al., 1993, *Science* 261:1303; Carrell et al., 1994, *Angew. Chem. Int. Ed. Engl.* 33:2059; Carell et al., 1994, *Angew. Chem. Int. Ed. Engl.* 33:2061; and Gallop et al., 1994, *J. Med. Chem.* 37:1233.

[0180] Libraries of compounds may be presented, e.g. presented in solution (e.g. Houghten, 1992, *Bio/Techniques* 13:412-421), or on beads (Lam, 1991, *Nature* 354:82-84), chips (Fodor, 1993, *Nature* 364:555-556), bacteria (U.S. Patent No. 5,223,409), spores (Patent Nos. 5,571,698; 5,403,484; and 5,223,409), plasmids (Cull et al., 1992, *Proc. Natl. Acad. Sci. USA* 89:1865-1869) or phage (Scott and Smith, 1990, *Science* 249:386-390; Devlin, 1990, *Science* 249:404-406; Cwirla et al., 1990, *Proc. Natl. Acad.*

Sci. USA 87:6378-6382; and Felici, 1991, J. Mol. Biol. 222:301-310).

[0181] In one example, agents that interact with (i.e. bind to) OGTA076, an OGTA076 fragment (e.g. a functionally active fragment), an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein are identified in a cell-based assay system. In accordance with this example, cells expressing OGTA076, a fragment of an OGTA076, an OGTA076-related polypeptide, a fragment of the OGTA076-related polypeptide, or an OGTA076 fusion protein are contacted with a candidate compound or a control compound and the ability of the candidate compound to interact with the OGTA076 is determined. If desired, this assay may be used to screen a plurality (e.g. a library) of candidate compounds. The cell, for example, can be of prokaryotic origin (e.g. *E. coli*) or eukaryotic origin (e.g. yeast or mammalian). Further, the cells can express OGTA076, a fragment of OGTA076, an OGTA076-related polypeptide, a fragment of the OGTA076-related polypeptide, or an OGTA076 fusion protein endogenously or be genetically engineered to express OGTA076, a fragment of OGTA076, an OGTA076-related polypeptide, a fragment of the OGTA076-related polypeptide, or an OGTA076 fusion protein. In certain instances, OGTA076, a fragment of OGTA076, an OGTA076-related polypeptide, a fragment of the OGTA076-related polypeptide, or an OGTA076 fusion protein or the candidate compound is labeled, for example with a radioactive label (such as ^{32}P , ^{35}S , and ^{125}I) or a fluorescent label (such as fluorescein isothiocyanate, rhodamine, phycoerythrin, phycoerythrin, allophycocyanin, o-phthaldehyde or fluorescamine) to enable detection of an interaction between OGTA076 and a candidate compound. The ability of the candidate compound to interact directly or indirectly with OGTA076, a fragment of an OGTA076, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein can be determined by methods known to those of skill in the art. For example, the interaction between a candidate compound and OGTA076, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein can be determined by flow cytometry, a scintillation assay, immunoprecipitation or western blot analysis.

[0182] In another example, agents that interact with (i.e. bind to) OGTA076, an OGTA076 fragment (e.g. a functionally active fragment), an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein are identified in a cell-free assay system. In accordance with this example, a native or recombinant OGTA076 or fragment thereof, or a native or recombinant OGTA076-related polypeptide or fragment thereof, or an OGTA076-fusion protein or fragment thereof, is contacted with a candidate compound or a control compound and the ability of the candidate compound to interact with OGTA076 or OGTA076-related polypeptide, or OGTA076 fusion protein is determined. If desired, this assay may be used to screen a plurality (e.g. a library) of candidate compounds. For example, OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076-fusion protein is first immobilized, by, for example, contacting OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein with an immobilized antibody (or other affinity reagent) which specifically recognizes and binds it, or by contacting a purified preparation of OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein with a surface designed to bind proteins. OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein may be partially or completely purified (e.g. partially or completely free of other polypeptides) or part of a cell lysate. Further, OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, or a fragment of an OGTA076-related polypeptide may be a fusion protein comprising OGTA076 or a biologically active portion thereof, or OGTA076-related polypeptide and a domain such as glutathione-S-transferase. Alternatively, OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide or an OGTA076 fusion protein can be biotinylated using techniques well known to those of skill in the art (e.g. biotinylation kit, Pierce Chemicals; Rockford, IL). The ability of the candidate compound to interact with OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein can be determined by methods known to those of skill in the art.

[0183] In another example, a cell-based assay system is used to identify agents that bind to or modulate the activity of a protein, such as an enzyme, or a biologically active portion thereof, which is responsible for the production or degradation of OGTA076 or is responsible for the post-translational modification of OGTA076. In a primary screen, a plurality (e.g. a library) of compounds are contacted with cells that naturally or recombinantly express: (i) OGTA076, an isoform of OGTA076, an OGTA076 homolog, an OGTA076-related polypeptide, an OGTA076 fusion protein, or a biologically active fragment of any of the foregoing; and (ii) a protein that is responsible for processing of OGTA076, an OGTA076 isoform, an OGTA076 homolog, an OGTA076-related polypeptide, an OGTA076 fusion protein, or a fragment in order to identify compounds that modulate the production, degradation, or post-translational modification of OGTA076, an OGTA076 isoform, an OGTA076 homolog, an OGTA076-related polypeptide, an OGTA076 fusion protein or fragment. If desired, compounds identified in the primary screen can then be assayed in a secondary screen against cells naturally or recombinantly expressing OGTA076. The ability of the candidate compound to modulate the production, degradation or post-translational modification of OGTA076, isoform, homolog, OGTA076-related polypeptide, or OGTA076 fusion protein can be determined by methods known to those of skill in the art, including without limitation, flow cytometry, a scintillation assay, immunoprecipitation and western blot analysis.

[0184] In another example, agents that competitively interact with (i.e. bind to) OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein are identified in a competitive binding assay. In accordance with this example, cells expressing OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein are contacted with a candidate compound and a compound known to interact with OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide or an OGTA076 fusion protein; the ability of the candidate compound to preferentially interact with OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein is then determined. Alternatively, agents that preferentially interact with (i.e. bind to) OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide or fragment of an OGTA076-related polypeptide are identified in a cell-free assay system by contacting OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein with a candidate compound and a compound known to interact with OGTA076, an OGTA076-related polypeptide or an OGTA076 fusion protein. As stated above, the ability of the candidate compound to interact with OGTA076, an OGTA076 fragment, an OGTA076-related polypeptide, a fragment of an OGTA076-related polypeptide, or an OGTA076 fusion protein can be determined by methods known to those of skill in the art. These assays, whether cell-based or cell-free, can be used to screen a plurality (e.g. a library) of candidate compounds.

[0185] In another example, agents that modulate (i.e. upregulate or downregulate) the expression or activity of OGTA076 or an OGTA076-related polypeptide are identified by contacting cells (e.g. cells of prokaryotic origin or eukaryotic origin) expressing OGTA076 or an OGTA076-related polypeptide with a candidate compound or a control compound (e.g. phosphate buffered saline (PBS)) and determining the expression of OGTA076, OGTA076-related polypeptide, or OGTA076 fusion protein, mRNA encoding OGTA076, or mRNA encoding the OGTA076-related polypeptide. The level of expression of OGTA076, OGTA076-related polypeptide, mRNA encoding OGTA076, or mRNA encoding the OGTA076-related polypeptide in the presence of the candidate compound is compared to the level of expression of OGTA076, OGTA076-related polypeptide, mRNA encoding OGTA076, or mRNA encoding the OGTA076-related polypeptide in the absence of the candidate compound (e.g. in the presence of a control compound). The candidate compound can then be identified as a modulator of the expression of OGTA076, or the OGTA076-related polypeptide based on this comparison. For example, when expression of OGTA076 or mRNA is significantly greater in the presence of the candidate compound than in its absence, the candidate compound is identified as a stimulator of expression of OGTA076 or mRNA. Alternatively, when expression of OGTA076 or mRNA is significantly less in the presence of the candidate compound than in its absence, the candidate compound is identified as an inhibitor of the expression of OGTA076 or mRNA. The level of expression of OGTA076 or the mRNA that encodes it can be determined by methods known to those of skill in the art. For example, mRNA expression can be assessed by Northern blot analysis or RT-PCR, and protein levels can be assessed by western blot analysis.

[0186] In another example, agents that modulate the activity of OGTA076 or an OGTA076-related polypeptide are identified by contacting a preparation containing OGTA076 or OGTA076-related polypeptide or cells (e.g. prokaryotic or eukaryotic cells) expressing OGTA076 or OGTA076-related polypeptide with a test compound or a control compound and determining the ability of the test compound to modulate (e.g. stimulate or inhibit) the activity of OGTA076 or OGTA076-related polypeptide. The activity of OGTA076 or an OGTA076-related polypeptide can be assessed by detecting induction of a cellular signal transduction pathway of OGTA076 or OGTA076-related polypeptide (e.g. intracellular Ca^{2+} , diacylglycerol, IP3, etc.), detecting catalytic or enzymatic activity of the target on a suitable substrate, detecting the induction of a reporter gene (e.g. a regulatory element that is responsive to OGTA076 or an OGTA076-related polypeptide and is operably linked to a nucleic acid encoding a detectable marker, e.g. luciferase), or detecting a cellular response, for example, cellular differentiation, or cell proliferation. Based on the present description, techniques known to those of skill in the art can be used for measuring these activities (see, e.g. U.S. Patent No. 5,401,639). The candidate compound can then be identified as a modulator of the activity of OGTA076 or an OGTA076-related polypeptide by comparing the effects of the candidate compound to the control compound. Suitable control compounds include phosphate buffered saline (PBS) and normal saline (NS).

[0187] In another example, agents that modulate (i.e. upregulate or downregulate) the expression, activity or both the expression and activity of OGTA076 or an OGTA076-related polypeptide are identified in an animal model. Examples of suitable animals include, but are not limited to, mice, rats, rabbits, monkeys, guinea pigs, dogs and cats. Preferably, the animal used represent a model of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer (e.g. xenografts of human colorectal cancer cell lines such as MDA-MB-345 in oestrogen-deprived SCID mice, Eccles et al. 1994 Cell Biophysics 24/25, 279; xenografts of renal cell cancer cell lines such as LABAZ1 in immune compromised mice, Zisman et al, Cancer Research 63, 4952-4959, August 15, 2003; xenografts of hepatocellular carcinoma cell lines such as MHCC97 in nude mice, Tian et al., Br J Cancer 1999 Nov;81(5):814-21; xenografts of non small cell lung cancer cell lines such as A549 and H460 and xenografts of small cell lung cancer cell lines such as NCI-

H345; xenografts of chronic lymphocytic leukaemia cell lines such as WSU-CLL in SCID mice, Mohammad et al, Leukaemia. 1996 Jan;10(1): 130-7; xenografts of ovarian cancer cell lines such as IGROV1 in nude mice, Benard et al, Cancer Res. 1985 Oct;45(10):4970-9 or xenografts of pancreatic cancer cell lines such as MIA PaCa-2 in nude mice, Marincola et al., J Surg Res 1989 Dec;47(6):520-9). These can be utilized to test compounds that modulate OGTA076 levels, since the pathology exhibited in these models is similar to that of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. In accordance with this example, the test compound or a control compound is administered (e.g. orally, rectally or parenterally such as intraperitoneally or intravenously) to a suitable animal and the effect on the expression, activity or both expression and activity of OGTA076 or OGTA076-related polypeptide is determined. Changes in the expression of OGTA076 or an OGTA076-related polypeptide can be assessed by the methods outlined above.

[0188] In yet another example, OGTA076 or an OGTA076-related polypeptide is used as a "bait protein" in a two-hybrid assay or three hybrid assay to identify other proteins that bind to or interact with OGTA076 or an OGTA076-related polypeptide (see, e.g. U.S. Patent No. 5,283,317; Zervos et al. (1993) Cell 72:223-232; Madura et al. (1993) J. Biol. Chem. 268:12046-12054; Bartel et al. (1993) Bio/Techniques 14:920-924; Iwabuchi et al. (1993) Oncogene 8:1693-1696; and PCT Publication No. WO 94/10300). As those skilled in the art will appreciate, such binding proteins are also likely to be involved in the propagation of signals by OGTA076 as, for example, upstream or downstream elements of a signaling pathway involving OGTA076.

[0189] This disclosure further provides novel agents identified by the above-described screening assays and uses thereof for treatments as described herein. In addition, the disclosure also provides the use of an agent which interacts with, or modulates the activity of, OGTA076 in the manufacture of a medicament for the treatment of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.

Therapeutic Use of OGTA076

[0190] The disclosure provides for treatment or prevention of various diseases and disorders by administration of a therapeutic compound. Such compounds include but are not limited to: OGTA076, OGTA076 analogs, OGTA076-related polypeptides and derivatives (including fragments) thereof; antibodies (or other affinity reagents) to the foregoing; nucleic acids encoding OGTA076, OGTA076 analogs, OGTA076-related polypeptides and fragments thereof; antisense nucleic acids to a gene encoding OGTA076 or an OGTA076-related polypeptide; and modulator (e.g. agonists and antagonists) of a gene encoding OGTA076 or an OGTA076-related polypeptide. An important feature of the present disclosure is the identification of genes encoding OGTA076 involved in colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer can be treated (e.g. to ameliorate symptoms or to retard onset or progression) or prevented by administration of a therapeutic compound that reduces function or expression of OGTA076 in the serum or tissue of subjects having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Thus provided herein are one or more of the aforementioned moieties for use in treatment or prophylaxis, for example in the treatment or prophylaxis of cancer, such as a cancer defined herein. The invention also extends to use of said moieties in the manufacture of a medicament for the treatment or prophylaxis of cancer, for example a cancer described herein and uses in a method of treating cancer, for example a cancer described herein.

[0191] In one embodiment, one or more antibodies (or other affinity reagents) each specifically binding to OGTA076 are administered alone or in combination with one or more additional therapeutic compounds or treatments.

[0192] In a further embodiment, a biological product such as an antibody (or other affinity reagent) is allogeneic to the subject to which it is administered. In another example, a human OGTA076 or a human OGTA076-related polypeptide, a nucleotide sequence encoding a human OGTA076 or a human OGTA076-related polypeptide, or an antibody (or other affinity reagent) to a human OGTA076 or a human OGTA076-related polypeptide, is administered to a human subject for therapy (e.g. to ameliorate symptoms or to retard onset or progression) or prophylaxis.

[0193] Without being limited by theory, it is conceived that the therapeutic activity of antibodies (or other affinity reagents) which specifically bind to OGTA076 may be achieved through the phenomenon of Antibody -Dependent Cell-mediated Cytotoxicity (ADCC) (see e.g. Janeway Jr. C.A. et al., Immunobiology, 5th ed., 2001, Garland Publishing, ISBN 0-8153-3642-X; Pier G.B. et al., Immunology, Infection, and Immunity, 2004, p246-5; Albanell J. et al., Advances in Experimental Medicine and Biology, 2003, 532:p2153-68 and Weng, W.-K. et al., Journal of Clinical Oncology, 2003, 21:p 3940-3947).

Treatment and Prevention of Colorectal Cancer, Kidney Cancer, Liver Cancer, Lung Cancer, Lymphoid Leukaemia (Particularly Chronic Lymphocytic Leukaemia), Ovarian Cancer or Pancreatic Cancer

[0194] Colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer is treated or prevented by administration to a subject suspected of having or known to have colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or to be at risk of developing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer of a compound that modulates (i.e. increases or decreases) the level or activity (i.e. function) of OGTA076 that is differentially present in the serum or tissue of subjects having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer compared with serum or tissue of subjects free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer. In one example, colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer is treated or prevented by administering to a subject suspected of having or known to have colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or to be at risk of developing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer a compound that upregulates (i.e. increases) the level or activity (i.e. function) of OGTA076 that are decreased in the serum or tissue of subjects having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Examples of such a compound include, but are not limited to, OGTA076 antisense oligonucleotides, ribozymes, antibodies (or other affinity reagents) directed against OGTA076, and compounds that inhibit the enzymatic activity of OGTA076. Other useful compounds e.g. OGTA076 antagonists and small molecule OGTA076 antagonists, can be identified using *in vitro* assays.

[0195] Colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer is also treated or prevented by administration to a subject suspected of having or known to have colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or to be at risk of developing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer of a compound that downregulates the level or activity (i.e. function) of OGTA076 that are increased in the serum or tissue of subjects having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Examples of such a compound include but are not limited to: OGTA076, OGTA076 fragments and OGTA076-related polypeptides; nucleic acids encoding OGTA076, an OGTA076 fragment and an OGTA076-related polypeptide (e.g. for use in gene therapy); and, for those OGTA076 or OGTA076-related polypeptides with enzymatic activity, compounds or molecules known to modulate that enzymatic activity. Other compounds that can be used, e.g. OGTA076 agonists, can be identified using *in vitro* assays.

[0196] In one example, therapy or prophylaxis is tailored to the needs of an individual subject. Thus, in specific examples, compounds that promote the level or function of OGTA076 are therapeutically or prophylactically administered to a subject suspected of having or known to have colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer, in whom the levels or functions of OGTA076 are absent or are decreased relative to a control or normal reference range. In further examples, compounds that promote the level or function of OGTA076 are therapeutically or prophylactically administered to a subject suspected of having or known to have colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in whom the levels or functions of OGTA076 are increased relative to a control or to a reference range. In further examples, compounds that decrease the level or function of OGTA076 are therapeutically or prophylactically administered to a subject suspected of having or known to have colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in whom the levels or functions of OGTA076 are increased relative to a control or to a reference range. In further examples, compounds that decrease the level or function of OGTA076 are therapeutically or prophylactically administered to a subject suspected of having or known to have colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in whom the levels or functions of OGTA076 are decreased relative to a control or to a reference range. The change in OGTA076 function or level due to the administration of such compounds can be readily detected, e.g. by obtaining a sample (e.g. blood or urine) and assaying *in vitro* the levels or activities of OGTA076, or the levels of mRNAs encoding OGTA076, or any combination of the foregoing. Such assays can be performed before and after the administration of the compound as described herein.

[0197] The compounds of the disclosure include but are not limited to any compound, e.g. a small organic molecule, protein, peptide, antibody (or other affinity reagent), nucleic acid, etc. that restores the OGTA076 profile towards normal. The compounds of the disclosure may be given in combination with any other chemotherapy drugs.

Vaccine Therapy

[0198] Also described herein is an immunogenic composition or a pharmaceutical composition, suitably a vaccine composition, comprising OGTA076 or an epitope containing fragment thereof, or nucleic acid encoding OGTA076 or a fragment thereof optionally together with an immunostimulant. The invention also extends to said composition(s) for use in treatment, for example for the treatment or prophylaxis of cancer, such as a cancer described herein. The invention further extends to use of said compositions for the manufacture of a medicament for the treatment or prophylaxis of cancer, for example a cancer described herein and the use of the composition(s) in treating same.

[0199] There is also provided a method of raising an immune response which comprises administering to a subject such compositions and a method for treating or preventing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer which comprises administering to a subject in need thereof a therapeutically effective amount of such compositions and such compositions for use in preventing or treating colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.

[0200] Thus, OGTA076 may be useful as antigenic material, and may be used in the production of vaccines for treatment or prophylaxis of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Such material can be "antigenic" and/or "immunogenic". Generally, "antigenic" is taken to mean that the protein is capable of being used to raise antibodies (or other affinity reagents) or indeed is capable of inducing an antibody response in a subject or experimental animal. "Immunogenic" is taken to mean that the protein is capable of eliciting an immune response, for example a protective immune response in a subject or experimental animal. Thus, in the latter case, the protein may be capable of not only generating an antibody response but, in addition, non-antibody based immune responses. "Immunogenic" also embraces whether the protein may elicit an immune-like response in an in-vitro setting e.g. a T-cell proliferation assay.

[0201] The skilled person will appreciate that homologues or derivatives of OGTA076 will also find use as antigenic/immunogenic material. Thus, for instance proteins which include one or more additions, deletions, substitutions or the like are encompassed by the present disclosure. In addition, it may be possible to replace one amino acid with another of similar "type". For instance, replacing one hydrophobic amino acid with another. One can use a program such as the CLUSTAL program to compare amino acid sequences. This program compares amino acid sequences and finds the optimal alignment by inserting spaces in either sequence as appropriate. It is possible to calculate amino acid identity or similarity (identity plus conservation of amino acid type) for an optimal alignment. A program like BLASTx will align the longest stretch of similar sequences and assign a value to the fit. It is thus possible to obtain a comparison where several regions of similarity are found, each having a different score. Both types of analysis are contemplated in the present disclosure.

[0202] In the case of homologues and derivatives, the degree of identity with a protein as described herein is less important than that the homologue or derivative should retain its antigenicity and/or immunogenicity. However, suitably, homologues or derivatives having at least 60% similarity (as discussed above) with the proteins or polypeptides described herein are provided. In one example, homologues or derivatives having at least 70% similarity, for example at least 80% similarity are provided, such as homologues or derivatives having at least 90% or even 95% similarity. Suitably, homologues or derivatives have at least 60% sequence identity with the proteins or polypeptides described herein. In one example homologues or derivatives have at least 70% identity, more preferably at least 80% identity. Most preferably, homologues or derivatives have at least 90% or even 95% identity.

[0203] In an alternative approach, the homologues or derivatives could be fusion proteins, incorporating moieties which render purification easier, for example by effectively tagging the desired protein or polypeptide. It may be necessary to remove the "tag" or it may be the case that the fusion protein itself retains sufficient antigenicity to be useful.

[0204] It is well known that it is possible to screen an antigenic protein or polypeptide to identify epitopic regions, i.e. those regions which are responsible for the protein or polypeptide's antigenicity or immunogenicity. Methods well known to the skilled

person can be used to test fragments and/or homologues and/or derivatives for antigenicity. Thus, the fragments of the present disclosure should include one or more such epitopic regions or be sufficiently similar to such regions to retain their antigenic/immunogenic properties. Thus, for fragments according to the present disclosure the degree of identity is perhaps irrelevant, since they may be 100% identical to a particular part of a protein or polypeptide, homologue or derivative as described herein. The key issue, once again, is that the fragment retains the antigenic/immunogenic properties of the protein from which it is derived.

[0205] What is important for homologues, derivatives and fragments is that they possess at least a degree of the antigenicity/immunogenicity of the protein or polypeptide from which they are derived. Thus, in an additional example, there is provided antigenic/or immunogenic fragments of OGTA076, or of homologues or derivatives thereof.

[0206] OGTA076, or antigenic fragments thereof, can be provided alone, as a purified or isolated preparation. They may be provided as part of a mixture with one or more other proteins of the disclosure, or antigenic fragments thereof. In a further example, therefore, the disclosure provides an antigen composition comprising OGTA076 and/or one or more antigenic fragments thereof. Such a composition can be used for the detection and/or diagnosis of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.

[0207] Vaccine compositions according to the disclosure may be either a prophylactic or therapeutic vaccine composition.

[0208] The vaccine compositions of the disclosure can include one or more adjuvants (immunostimulants). Examples well-known in the art include inorganic gels, such as aluminium hydroxide, and water-in-oil emulsions, such as incomplete Freund's adjuvant. Other useful adjuvants will be well known to the skilled person.

[0209] Suitable adjuvants for use in vaccine compositions for the treatment of cancer include: 3De-O-acylated monophosphoryl lipid A (known as 3D-MPL or simply MPL see WO92/116556), a saponin, for example QS21 or QS7, and TLR4 agonists such as a CpG containing molecule, for example as disclosed in WO95/26204.

[0210] The adjuvants employed may be a combination of components, for example MPL and QS21 or MPL, QS21 and a CpG containing moiety.

[0211] Adjuvants may be formulated as oil-in-water emulsions or liposomal formulations.

[0212] Such preparations may include other vehicles.

[0213] In another example, a preparation of oligonucleotides comprising 10 or more consecutive nucleotides complementary to a nucleotide sequence encoding OGTA076 or an OGTA076 peptide fragments is used as vaccines for the treatment of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Such preparations may include adjuvants or other vehicles.

Inhibition of OGTA076 to Treat Colorectal Cancer, Kidney Cancer, Liver Cancer, Lung Cancer, Lymphoid Leukaemia (Particularly Chronic Lymphocytic Leukaemia), Ovarian Cancer or Pancreatic Cancer

[0214] In one example, colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer is treated or prevented by administration of a compound that antagonizes (inhibits) the level(s) and/or function(s) of OGTA076 which are elevated in the serum or tissue of subjects having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer as compared with serum or tissue of subjects free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer.

[0215] Compounds useful for this purpose include but are not limited to anti-OGTA076 antibodies (or other affinity reagents, and fragments and derivatives containing the binding region thereof), OGTA076 antisense or ribozyme nucleic acids, and nucleic acids encoding dysfunctional OGTA076 that are used to "knockout" endogenous OGTA076 function by homologous recombination (see, e.g. Capecchi, 1989, Science 244:1288-1292). Other compounds that inhibit OGTA076 function can be identified by use of known *in vitro* assays, e.g. assays for the ability of a test compound to inhibit binding of OGTA076 to another protein or a binding partner, or to inhibit a known OGTA076 function. Preferably such inhibition is assayed *in vitro* or in cell culture, but genetic assays may also be employed. The Preferred Technologies can also be used to detect levels of OGTA076

before and after the administration of the compound. Preferably, suitable *in vitro* or *in vivo* assays are utilized to determine the effect of a specific compound and whether its administration is indicated for treatment of the affected tissue, as described in more detail below.

[0216] In a specific example, a compound that inhibits OGTA076 function (activity) is administered therapeutically or prophylactically to a subject in whom an increased serum or tissue level or functional activity of OGTA076 (e.g. greater than the normal level or desired level) is detected as compared with serum or tissue of subjects free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer or a predetermined reference range. Methods standard in the art can be employed to measure the increase in OGTA076 level or function, as outlined above. Preferred OGTA076 inhibitor compositions include small molecules, i.e. molecules of 1000 daltons or less. Such small molecules can be identified by the screening methods described herein.

Assays for Therapeutic or Prophylactic Compounds

[0217] The present disclosure also provides assays for use in drug discovery in order to identify or verify the efficacy of compounds for treatment or prevention of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.

[0218] Thus there is provided a method of screening for compounds that modulate the activity of OGTA076, the method comprising: (a) contacting OGTA076 or a biologically active portion thereof with a candidate compound; and (b) determining whether activity of OGTA076 is thereby modulated. Such a process may comprise (a) contacting OGTA076 or a biologically active portion thereof with a candidate compound in a sample; and (b) comparing the activity of OGTA076 or a biologically active portion thereof in said sample after contact with said candidate compound with the activity of OGTA076 or a biologically active portion thereof in said sample before contact with said candidate compound, or with a reference level of activity.

[0219] The method of screening may be a method of screening for compounds that inhibit activity of OGTA076.

[0220] OGTA076 or a biologically active portion thereof may, for example be expressed on or by a cell. OGTA076 or a biologically active portion thereof may, for example, be isolated from cells which express it. OGTA076 or a biologically active portion thereof may, for example, be immobilised onto a solid phase.

[0221] There is also provided a method of screening for compounds that modulate the expression of OGTA076 or nucleic acid encoding OGTA076, the method comprising: (a) contacting cells expressing OGTA076 or nucleic acid encoding OGTA076 with a candidate compound; and (b) determining whether expression of OGTA076 or nucleic acid encoding OGTA076 is thereby modulated. Such a process may comprise (a) contacting cells expressing OGTA076 or nucleic acid encoding OGTA076 with a candidate compound in a sample; and (b) comparing the expression of OGTA076 or nucleic acid encoding OGTA076 by cells in said sample after contact with said candidate compound with the expression of OGTA076 or nucleic acid encoding OGTA076 of cells in said sample before contact with said candidate compound, or with a reference level of expression.

[0222] The method may be a method of screening for compounds that inhibit expression of OGTA076 or nucleic acid encoding OGTA076.

[0223] Other examples of the disclosure include: a compound obtainable by an aforementioned screening method, a compound which modulates the activity or expression of OGTA076 or nucleic acid encoding OGTA076, for example a compound which inhibits the activity or expression of OGTA076 or nucleic acid encoding OGTA076.

[0224] Such a compound is provided for use in treating or preventing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. There is also provided a method for treating or preventing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer which comprises administering to a subject in need thereof a therapeutically effective amount of such a compound.

[0225] Test compounds can be assayed for their ability to restore OGTA076 levels in a subject having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer towards levels found in subjects free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer or to produce similar changes in experimental

animal models of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Compounds able to restore OGTA076 levels in a subject having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer towards levels found in subjects free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer or to produce similar changes in experimental animal models of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer can be used as lead compounds for further drug discovery, or used therapeutically. OGTA076 expression can be assayed by the Preferred Technologies, immunoassays, gel electrophoresis followed by visualization, detection of OGTA076 activity, or any other method taught herein or known to those skilled in the art. Such assays can be used to screen candidate drugs, in clinical monitoring or in drug development, where abundance of OGTA076 can serve as a surrogate marker for clinical disease.

[0226] In various specific examples, *in vitro* assays can be carried out with cells representative of cell types involved in a subject's disorder, to determine if a compound has a desired effect upon such cell types.

[0227] Compounds for use in therapy can be tested in suitable animal model systems prior to testing in humans, including but not limited to rats, mice, chicken, cows, monkeys, rabbits, etc. For *in vivo* testing, prior to administration to humans, any animal model system known in the art may be used. Examples of animal models of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer include, but are not limited to xenografts of human colorectal cancer cell lines such as MDA-MB-345 in oestrogen-deprived SCID mice, Eccles et al. 1994 Cell Biophysics 24/25, 279; xenografts of renal cell cancer cell lines such as LABAZ1 in immune compromised mice, Zisman et al, Cancer Research 63, 4952-4959, August 15, 2003; xenografts of hepatocellular carcinoma cell lines such as MHCC97 in nude mice, Tian et al., Br J 5 Cancer 1999 Nov;81(5):814-21; xenografts of non small cell lung cancer cell lines such as A549 and H460 and xenografts of small cell lung cancer cell lines such as NCI-H345; xenografts of chronic lymphocytic leukaemia cell lines such as WSU-CLL in SCID mice, Mohammad et al, Leukaemia. 1996 Jan; 10(1):130-7; xenografts of ovarian cancer cell lines such as IGROV1 in nude mice, Benard et al, Cancer Res. 1985 Oct;45(10):4970-9 or xenografts of pancreatic cancer cell lines such as MIA PaCa-2 in nude mice, Marincola et al., J Surg Res 1989 Dec;47(6):520-9. These can be utilized to test compounds that modulate OGTA076 levels, since the pathology exhibited in these models is similar to that of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. It is also apparent to the skilled artisan that based upon the present disclosure, transgenic animals can be produced with "knock-out" mutations of the gene or genes encoding OGTA076. A "knock-out" mutation of a gene is a mutation that causes the mutated gene to not be expressed, or expressed in an aberrant form or at a low level, such that the activity associated with the gene product is nearly or entirely absent. Preferably, the transgenic animal is a mammal; more preferably, the transgenic animal is a mouse.

[0228] In one example, test compounds that modulate the expression of OGTA076 are identified in non-human animals (e.g. mice, rats, monkeys, rabbits, and guinea pigs), for example non-human animal models for colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer, expressing OGTA076. In accordance with this example, a test compound or a control compound is administered to the animals, and the effect of the test compound on expression of OGTA076 is determined. A test compound that alters the expression of OGTA076 can be identified by comparing the level of OGTA076 (or mRNA encoding the same) in an animal or group of animals treated with a test compound with the level of OGTA076 or mRNA in an animal or group of animals treated with a control compound. Techniques known to those of skill in the art can be used to determine the mRNA and protein levels, for example, *in situ* hybridization. The animals may or may not be sacrificed to assay the effects of a test compound.

[0229] In another example, test compounds that modulate the activity of OGTA076 or a biologically active portion thereof are identified in non-human animals (e.g. mice, rats, monkeys, rabbits, and guinea pigs), for example non-human animal models for colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer, expressing OGTA076. In accordance with this example, a test compound or a control compound is administered to the animals, and the effect of a test compound on the activity of OGTA076 is determined. A test compound that alters the activity of OGTA076 can be identified by assaying animals treated with a control compound and animals treated with the test compound. The activity of OGTA076 can be assessed by detecting induction of a cellular second messenger of OGTA076 (e.g. intracellular Ca^{2+} , diacylglycerol, IP3, etc.), detecting catalytic or enzymatic activity of OGTA076 or binding partner thereof, detecting the induction of a reporter gene (e.g. a regulatory element that is responsive to OGTA076 operably linked to a nucleic acid encoding a detectable marker, such as luciferase or green fluorescent protein), or detecting a cellular response (e.g. cellular differentiation or cell proliferation). Techniques known to those of skill in the art can be utilized to detect changes in the activity of OGTA076 (see, e.g. U.S. Patent No. 5,401,639).

[0230] In yet another example, test compounds that modulate the level or expression of OGTA076 are identified in human subjects having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer, for example those having severe colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. In accordance with this example, a test compound or a control compound is administered to the human subject, and the effect of a test compound on OGTA076 expression is determined by analyzing the expression of OGTA076 or the mRNA encoding the same in a biological sample (e.g. serum, plasma, or urine). A test compound that alters the expression of OGTA076 can be identified by comparing the level of OGTA076 or mRNA encoding the same in a subject or group of subjects treated with a control compound to that in a subject or group of subjects treated with a test compound. Alternatively, alterations in the expression of OGTA076 can be identified by comparing the level of OGTA076 or mRNA encoding the same in a subject or group of subjects before and after the administration of a test compound. Techniques known to those of skill in the art can be used to obtain the biological sample and analyze the mRNA or protein expression. For example, the Preferred Technologies described herein can be used to assess changes in the level of OGTA076.

[0231] In another example, test compounds that modulate the activity of OGTA076 are identified in human subjects having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer (preferably those with severe colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer). In this example, a test compound or a control compound is administered to the human subject, and the effect of a test compound on the activity of OGTA076 is determined. A test compound that alters the activity of OGTA076 can be identified by comparing biological samples from subjects treated with a control compound to samples from subjects treated with the test compound. Alternatively, alterations in the activity of OGTA076 can be identified by comparing the activity of OGTA076 in a subject or group of subjects before and after the administration of a test compound. The activity of OGTA076 can be assessed by detecting in a biological sample (e.g. serum, plasma, or urine) induction of a cellular signal transduction pathway of OGTA076 (e.g. intracellular Ca^{2+} , diacylglycerol, IP3, etc.), catalytic or enzymatic activity of OGTA076 or a binding partner thereof, or a cellular response, for example, cellular differentiation, or cell proliferation. Techniques known to those of skill in the art can be used to detect changes in the induction of a second messenger of OGTA076 or changes in a cellular response. For example, RT-PCR can be used to detect changes in the induction of a cellular second messenger.

[0232] In one example, a test compound that changes the level or expression of OGTA076 towards levels detected in control subjects (e.g. humans free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer) is selected for further testing or therapeutic use. In another example, a test compound that changes the activity of OGTA076 towards the activity found in control subjects (e.g. humans free from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer and pancreatic cancer) is selected for further testing or therapeutic use.

[0233] In another example, test compounds that reduce the severity of one or more symptoms associated with colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer are identified in human subjects having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer, for example subjects with severe colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. In accordance with this example, a test compound or a control compound is administered to the subjects, and the effect of a test compound on one or more symptoms of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer is determined. A test compound that reduces one or more symptoms can be identified by comparing the subjects treated with a control compound to the subjects treated with the test compound. Techniques known to physicians familiar with colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer can be used to determine whether a test compound reduces one or more symptoms associated with colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. For example, a test compound that reduces tumour burden in a subject having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer will be beneficial for subjects having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.

[0234] In one example, a test compound that reduces the severity of one or more symptoms associated with colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or

pancreatic cancer in a human having colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer is selected for further testing or therapeutic use.

Therapeutic and Prophylactic Compositions and their Use

[0235] The invention provides an antibody which is capable of specific binding to Lymphocyte antigen 75 for use in treatment of pancreatic cancer wherein said antibody contains or is conjugated to a therapeutic moiety.

[0236] Also disclosed are methods of treatment (and prophylaxis) comprising administering to a subject an effective amount of a compound of the disclosure. In one example, the compound is substantially purified (e.g. substantially free from substances that limit its effect or produce undesired side-effects). The subject is for example an animal, including but not limited to animals such as cows, pigs, horses, chickens, cats, dogs, etc., such as a mammal, and particularly a human. In a specific embodiment, a non-human mammal is the subject.

[0237] Formulations and methods of administration that can be employed when the compound comprises a nucleic acid are described above; additional appropriate formulations and routes of administration are described below.

[0238] Various delivery systems are known and can be used to administer a compound of the invention, e.g. encapsulation in liposomes, microparticles, microcapsules, recombinant cells capable of expressing the compound, receptor-mediated endocytosis (see, e.g. Wu and Wu, 1987, *J. Biol. Chem.* 262:4429-4432), construction of a nucleic acid as part of a retroviral or other vector, etc. Methods of introduction can be enteral or parenteral and include but are not limited to intradermal, intramuscular, intraperitoneal, intravenous, subcutaneous, intranasal, epidural, and oral routes. The compounds may be administered by any convenient route, for example by infusion or bolus injection, by absorption through epithelial or mucocutaneous linings (e.g. oral mucosa, rectal and intestinal mucosa, etc.) and may be administered together with other biologically active agents. Administration can be systemic or local. In addition, it may be desirable to introduce the pharmaceutical compositions of the invention into the central nervous system by any suitable route, including intraventricular and intrathecal injection; intraventricular injection may be facilitated by an intraventricular catheter, for example, attached to a reservoir, such as an Ommaya reservoir. Pulmonary administration can also be employed, e.g. by use of an inhaler or nebulizer, and formulation with an aerosolizing agent.

[0239] In a specific example, it may be desirable to administer the pharmaceutical compositions of the invention locally to the area in need of treatment; this may be achieved, for example, and not by way of limitation, by local infusion during surgery, topical application, e.g. by injection, by means of a catheter, or by means of an implant, said implant being of a porous, non-porous, or gelatinous material, including membranes, such as sialastic membranes, or fibers. In one example, administration can be by direct injection into colorectal, kidney, liver, lung, lymphoid, ovarian or pancreatic tissue or at the site (or former site) of a malignant tumor or neoplastic or pre-neoplastic tissue.

[0240] In another example, the compound can be delivered in a vesicle, in particular a liposome (see Langer, 1990, *Science* 249:1527-1533; Treat et al., in *Liposomes in the Therapy of Infectious Disease and Cancer*, Lopez-Berestein and Fidler (eds.), Liss, New York, pp. 353-365 (1989); Lopez-Berestein, *ibid.*, pp. 317-327; see generally *ibid.*)

[0241] In yet another example, the compound can be delivered in a controlled release system. In one example, a pump may be used (see Langer, *supra*; Sefton, 1987, *CRC Crit. Ref. Biomed. Eng.* 14:201; Buchwald et al., 1980, *Surgery* 88:507; Saudek et al., 1989, *N. Engl. J. Med.* 321:574). In another example, polymeric materials can be used (see *Medical Applications of Controlled Release*, Langer and Wise (eds.), CRC Pres., Boca Raton, Florida (1974); *Controlled Drug Bioavailability, Drug Product Design and Performance*, Smolen and Ball (eds.), Wiley, New York (1984); Ranger and Peppas, J., 1983, *Macromol. Sci. Rev. Macromol. Chem.* 23:61; see also Levy et al., 1985, *Science* 228:190; During et al., 1989, *Ann. Neurol.* 25:351; Howard et al., 1989, *J. Neurosurg.* 71:105). In yet another example, a controlled release system can be placed in proximity of the therapeutic target, i.e. the colon, kidney, liver, lung, ovary or pancreas thus requiring only a fraction of the systemic dose (see, e.g. Goodson, in *Medical Applications of Controlled Release*, *supra*, vol. 2, pp. 115-138 (1984)).

[0242] Other controlled release systems are discussed in the review by Langer (1990, *Science* 249:1527-1533).

[0243] In a specific example where the compound of the disclosure is a nucleic acid encoding a protein, the nucleic acid can be administered *in vivo* to promote expression of its encoded protein, by constructing it as part of an appropriate nucleic acid expression vector and administering it so that it becomes intracellular, e.g. by use of a retroviral vector (see U.S. Patent No. 4,980,286), or by direct injection, or by use of microparticle bombardment (e.g. a gene gun; Biostatic, Dupont), or coating with

lipids or cell-surface receptors or transfecting agents, or by administering it in linkage to a homeobox-like peptide which is known to enter the nucleus (see e.g. Joliot et al., 1991, Proc. Natl. Acad. Sci. USA 88:1864-1868), etc. Alternatively, a nucleic acid can be introduced intracellularly and incorporated within host cell DNA for expression, by homologous recombination.

[0244] The present disclosure also provides pharmaceutical compositions. Such compositions comprise a therapeutically effective amount of a compound, and a pharmaceutically acceptable carrier. In a specific example, the term "pharmaceutically acceptable" means approved by a regulatory agency of the Federal or a state government or listed in the U.S. Pharmacopeia or other generally recognized pharmacopeia for use in animals, and more particularly in humans. The term "carrier" refers to a diluent, adjuvant, excipient, or vehicle with which the therapeutic is administered. Such pharmaceutical carriers can be sterile liquids, such as water and oils, including those of petroleum, animal, vegetable or synthetic origin, such as peanut oil, soybean oil, mineral oil, sesame oil and the like. Water may be a suitable carrier when the pharmaceutical composition is administered intravenously. Saline solutions and aqueous dextrose and glycerol solutions can also be employed as liquid carriers, particularly for injectable solutions. Suitable pharmaceutical excipients include starch, glucose, lactose, sucrose, gelatin, malt, rice, flour, chalk, silica gel, sodium stearate, glycerol monostearate, talc, sodium chloride, dried skim milk, glycerol, propylene, glycol, water, ethanol and the like. The composition, if desired, can also contain minor amounts of wetting or emulsifying agents, or pH buffering agents. These compositions can take the form of solutions, suspensions, emulsion, tablets, pills, capsules, powders, sustained-release formulations and the like. The composition can be formulated as a suppository, with traditional binders and carriers such as triglycerides. Oral formulation can include standard carriers such as pharmaceutical grades of mannitol, lactose, starch, magnesium stearate, sodium saccharine, cellulose, magnesium carbonate, etc. Examples of suitable pharmaceutical carriers are described in "Remington's Pharmaceutical Sciences" by E.W. Martin. Such compositions will contain a therapeutically effective amount of the compound, for example in purified form, together with a suitable amount of carrier so as to provide the form for proper administration to the subject. The formulation should suit the mode of administration.

[0245] In one example, the composition is formulated in accordance with routine procedures as a pharmaceutical composition adapted for intravenous administration to human beings. Typically, compositions for intravenous administration are solutions in sterile isotonic aqueous buffer. Where necessary, the composition may also include a solubilizing agent and a local anesthetic such as lidocaine to ease pain at the site of the injection. Generally, the ingredients are supplied either separately or mixed together in unit dosage form, for example, as a dry lyophilized powder or water free concentrate in a hermetically sealed container such as an ampoule or sachette indicating the quantity of active agent. Where the composition is to be administered by infusion, it can be dispensed with an infusion bottle containing sterile pharmaceutical grade water or saline. Where the composition is administered by injection, an ampoule of sterile water for injection or saline can be provided so that the ingredients may be mixed prior to administration.

[0246] The compounds of the disclosure can be formulated as neutral or salt forms. Pharmaceutically acceptable salts include those formed with free amino groups such as those derived from hydrochloric, phosphoric, acetic, oxalic, tartaric acids, etc., and those formed with free carboxyl groups such as those derived from sodium, potassium, ammonium, calcium, ferric hydroxides, isopropylamine, triethylamine, 2-ethylamino ethanol, histidine, procaine, etc.

[0247] The amount of the compound of the disclosure which will be effective in the treatment of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer can be determined by standard clinical techniques. In addition, *in vitro* assays may optionally be employed to help identify optimal dosage ranges. The precise dose to be employed in the formulation will also depend on the route of administration, and the seriousness of the disease or disorder, and should be decided according to the judgment of the practitioner and each subject's circumstances. However, suitable dosage ranges for intravenous administration are generally about 20-500 micrograms of active compound per kilogram body weight. Suitable dosage ranges for intranasal administration are generally about 0.01 pg/kg body weight to 100 for example 10 such as 1 mg/kg body weight. Effective doses may be extrapolated from dose-response curves derived from *in vitro* or animal model test systems.

[0248] Suppositories generally contain active ingredient in the range of 0.5% to 10% by weight; oral formulations preferably contain 10% to 95% active ingredient.

[0249] The disclosure also provides a pharmaceutical pack or kit comprising one or more containers filled with one or more of the ingredients of the pharmaceutical compositions of the disclosure. Optionally associated with such container(s) can be a notice in the form prescribed by a governmental agency regulating the manufacture, use or sale of pharmaceuticals or biological products, which notice reflects (a) approval by the agency of manufacture, use or sale for human administration, (b) directions for use, or both.

Determining Abundance of OGTA076 by Imaging Technology

[0250] An advantage of determining abundance of OGTA076 by imaging technology may be that such a method is non-invasive (save that reagents may need to be administered) and there is no need to extract a sample from the subject.

[0251] Suitable imaging technologies include positron emission tomography (PET) and single photon emission computed tomography (SPECT). Visualisation of OGTA076 using such techniques requires incorporation or binding of a suitable label e.g. a radiotracer such as ^{18}F , ^{11}C or ^{123}I (see e.g. NeuroRx - The Journal of the American Society for Experimental NeuroTherapeutics (2005) 2(2), 348-360 and *idem* pages 361-371 for further details of the techniques). Radiotracers or other labels may be incorporated into OGTA076 by administration to the subject (e.g. by injection) of a suitably labelled specific ligand. Alternatively they may be incorporated into a binding affinity reagent (e.g. antibody) specific for OGTA076 which may be administered to the subject (e.g. by injection). For discussion of use of Affibodies for imaging see e.g. Orlova A, Magnusson M, Eriksson TL, Nilsson M, Larsson B, Hoiden-Guthenberg I, Widstrom C, Carlsson J, Tolmachev V, Stahl S, Nilsson FY, Tumor imaging using a picomolar affinity HER2 binding Affibody molecule, Cancer Res. 2006 Apr 15;66(8):4339-48.

Diagnosis and Treatment of Colorectal Cancer, Kidney Cancer, Liver Cancer, Lung Cancer, Lymphoid Leukaemia (Particularly Chronic Lymphocytic Leukaemia), Ovarian Cancer or Pancreatic Cancer using Immunohistochemistry

[0252] Immunohistochemistry is an excellent detection technique and may therefore be very useful in the diagnosis and treatment of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer. Immunohistochemistry may be used to detect, diagnose, or monitor colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer through the localization of OGTA076 antigens in tissue sections by the use of labeled antibodies (or other affinity reagents), derivatives and analogs thereof, which specifically bind to OGTA076, as specific reagents through antigen-antibody interactions that are visualized by a marker such as fluorescent dye, enzyme, radioactive element or colloidal gold.

[0253] The advancement of monoclonal antibody technology has been of great significance in assuring the place of immunohistochemistry in the modern accurate microscopic diagnosis of human neoplasms. The identification of disseminated neoplastically transformed cells by immunohistochemistry allows for a clearer picture of cancer invasion and metastasis, as well as the evolution of the tumour cell associated immunophenotype towards increased malignancy. Future antineoplastic therapeutical approaches may include a variety of individualized immunotherapies, specific for the particular immunophenotypical pattern associated with each individual patient's neoplastic disease. For further discussion see e.g. Bodey B, The significance of immunohistochemistry in the diagnosis and therapy of neoplasms, Expert Opin Biol Ther. 2002 Apr;2(4):371-93.

EXAMPLE 1: IDENTIFICATION OF MEMBRANE PROTEINS EXPRESSED IN COLORECTAL CANCER, CHRONIC LYMPHOCYTIC LEUKAEMIA AND PANCREATIC CANCER BLOOD AND TISSUE SAMPLES USING ID GEL ELECTROPHORESIS

[0254] Using the following Reference Protocol, membrane proteins extracted from colorectal cancer, chronic lymphocytic leukaemia and pancreatic cancer tissue samples were separated by 1D gel and analysed.

1.1 MATERIALS AND METHODS

1.1.1 - Plasma Membrane Fractionation

[0255] The cells recovered from the epithelium of a colorectal cancer, chronic lymphocytic leukaemia or pancreatic cancer were lysed and submitted to centrifugation at 1000G. The supernatant was taken, and it was subsequently centrifuged at 3000G. Once again, the supernatant was taken, and it was then centrifuged at 100 000G.

[0256] The resulting pellet was recovered and put on 15-60% sucrose gradient.

[0257] A Western blot was used to identify sub cellular markers, and the Plasma Membrane fractions were pooled.

[0258] The pooled solution was either run directly on 1D gels (see section 1.1.4 below), or further fractionated into heparin binding and nucleotide binding fractions as described below.

1.1.2 - *Plasma Membrane Heparin-binding Fraction*

[0259] The pooled solution from 1.1.1 above was applied to a Heparin column, eluted from column and run on 1D gels (see section 1.1.4 below).

1.1.3 - *Plasma Nucleotide-binding Fraction*

[0260] The pooled solution from 1.1.1 above was applied to a Cibacrom Blue 3GA column, eluted from column and run on 1D gels (see section 1.1.4 below).

1.1.4 - *1D gel technology*

[0261] Protein or membrane pellets were solubilised in 1D sample buffer (1-2 µg/µl). The sample buffer and protein mixture was then heated to 95°C for 3 min.

[0262] A 9-16% acrylamide gradient gel was cast with a stacking gel and a stacking comb according to the procedure described in Ausubel F.M. et al., eds., 1989, Current Protocols in Molecular Biology, Vol. II, Green Publishing Associates, Inc., and John Wiley & Sons, Inc., New York, section 10.2.

[0263] 30-50 micrograms of the protein mixtures obtained from detergent and the molecular weight standards (66, 45 ,31, 21,14 kDa) were added to the stacking gel wells using a 10 microlitre pipette tip and the samples run at 40mA for 5 hours.

[0264] The plates were then prised open, the gel placed in a tray of fixer (10% acetic acid, 40% ethanol, 50% water) and shaken overnight. Following this, the gel was primed by 30 minutes shaking in a primer solution (7.5% acetic acid (75ml), 0.05% SDS (5ml of 10%)). The gel was then incubated with a fluorescent dye (7.5% acetic acid, 0.06% OGS in-house dye (600µl)) with shaking for 3 hrs. Sypro Red (Molecular Probes, Inc., Eugene, Oregon) is a suitable dye for this purpose. A preferred fluorescent dye is disclosed in U.S. Application No. 09/412,168, filed on October 5.

[0265] A computer-readable output was produced by imaging the fluorescently stained gels with an Apollo 3 scanner (Oxford Glycosciences, Oxford, UK). This scanner is developed from the scanner described in WO 96/36882 and in the Ph.D. thesis of David A. Basiji, entitled "Development of a High-throughput Fluorescence Scanner Employing Internal Reflection Optics and Phase-sensitive Detection (Total Internal Reflection, Electrophoresis)", University of Washington (1997), Volume 58/12-B of Dissertation Abstracts International, page 6686. The latest embodiment of this instrument includes the following improvements: The gel is transported through the scanner on a precision lead-screw drive system. This is preferable to laying the glass plate on the belt-driven system that is defined in the Basiji thesis as it provides a reproducible means of accurately transporting the gel past the imaging optics.

[0266] The gel is secured into the scanner against three alignment stops that rigidly hold the glass plate in a known position. By doing this in conjunction with the above precision transport system and the fact that the gel is bound to the glass plate, the absolute position of the gel can be predicted and recorded. This ensures that accurate co-ordinates of each feature on the gel can be communicated to the cutting robot for excision. This cutting robot has an identical mounting arrangement for the glass plate to preserve the positional accuracy.

[0267] The carrier that holds the gel in place has integral fluorescent markers (Designated M1, M2, M3) that are used to correct the image geometry and are a quality control feature to confirm that the scanning has been performed correctly.

[0268] The optical components of the system have been inverted. The laser, mirror, waveguide and other optical components are now above the glass plate being scanned. The embodiment of the Basiji thesis has these underneath. The glass plate is therefore mounted onto the scanner gel side down, so that the optical path remains through the glass plate. By doing this, any particles of gel that may break away from the glass plate will fall onto the base of the instrument rather than into the optics.

[0269] In scanning the gels, they were removed from the stain, rinsed with water and allowed to air dry briefly and imaged on the Apollo 3. After imaging, the gels were sealed in polyethylene bags containing a small volume of staining solution, and then stored at 4°C.

[0270] Apparent molecular weights were calculated by interpolation from a set of known molecular weight markers run alongside the samples.

1.1.5 - Recovery and analysis of selected proteins

[0271] Proteins were robotically excised from the gels by the process described in U.S. Patent No. 6,064,754, Sections 5.4 and 5.6, 5.7, 5.8, as is applicable to 1D-electrophoresis, with modification to the robotic cutter as follows: the cutter begins at the top of the lane, and cuts a gel disc 1.7mm in diameter from the left edge of the lane. The cutter then moves 2mm to the right, and 0.7mm down and cuts a further disc. This is then repeated. The cutter then moves back to a position directly underneath the first gel cut, but offset by 2.2mm downwards, and the pattern of three diagonal cuts are repeated. This is continued for the whole length of the gel.

[0272] NOTE: If the lane is observed to broaden significantly then a correction can be made also sideways i.e. instead of returning to a position directly underneath a previous gel cut, the cut can be offset slightly to the left (on the left of the lane) and/or the right (on the right of the lane). The proteins contained within the gel fragments were processed to generate tryptic peptides; partial amino acid sequences of these peptides were determined by mass spectroscopy as described in WO98/53323 and Application No. 09/094,996, filed June 15, 1998.

[0273] Proteins were processed to generate tryptic digest peptides. Tryptic peptides were analyzed by mass spectrometry using a PerSeptive Biosystems Voyager- DETM STR Matrix-Assisted Laser Desorption Ionization Time-of-Flight (MALDI-TOF) mass spectrometer, and selected tryptic peptides were analyzed by tandem mass spectrometry (MS/MS) using a Micromass Quadrupole Time-of-Flight (Q-TOF) mass spectrometer (Micromass, Altrincham, U.K.) equipped with a nanoflowTM electrospray Z-spray source. For partial amino acid sequencing and identification of OGTA076, uninterpreted tandem mass spectra of tryptic peptides were searched using the SEQUEST search program (Eng et al., 1994, J. Am. Soc. Mass Spectrom. 5:976-989), version v.C.1. Criteria for database identification included: the cleavage specificity of trypsin; the detection of a suite of a, b and y ions in peptides returned from the database, and a mass increment for all Cys residues to account for carbamidomethylation. The database searched was a database constructed of protein entries in the non-redundant database held by the National Centre for Biotechnology Information (NCBI) which is accessible at www.ncbi.nlm.nih.gov. Following identification of proteins through spectral-spectral correlation using the SEQUEST program, masses detected in MALDI-TOF mass spectra were assigned to tryptic digest peptides within the proteins identified. In cases where no amino acid sequences could be identified through searching with uninterpreted MS/MS spectra of tryptic digest peptides using the SEQUEST program, tandem mass spectra of the peptides were interpreted manually, using methods known in the art. (In the case of interpretation of low-energy fragmentation mass spectra of peptide ions see Gaskell et al., 1992, Rapid Commun. Mass Spectrom. 6:658-662).

1.1.6 - Discrimination of colorectal cancer, chronic lymphocytic leukaemia or pancreatic cancer associated proteins

[0274] The process to identify OGTA076 uses the peptide sequences obtained experimentally by mass spectrometry described above of naturally occurring human proteins to identify and organize coding exons in the published human genome sequence.

[0275] Recent dramatic advances in defining the chemical sequence of the human genome have led to the near completion of this immense task (Venter, J.C. et al. (2001). The sequence of the human genome. *Science* 16: 1304-51; International Human Genome Sequencing Consortium. (2001). Initial sequencing and analysis of the human genome *Nature* 409: 860-921). There is little doubt that this sequence information will have a substantial impact on our understanding of many biological processes, including molecular evolution, comparative genomics, pathogenic mechanisms and molecular medicine. For the full medical value inherent in the sequence of the human genome to be realised, the genome needs to be 'organised' and annotated. By this, is meant at least the following three things: (i) The assembly of the sequences of the individual portions of the genome into a coherent, continuous sequence for each chromosome. (ii) The unambiguous identification of those regions of each chromosome that contain genes. (iii) Determination of the fine structure of the genes and the properties of its mRNA and protein products. While the definition of a 'gene' is an increasingly complex issue (H Pearson: What is a gene? *Nature* (2006) 24: 399 - 401), what is of immediate interest for drug discovery and development is a catalogue of those genes that encode functional, expressed

proteins. A subset of these genes will be involved in the molecular basis of most if not all pathologies. Therefore an important and immediate goal for the pharmaceutical industry is to identify all such genes in the human genome and describe their fine structure.

Processing and integration of peptide masses, peptide signatures, ESTs and Public Domain Genomic Sequence Data to form OGAP® database

[0276] Discrete genetic units (exons, transcripts and genes) were identified using the following sequential steps:

1. 1. A 'virtual transcriptome' is generated, containing the tryptic peptides which map to the human genome by combining the gene identifications available from Ensembl and various gene prediction programs. This also incorporates SNP data (from dbSNP) and all alternate splicing of gene identifications. Known contaminants were also added to the virtual transcriptome.
2. 2. All tandem spectra in the OGeS Mass Spectrometry Database are interpreted in order to produce a peptide that can be mapped to one in the virtual transcriptome. A set of automated spectral interpretation algorithms were used to produce the peptide identifications.
3. 3. The set of all mass-matched peptides in the OGeS Mass Spectrometry Database is generated by searching all peptides from transcripts hit by the tandem peptides using a tolerance based on the mass accuracy of the mass spectrometer, typically 20ppm.
4. 4. All tandem and mass-matched peptides are combined in the form of "protein clusters". This is done using a recursive process which groups sequences into clusters based on common peptide hits. Biological sequences are considered to belong to the same cluster if they share one or more tandem or mass-matched peptide.
5. 5. After initial filtering to screen out incorrectly identified peptides, the resulting clusters are then mapped on the human genome.
6. 6. The protein clusters are then aggregated into regions that define preliminary gene boundaries using their proximity and the co-observation of peptides within protein clusters. Proximity is defined as the peptide being within 80,000 nucleotides on the same strand of the same chromosome. Various elimination rules, based on cluster observation scoring and multiple mapping to the genome are used to refine the output. The resulting 'confirmed genes' are those which best account for the peptides and masses observed by mass spectrometry in each cluster. Nominal co-ordinates for the gene are also an output of this stage.
7. 7. The best set of transcripts for each confirmed gene are created from the protein clusters, peptides, ESTs, candidate exons and molecular weight of the original protein spot.
8. 8. Each identified transcript was linked to the sample providing the observed peptides.
9. 9. Use of an application for viewing and mining the data. The result of steps 1-8 was a database containing genes, each of which consisted of a number of exons and one or more transcripts. An application was written to display and search this integrated genome / proteome data. Any features (OMIM disease locus, InterPro etc.) that had been mapped to the same Golden Path co-ordinate system by Ensembl could be cross-referenced to these genes by coincidence of location and fine structure.

Results

[0277] The process was used to generate approximately 1 million peptide sequences to identify protein-coding genes and their exons resulted in the identification of protein sequences for 18083 genes across 67 different tissues and 57 diseases including 506 genes in bladder cancer, 4,713 genes in breast cancer, 766 genes in burkitt's lymphoma, 1,371 genes in cervical cancer, 2,424 genes in chronic lymphocytic leukaemia, 949 genes in colorectal cancer, 1,782 genes in hepatocellular carcinoma, 1,004 genes in kidney cancer, 978 genes in lung cancer, 1,764 genes in melanoma, 1,033 genes in ovarian cancer, 2,961 genes in pancreatic cancer and 3,307 genes in prostate cancer, illustrated here by OGTA076 isolated and identified from colorectal cancer, chronic lymphocytic leukaemia, kidney cancer, liver cancer, lung cancer, ovarian cancer and pancreatic cancer samples. Following comparison of the experimentally determined sequences with sequences in the OGAP® database, OGTA076 showed a high degree of specificity to colorectal cancer, chronic lymphocytic leukaemia, kidney cancer, liver cancer, lung cancer, ovarian cancer and pancreatic cancer indicative of the prognostic and diagnostic nature.

1.2 RESULTS

[0278] These experiments identified OGTA076, in its three different isoforms, as further described herein. The full-length OGTA076 was detected in the plasma membrane of colorectal cancer, chronic lymphocytic leukaemia and pancreatic cancer samples and was not detected in the cytosol.

[0279] Figure 2 shows the Protein Index for OGTA076. For each gene, the protein index uses the mass spectrometry data to assign a score to each disease, relative to the global database. The Protein Index can then be used to identify cancer specific genes with a high score in cancer indications and low/negligible scores in normal and other diseases. The index contains ~ 1 million peptides sequenced via mass spectrometry from 56 diseases. For each gene, this yields a score for each disease and subcellular location. The results are summarized below:

Protein Index Report for OGTA076

Indications positive:

[0280]

Colorectal cancer

Chronic lymphocytic leukaemia

Kidney cancer

Liver cancer

Lung cancer

Ovarian cancer

Pancreatic cancer

Disease controls

Acute monocytic leukaemia	Diabetes and Obesity	Migraine, acute
Acute T-cell leukaemia	Diverticulitis	Multiple sclerosis
Alzheimer's Disease	Dyslipidaemia	Neuroblastoma
Arthritis	Emphysema	Normal
Asthma	Focal apocrine metaplasia	Obesity
Atherosclerosis	Gastric cancer	Osteoarthritis
B-cell non-Hodgkin's lymphoma	Gaucher disease	Osteosarcoma
Bladder carcinoma	Glioblastoma	Ovarian cancer
Breast cancer	Hepatoblastoma	Pancreatic cancer
Breast diseases, benign	Hypertension	Prostate cancer
Burkitt's lymphoma	Lactational foci	Prostatic diseases, benign
Bursitis	Kidney cancer	Prostatitis
Cancer, unspecified	Leukaemia, unspecified	Retinoblastoma
Cervical cancer	Liver cancer	Schizophrenia
Chronic lymphocytic leukaemia	Liver cirrhosis	Skin ulcer
Chronic obstructive pulmonary disease	Lung cancer	Smoker
Colorectal cancer	Lymphoma, histiocytic	Teratocarcinoma
Dementia, vascular	Melanoma	

Depression	Metabolic syndrome X	
Subcellular location		
Birbeck Granules	Membrane	Secreted
Cell surface digest	Membrane Glycoprotein Binding Fraction	Soluble Fraction
Chromatin Fraction	Mitochondria	Supernatant
Crude Cell Membrane	Nucleus	Whole Cell
Cytosol	Peroxisomes	
Golgi/Mitochondrial	Plasma Membrane	

[0281] Figure 2 shows the Protein Index for OGTA076 is high in colorectal cancer, medium in chronic lymphocytic leukaemia, medium in kidney cancer, low in liver cancer, high in lung cancer, medium in ovarian cancer and low in pancreatic cancer plasma membrane and very low in normal plasma membrane and membrane. OGTA076 was not detected in any other diseases. This indicates that OGTA076 is potentially a good marker for colorectal cancer, kidney cancer, liver cancer, lung cancer, chronic lymphocytic leukaemia, ovarian cancer and pancreatic cancer.

EXAMPLE 2: IDENTIFICATION OF MEMBRANE PROTEINS EXPRESSED IN COLORECTAL CANCER, KIDNEY CANCER, LIVER CANCER, LUNG CANCER OR OVARIAN CANCER BLOOD AND TISSUE SAMPLES

[0282] Using the following Reference Protocol, membrane proteins extracted from colorectal cancer, kidney cancer, liver cancer, lung cancer and ovarian cancer tissue and normal adjacent colorectal, kidney, liver, lung and ovarian tissue samples were digested, labelled with Isotope Tagging for Absolute & Relative Quantitation reagents (iTRAQ; Applied Biosystems, Foster City, CA, USA) and resulting peptides sequenced by tandem mass spectrometry.

2.1 MATERIALS AND METHODS

2.1.1 - *Plasma Membrane Fractionation*

[0283] The cells recovered from a colorectal cancer, kidney cancer, liver cancer, lung cancer or ovarian cancer or normal adjacent colorectal, kidney, liver, lung or ovarian tissue were lysed and submitted to centrifugation at 1000G. The supernatant was taken, and it was subsequently centrifuged at 3000G. Once again, the supernatant was taken, and it was then centrifuged at 100 000G.

[0284] The resulting pellet was recovered and put on 15-60% sucrose gradient.

[0285] A Western blot was used to identify sub cellular markers, and the Plasma Membrane fractions were pooled.

[0286] The pooled solution was then analysed directly by iTRAQ (see section 2.1.2 below).

2.1.2 - *iTRAQ methodology*

[0287] Membrane protein pellets from colorectal cancer, kidney cancer, liver cancer, lung cancer or ovarian cancer and normal adjacent colorectal, kidney, liver, lung or ovarian tissue were solubilised in sample buffer (2-4 µg/µl in 0.5% SDS) by the addition of buffer and then heating to 95°C for 3 min.

[0288] To a volume of each protein solution equating to 50µg, 150µl or 0.5M triethylammonium bicarbonate (TEAB) solution was added. To each sample, 3µl of 50mM tris-(2-carboxyethyl)phosphine was added and the mixture was incubated at 60°C for 1 hour. 1µl of cysteine blocking reagent, 200mM methyl methanethiosulphonate (MMTS) in isopropanol, was then added. After incubation at room temperature for 10 minutes, 15µl of 1µg/µl trypsin was added to each sample followed by incubation at 37°C overnight.

[0289] The digested samples were dried under a vacuum and re-constituted with 30µl of 0.5M TEAB solution. 70µl ethanol was added to each of the four iTRAQ reagents (114/115/116/117) and one reagent added to each of the four samples analysed (two colorectal cancer, kidney cancer, liver cancer, lung cancer or ovarian cancer samples and two corresponding normal adjacent tissue samples) and left at room temperature for 1 hour. The specific reagent added to each sample was recorded. The four labeled samples were combined & vortexed.

[0290] The combined sample was reduced to dryness under a vacuum and de-salted by loading onto a C18 spin column, washing with aqueous solvent and then eluting with 70% acetonitrile. The sample fraction was again reduced to dryness and then re-dissolved in 40µl of solvent A (97.9 water, 2% acetonitrile, 0.1% formic acid) prior to ion exchange fractionation.

2.1.3 - Fractionation and analysis of labeled peptides

[0291] The sample was fractionated by strong cation exchange chromatography using an Agilent 1200 chromatograph (Agilent, Santa Clara, CA, USA). Samples were eluted off an Agilent Zorbax Bio-SCXII column (3.5µm; 50 x 0.8mm) using a 20µl/min gradient of 0-100mM sodium acetate over 20 minutes and then to 1M over 10 minutes. 1 minute fractions were collected over the 30 minute run.

[0292] Each fraction was analysed by liquid chromatography/mass spectrometry using an Agilent 1200 chromatograph fitted with a Zorbax 300SB-C18 (150mm x 75µm) and an Agilent 6510 quadrupole - time-of-flight instrument (Agilent, Santa Clara, CA, USA). Peptides were eluted with a 300nl/min gradient increasing from 15% to 45% acetonitrile in 60 minutes. Data was acquired in auto MS/MS mode such that up to 3 precursor ions above the intensity threshold were selected and product ion spectra accumulated to facilitate the sequencing of the labeled peptides. Raw was processed to create peak lists using Spectrum Mill software (Agilent, Santa Clara, CA, USA).

2.1.4 - Amino acid sequence analysis of labeled peptides

[0293] For partial amino acid sequencing and identification of OGTA076, uninterpreted tandem mass spectra of tryptic peptides were searched using the SEQUEST search program (Eng et al., 1994, J. Am. Soc. Mass Spectrom. 5:976-989). Criteria for database identification included: the cleavage specificity of trypsin; the detection of a suite of a, b and y ions in peptides returned from the database, and a mass increment for all cysteine residues to account for modification with methyl methanethiosulphonate and the addition of iTRAQ labels to free amines (N-terminus & lysine). The data was searched through IPI Human v3.23 (www.ebi.ac.uk/IPI/IPIhuman.html).

2.1. 5 - Discrimination of colorectal cancer, kidney cancer, liver cancer, lung cancer or ovarian cancer associated proteins

[0294] The process described in Example 1 section 1.1.6 was employed to discriminate the colorectal cancer, kidney cancer, liver cancer, lung cancer and ovarian cancer associated proteins in the experimental samples.

2.2 RESULTS

[0295] These experiments identified OGTA076, in its three different isoforms, as further described herein. The full-length OGTA076 was detected in the plasma membrane of colorectal cancer, kidney cancer, liver cancer, lung cancer and ovarian cancer samples. The iTRAQ analysis showed that levels of OGTA076 in the cancer samples were higher than in the matched normal adjacent tissue samples.

[0296] Figure 2 shows the Protein Index for OGTA076. See Example 1 section 1.2 for a description of the Protein Index for OGTA076.

[0297] Throughout the specification and the claims which follow, unless the context requires otherwise, the word 'comprise', and variations such as 'comprises' and 'comprising', will be understood to imply the inclusion of a stated integer, step, group of

integers or group of steps but not to the exclusion of any other integer, step, group of integers or group of steps.

Sequence	Seq. ID
MRTGWATPRRPAGLMLLFWFFDIAEPSGRAANDPFTIVHGNTGKCIKPVYGVIVADDCEDEDKLWKWS QHRLFLHSQKCLGLDITKSVNELRMSFCDSSAMLWKCBEHHSLYGAARYLALKDGHGTAISASDWKK GGSEESLCQDPYHEIYTRDGNSYGRPCEFPFLIDGTWHDCILDEDHSGPWCATTINYEYDRKGICLKPE NGCEDNWEKNEQFGSCYQFNQTALSWKEAYVSCQNGADLLSINSAEALTYKEKEGIAKIFWIGLNLQY SARGNEWSDHKPLNLFNWDPDRSAPTIIGGSSCARMDAESGLWQSFSCAEOLPYVCRKPLNNTEVLTDWWT YSDTRCDAGWLPNNGFCYLLVNESENNSWDKAHACKAFAFSSDLISIHSLADVEVVVTKLHNEDIKEEWWIGLK NINIPTLFWQSDGKWRHGETCYKIEDEVPFGTNCNLTITSRFEQEYLNDLMKYDKSLRKYFWTGLRDVDSCE DPMCPDEGWKRHGETCYKIEDEVPFGTNCNLTITSRFEQEYLNDLMKYDKSLRKYFWTGLRDVDSCE YMWATGRRRAVFTFSNFWLPEPASPGCVAAMSTGKSVGKWEVKDCRSFKALSIKKMSGPLGPEEASPKP DPPCGWQSFPSLSCYKVFHAERIVRKRNNFVLSQHSLDGSSESSFWSDGSTFDYIPWKGQTSPGNCVLL DPKGWTKHEKCNVKDGAICYKPTSKKLSRLTYSSRCPAAKENGSRWIQYKHCYKSDQALHSFSEAKKL CSKHDHSATIVS1KDEDENKFVSRMLRENNNITMRVWLGLSQHSVDQSWSLDGEVTVKWEENKSXGVG RCSEMLIASNETWKVCEHGFGRVVKVPLGPDTAIIAIVATLSILVLMGGLIWFQHRRLHLAGFSSV RYAQGVNEDEIMLPFHD	1
MRTGWATPRRPAGLMLLFWFFDIAEPSGRAANDPFTIVHGNTGKCIKPVYGVIVADDCEDEDKLWKWS QHRLFLHSQKCLGLDITKSVNELRMSFCDSSAMLWKCBEHHSLYGAARYLALKDGHGTAISASDWKK GGSEESLCQDPYHEIYTRDGNSYGRPCEFPFLIDGTWHDCILDEDHSGPWCATTINYEYDRKGICLKPE NGCEDNWEKNEQFGSCYQFNQTALSWKEAYVSCQNGADLLSINSAEALTYKEKEGIAKIFWIGLNLQY SARGNEWSDHKPLNLFNWDPDRSAPTIIGGSSCARMDAESGLWQSFSCAEOLPYVCRKPLNNTEVLTDWWT YSDTRCDAGWLPNNGFCYLLVNESENNSWDKAHACKAFAFSSDLISIHSLADVEVVVTKLHNEDIKEEWWIGLK NINIPTLFWQSDGKWRHGETCYKIEDEVPFGTNCNLTITSRFEQEYLNDLMKYDKSLRKYFWTGLRDVDSCE DPMCPDEGWKRHGETCYKIEDEVPFGTNCNLTITSRFEQEYLNDLMKYDKSLRKYFWTGLRDVDSCE YMWATGRRRAVFTFSNFWLPEPASPGCVAAMSTGKSVGKWEVKDCRSFKALSIKKMSGPLGPEEASPKP DPPCGWQSFPSLSCYKVFHAERIVRKRNNFVLSQHSLDGSSESSFWSDGSTFDYIPWKGQTSPGNCVLL DPKGWTKHEKCNVKDGAICYKPTSKKLSRLTYSSRCPAAKENGSRWIQYKHCYKSDQALHSFSEAKKL CSKHDHSATIVS1KDEDENKFVSRMLRENNNITMRVWLGLSQHSVDQSWSLDGEVTVKWEENKSXGVG RCSEMLIASNETWKVCEHGFGRVVKVPLGPDTAIIAIVATLSILVLMGGLIWFQHRRLHLAGFSSV RYAQGVNEDEIMLPFHD	2
MRTGWATPRRPAGLMLLFWFFDIAEPSGRAANDPFTIVHGNTGKCIKPVYGVIVADDCEDEDKLWKWS QHRLFLHSQKCLGLDITKSVNELRMSFCDSSAMLWKCBEHHSLYGAARYLALKDGHGTAISASDWKK GGSEESLCQDPYHEIYTRDGNSYGRPCEFPFLIDGTWHDCILDEDHSGPWCATTINYEYDRKGICLKPE NGCEDNWEKNEQFGSCYQFNQTALSWKEAYVSCQNGADLLSINSAEALTYKEKEGIAKIFWIGLNLQY SARGNEWSDHKPLNLFNWDPDRSAPTIIGGSSCARMDAESGLWQSFSCAEOLPYVCRKPLNNTEVLTDWWT YSDTRCDAGWLPNNGFCYLLVNESENNSWDKAHACKAFAFSSDLISIHSLADVEVVVTKLHNEDIKEEWWIGLK NINIPTLFWQSDGKWRHGETCYKIEDEVPFGTNCNLTITSRFEQEYLNDLMKYDKSLRKYFWTGLRDVDSCE DPMCPDEGWKRHGETCYKIEDEVPFGTNCNLTITSRFEQEYLNDLMKYDKSLRKYFWTGLRDVDSCE YMWATGRRRAVFTFSNFWLPEPASPGCVAAMSTGKSVGKWEVKDCRSFKALSIKKMSGPLGPEEASPKP DPPCGWQSFPSLSCYKVFHAERIVRKRNNFVLSQHSLDGSSESSFWSDGSTFDYIPWKGQTSPGNCVLL DPKGWTKHEKCNVKDGAICYKPTSKKLSRLTYSSRCPAAKENGSRWIQYKHCYKSDQALHSFSEAKKL CSKHDHSATIVS1KDEDENKFVSRMLRENNNITMRVWLGLSQHSVDQSWSLDGEVTVKWEENKSXGVG RCSEMLIASNETWKVCEHGFGRVVKVPLGPDTAIIAIVATLSILVLMGGLIWFQHRRLHLAGFSSV RYAQGVNEDEIMLPFHD	3
MRTGWATPRRPAGLMLLFWFFDIAEPSGRAANDPFTIVHGNTGKCIKPVYGVIVADDCEDEDKLWKWS QHRLFLHSQKCLGLDITKSVNELRMSFCDSSAMLWKCBEHHSLYGAARYLALKDGHGTAISASDWKK GGSEESLCQDPYHEIYTRDGNSYGRPCEFPFLIDGTWHDCILDEDHSGPWCATTINYEYDRKGICLKPE NGCEDNWEKNEQFGSCYQFNQTALSWKEAYVSCQNGADLLSINSAEALTYKEKEGIAKIFWIGLNLQY SARGNEWSDHKPLNLFNWDPDRSAPTIIGGSSCARMDAESGLWQSFSCAEOLPYVCRKPLNNTEVLTDWWT YSDTRCDAGWLPNNGFCYLLVNESENNSWDKAHACKAFAFSSDLISIHSLADVEVVVTKLHNEDIKEEWWIGLK NINIPTLFWQSDGKWRHGETCYKIEDEVPFGTNCNLTITSRFEQEYLNDLMKYDKSLRKYFWTGLRDVDSCE DPMCPDEGWKRHGETCYKIEDEVPFGTNCNLTITSRFEQEYLNDLMKYDKSLRKYFWTGLRDVDSCE YMWATGRRRAVFTFSNFWLPEPASPGCVAAMSTGKSVGKWEVKDCRSFKALSIKKMSGPLGPEEASPKP DPPCGWQSFPSLSCYKVFHAERIVRKRNNFVLSQHSLDGSSESSFWSDGSTFDYIPWKGQTSPGNCVLL DPKGWTKHEKCNVKDGAICYKPTSKKLSRLTYSSRCPAAKENGSRWIQYKHCYKSDQALHSFSEAKKL CSKHDHSATIVS1KDEDENKFVSRMLRENNNITMRVWLGLSQHSVDQSWSLDGEVTVKWEENKSXGVG RCSEMLIASNETWKVCEHGFGRVVKVPLGPDTAIIAIVATLSILVLMGGLIWFQHRRLHLAGFSSV RYAQGVNEDEIMLPFHD	4

Sequence	Seq. ID
AFSSDLISIHSLADVEVVVTK	5
CEHHSLYGAAR	6
CLGLDITK	7
CSMLIASNETWKK	8
DGAICYKPTK	9
DGHGTAISNASDVWKK	10
DVDSCGEYNWATVGGRRR	11
EEVWIGLK	12
EFMLRPFACDTK	13
EGIAK	14
ELTYSNFHPLLVSGR	15
ENNNTMR	16
EVKPVDSVK	17
FEQEYLNDLMK	18
FPVTFGEECLYMSAK	19
GNCEVSSVEGLCK	20
GWHFYDDR	21
HDHSATIVSIK	22
HFVSLCQK	23
HGETCYK	24
HMATTQDEVHTK	25
IANISGDGQK	26
IEMVDYK	27
IIPK	28
IPENFFEEESR	29
ISEWPIDDHFTYSR	30
KGNCEVSSVEGLCK	31
KRNWEEAER	32
KVECEHGFGR	33
KYFWTGLR	34
LALK	35
LFHLHSQK	36
LHLAGFSSVR	37
LHNEDIK	38
LNDASSDK	39
LNPK	40
LPFICEK	41
MCPPDEGWKR	42
MFSCDSSAMLWWK	43
MSGPLGPEEASPK	44
NNSLMWFDK	45
NWEEAER	46
QTLQNASETVK	47

Sequence	Seq. ID
RGWHFYDDR	48
RHGETCYK	49
RLHFSR	50
RNWEEAER	51
SDQALHSFSEAK	52
SHILSIR	53
SNFHPLLVSGR	54
SPDLQGSWQWSDR	55
TLTWHSAK	56
TPDWYNPDR	57
TPLSYTHWR	58
TPVSTIIMPNEFQQDYDIR	59
VECEHGFGR	60
VFHRPWR	61
VIEEAVYFHQH	62
VQCSEQWIPFQNK	63
WVSQHR	64
YFWTGLR	65
YLNNLYK	66

PREFERRED EXAMPLES.

[0298] The following are preferred examples of the disclosure:

1. 1. OGTA076, or a fragment thereof, a nucleic acid encoding same or an immunogenic composition comprising OGTA076 or an epitope containing fragment thereof, or nucleic acid encoding OGTA076 or a fragment thereof optionally together with an immunostimulant.
2. 2. OGTA076, or a fragment thereof, a nucleic acid encoding same or an immunogenic composition as defined in claim 1, wherein composition is a vaccine.
3. 3. OGTA076, or a fragment thereof, a nucleic acid encoding same or an immunogenic composition as defined in claim 1 or 2 for use in treating or preventing disease.
4. 4. OGTA076, or a fragment thereof, a nucleic acid encoding same or an immunogenic composition according to claim 3 wherein the disease is cancer.
5. 5. OGTA076, or a fragment thereof, a nucleic acid encoding same or a composition according to any one of claims 4 for use in preventing or treating colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer
6. 6. A method for treating or preventing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or of raising an immune response, which comprises administering to a subject in need thereof a therapeutically effective amount of:
 1. a) OGTA076, or an immunogenic fragment thereof,
 2. b) nucleic acid encoding OGTA076 or an immunogenic fragment thereof or
 3. c) an immunogenic composition comprising OGTA076 or an epitope containing fragment thereof, or nucleic acid encoding OGTA076 or a fragment thereof optionally together with an immunostimulant.
7. 7. A method according to the claim 6, wherein the immunogenic fragment is an epitope containing fragment.
8. 8. A method according to claim 6, wherein the immunostimulant is an adjuvant.
9. 9. An affinity reagent capable of specific binding to OGTA076 or a fragment thereof.
10. 10. An affinity reagent according to claim 9 which contains or is conjugated to a detectable label.
11. 11. An affinity reagent according to claim 9 which contains or is conjugated to a therapeutic moiety.
12. 12. An affinity reagent according to claim 11 wherein the therapeutic moiety is a cytotoxic moiety or a radioactive isotope.

13. 13. An affinity reagent according to any one of claims 9 to 12 which is an antibody.
14. 14. An antibody according to claim 13, which is an isolated monoclonal antibody, or an antigen-binding portion thereof, an antibody fragment, or an antibody mimetic.
15. 15. An isolated monoclonal antibody according to claim 14, wherein said antibody is a full-length antibody of an IgG1, IgG2, IgG3, or IgG4 isotype.
16. 16. An isolated monoclonal antibody according to claim 14, wherein said antibody is selected from the group consisting of: a whole antibody, an antibody fragment, a humanised antibody, a single chain antibody, an immunoconjugate, a defucosylated antibody, and a bispecific antibody.
17. 17. An antibody fragment according to claim 14, wherein the fragment is selected from the group consisting of: a UniBody, a domain antibody and a Nanobody.
18. 18. An antibody mimetic according to claim 14, wherein the mimetic is an Affibody.
19. 19. A monoclonal antibody according to claim 14, which has cytotoxicity against OGTA076 antigen expressing cells in the presence of a human complement.
20. 20. A monoclonal antibody according to claim 14, which has cytotoxicity against OGTA076 antigen expressing cells in the presence of human immune effector cells.
21. 21. An affinity reagent or an antibody as defined in any one of claims 9 to 20 for use in treatment or prophylaxis, for example treatment or prophylaxis of cancer such as colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer
22. 22. A pharmaceutical composition comprising a therapeutically effective amount of an affinity reagent or a fragment thereof or an antibody or a fragment thereof as defined in any one of claims 9 to 21, and a pharmaceutically acceptable diluent or carrier
23. 23. A pharmaceutical composition according to claim 22 comprising one or more affinity reagents and/or antibodies as defined in any one of claims 9 to 21 and a pharmaceutically acceptable excipient.
24. 24. An agent as defined in any one of claims 9 to 21 or a composition as defined in claim 22 or claim 23 for use in treating or preventing disease.
25. 25. An agent according to claim 24 wherein the disease is cancer.
26. 26. An agent according to claim 25 wherein the cancer is colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.
27. 27. A method for treating or preventing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer which comprises administering to a subject in need thereof a therapeutically effective amount of an agent as defined in any one of claims 9 to 21 or a composition as defined in claim 22 or claim 23.
28. 28. An isolated nucleic acid molecule encoding the isolated antibody or antigen-binding portion thereof of claim 14.
29. 29. An expression vector comprising the nucleic acid molecule of claim 28.
30. 30. A host cell comprising the expression vector of claim 29.
31. 31. A method for preparing an anti-OGTA076 antibody, said method comprising the steps of:
 1. a) obtaining a host cell that contains one or more nucleic acid molecules encoding the antibody of claim 13;
 2. b) growing the host cell in a host cell culture;
 3. c) providing host cell culture conditions wherein the one or more nucleic acid molecules are expressed; and
 4. d) recovering the antibody from the host cell or from the host cell culture.
32. 32. A method of making the antibody of claim 14, comprising the steps of:
 1. a) immunising a transgenic animal comprising human immunoglobulin genes with an OGTA076 peptide;
 2. b) recovering B-cells from said transgenic animal;
 3. c) making hybridomas from said B-cells;
 4. d) selecting hybridomas that express antibodies that bind OGTA076; and
 5. e) recovering said antibodies that bind OGTA076 from said selected hybridomas.
33. 33. A method of making anti-OGTA076 antibodies, comprising the steps of:
 1. a) immunising a transgenic animal comprising human immunoglobulin genes with an OGTA076 peptide;
 2. b) recovering mRNA from the B-cells of said transgenic animal;
 3. c) converting said mRNA to cDNA;
 4. d) expressing said cDNA in phages such that anti-OGTA076 antibodies encoded by said cDNA are presented on the surface of said phages;
 5. e) selecting phages that present anti-OGTA076 antibodies;
 6. f) recovering nucleic acid molecules from said selected phages that encode said anti-OGTA076 immunoglobulins;
 7. g) expressing said recovered nucleic acid molecules in a host cell; and recovering antibodies from said host cell that bind OGTA076.
34. 34. A process for preparing a monoclonal antibody which is capable of immunospecific binding to OGTA076 which comprises the step of immunising a non-human animal with a protein which is OGTA076 or an immunogenic fragment

thereof or a fusion protein containing OGTA076 or an immunogenic fragment thereof or immunising a non-human animal with cells which express such a protein in either case optionally together with an immunostimulant.

35. 35. A process according to claim 34 which further comprises the step of isolating antibody producing cells from said animal and immortalising them by fusing them with immortal cells to produce an antibody-producing hybridoma.
36. 36. A process according to claim 33 which further comprises the step of isolating antibodies from said hybridoma.
37. 37. An isolated monoclonal antibody which is capable of immunospecific binding to the stem of OGTA076 obtainable by the process of any one of claims 34 to 36.
38. 38. An isolated monoclonal antibody according to claim 37 which is a human monoclonal antibody by virtue of the non-human animal being transgenic and adapted to express human immunoglobulin heavy and light chain genes and not endogenous immunoglobulin heavy and light chain genes.
39. 39. An isolated humanised monoclonal antibody which is capable of immunospecific binding to the stem of OGTA076 characterised by having one or more CDRs of a monoclonal antibody obtainable by the process of any one of claims 34 to 36.
40. 40. A fragment or derivative of a monoclonal antibody according to any one of claims 37-39.
41. 41. A kit containing one or more affinity reagents or antibody according to any one of claims 9 to 21 or a composition as defined in claim 22 or 23, wherein said affinity agent is suitable for use in treatment and/or diagnosis.
42. 42. A kit as defined claim 41, which further comprises instructions for use of said affinity agent as defined in any one of claims 243 to 26.
43. 43. A kit according to claim 41 or 42 wherein the affinity agent is lyophilised for later reconstitution.
44. 44. A kit according to claim 43, wherein the kit further comprises a solution for reconstituting said affinity agent.
45. 45. A kit according to claim 44, wherein the solution is an isotonic aqueous solution.
46. 46. A kit as defined in any one of claims 41 to 45 which further comprises a hybridising agent.
47. 47. A method of screening for compounds that modulate the activity of OGTA076, the method comprising: (a) contacting OGTA076 or a biologically active portion thereof with a candidate compound; and (b) determining whether activity of OGTA076 is thereby modulated.
48. 48. A method according to claim 47 which comprises (a) contacting OGTA076 or a biologically active portion thereof with a candidate compound in a sample; and (b) comparing the activity of OGTA076 or a biologically active portion thereof in said sample after contact with said candidate compound with the activity of OGTA076 or a biologically active portion thereof in said sample before contact with said candidate compound, or with a reference level of activity.
49. 49. A method according to claim 47 or claim 48 which is a method of screening for compounds that inhibit activity of OGTA076.
50. 50. A method according to any one of claims 47 to 49 wherein OGTA076 or a biologically active portion thereof is expressed on or by a cell.
51. 51. A method according to any one of claims 47 to 49 wherein OGTA076 or a biologically active portion thereof is isolated from cells which express it.
52. 52. A method according to claim 51 wherein OGTA076 or a biologically active portion thereof is immobilised onto a solid phase.
53. 53. A method of screening for compounds that modulate the expression of OGTA076 or nucleic acid encoding OGTA076, the method comprising: (a) contacting cells expressing OGTA076 or nucleic acid encoding OGTA076 with a candidate compound; and (b) determining whether expression of OGTA076 or nucleic acid encoding OGTA076 is thereby modulated.
54. 54. A method according to claim 53 which comprises (a) contacting cells expressing OGTA076 or nucleic acid encoding OGTA076 with a candidate compound in a sample; and (b) comparing the expression of OGTA076 or nucleic acid encoding OGTA076 by cells in said sample after contact with said candidate compound with the expression of OGTA076 or nucleic acid encoding OGTA076 of cells in said sample before contact with said candidate compound, or with a reference level of expression.
55. 55. A method according to claim 52 or claim 53 which is a method of screening for compounds that inhibit expression of OGTA076 or nucleic acid encoding OGTA076.
56. 56. A compound obtainable by a method according to any one of claims 47 to 55.
57. 57. A compound which modulates the activity or expression of OGTA076 or nucleic acid encoding OGTA076.
58. 58. A compound according to claim 57 which inhibits the activity or expression of OGTA076 or nucleic acid encoding OGTA076.
59. 59. A compound according to any one of claims 56 to 58 for use in treating or preventing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.
60. 60. A method for treating or preventing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer which comprises administering to a subject in need thereof a therapeutically effective amount of a compound according to any one of claims 56 to 58.

61. 61. A hybridizing agent capable of hybridizing to nucleic acid encoding OGTA076.
62. 62. A hybridizing agent according to claim 61 which contains or is conjugated to a detectable label.
63. 63. A pharmaceutical composition comprising one or more hybridizing agents as defined in claim 60 or claim 61 and a pharmaceutically acceptable diluent or carrier.
64. 64. A kit containing one or more hybridizing agents according to any one of claims 61-62 or a composition according to claim 63 wherein said hybridising agent is suitable for use in treatment and/or diagnosis.
65. 65. A kit according to claim 64 further containing reagents capable of detecting and reporting the binding of said hybridizing agents to their binding partners.
66. 66. A kit containing OGTA076 and/or one or more fragments thereof wherein OGTA076 is suitable for use in treatment and/or diagnosis.
67. 67. A hybridizing agent as defined in any one of claim 61 or claim 62 for use in treatment.
68. 68. A hybridizing agent according to claim 67 wherein the treatment is for cancer.
69. 69. A hybridizing agent according to claim 68, wherein the cancer is selected from colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer.
70. 70. A method for treating or preventing colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer which comprises administering to a subject in need thereof a therapeutically effective amount of a composition comprising a hybridizing agent capable of hybridizing to nucleic acid encoding OGTA076, and a pharmaceutically acceptable diluent or carrier.
71. 71. A method of detecting, diagnosing and/or screening for or monitoring the progression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or of monitoring the effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy in a subject which comprises detecting the presence or level of OGTA076, or one or more fragments thereof, or the presence or level of nucleic acid encoding OGTA076 or the presence or level of the activity of OGTA076 or which comprises detecting a change in the level thereof in said subject.
72. 72. A method of detecting, diagnosing and/or screening for colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in a candidate subject which comprises detecting the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 in said candidate subject, in which either (a) the presence of an elevated level of OGTA076 or said one or more fragments thereof or an elevated level of nucleic acid encoding OGTA076 or the presence of an elevated level of OGTA076 activity in the candidate subject as compared with the level in a healthy subject or (b) the presence of a detectable level of OGTA076 or said one or more fragments thereof or a detectable level of nucleic acid encoding OGTA076 or the presence of a detectable level of OGTA076 activity in the candidate subject as compared with a corresponding undetectable level in a healthy subject indicates the presence of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in said subject.
73. 73. A method of monitoring the progression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in a subject or of monitoring the effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy which comprises detecting the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 in said candidate subject at a first time point and at a later time point, the presence of an elevated or lowered level of OGTA076 or said one or more fragments thereof or an elevated or lowered level of nucleic acid encoding OGTA076 or the presence of an elevated or lowered level of OGTA076 activity in the subject at the later time point as compared with the level in the subject at said first time point, indicating the progression or regression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or indicating the effect or non-effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy in said subject.
74. 74. A method according to any one of claims 70 to 73 wherein the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 is detected by analysis of a biological sample obtained from said subject.
75. 75. A method according to claim 74 which includes the step of obtaining said sample for analysis from said subject.
76. 76. A method according to claim 74 or claim 75 wherein the sample is a sample of colorectal, kidney, liver, lung, lymphoid, ovarian or pancreatic tissue.
77. 77. A method according to any one of claims 70 to 76 wherein the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 is detected quantitatively.

78. 78. A method according to claim 77 wherein the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 is detected quantitatively by means involving use of an imaging technology.

79. 79. A method according to any one of claims 70 to 77 involving use of immunohistochemistry on tissue sections in order to determine the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076, and thereby to localise colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer cells.

80. 80. A method according to any one of claims 70 to 77 wherein the presence of OGTA076, or one or more fragments thereof, or the presence of nucleic acid encoding OGTA076 or the presence of the activity of OGTA076 is detected by analysis in situ.

81. 81. A method according to any one of claims 70 to 80 wherein the presence of OGTA076 or one or more epitope-containing fragments thereof is detected directly or indirectly.

82. 82. A method according to claim 81 wherein the presence of OGTA076 or one or more fragments thereof is detected using an affinity reagent capable of specific binding to OGTA076 or one or more fragments thereof.

83. 83. A method according to claim 82 wherein the affinity reagent is an antibody.

84. 84. A method according to any one of claims 70 to 80 wherein nucleic acid encoding OGTA076 is detected.

85. 85. A method according to claim 84 wherein nucleic acid encoding OGTA076 is detected using a hybridizing agent capable of hybridizing to nucleic acid encoding OGTA076.

86. 86. A method according to any one of claims 70 to 80 wherein the activity of OGTA076 is detected.

87. 87. A method of detecting, diagnosing and/or screening for or monitoring the progression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or of monitoring the effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy in a subject which comprises detecting the presence or level of antibodies capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof or which comprises detecting a change in the level thereof in said subject.

88. 88. A method of detecting, diagnosing and/or screening for colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in a subject which comprises detecting the presence of antibodies capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof in said subject, in which (a) the presence of an elevated level of antibodies capable of immunospecific binding to OGTA076 or said one or more epitope-containing fragments thereof in said subject as compared with the level in a healthy subject or (b) the presence of a detectable level of antibodies capable of immunospecific binding to OGTA076 or said one or more epitope-containing fragments thereof in said subject as compared with a corresponding undetectable level in a healthy subject indicates the presence of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer in said subject.

89. 89. A method of monitoring the progression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or of monitoring the effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy in a subject which comprises detecting the presence of antibodies capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof in said subject at a first time point and at a later time point, the presence of an elevated or lowered level of antibodies capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof in said subject at the later time point as compared with the level in said subject at said first time point, indicating the progression or regression of colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic leukaemia), ovarian cancer or pancreatic cancer or the effect or non-effect of an anti-colorectal cancer, anti-kidney cancer, anti-liver cancer, anti-lung cancer, anti-lymphoid leukaemia (particularly chronic lymphocytic leukaemia), anti-ovarian cancer or anti-pancreatic cancer drug or therapy in said subject.

90. 90. A method according to any one of claims 77 to 89 wherein the presence of antibodies capable of immunospecific binding to OGTA076, or one or more epitope-containing fragments thereof is detected by analysis of a biological sample obtained from said subject.

91. 91. A method according to claim 90 which includes the step of obtaining said sample for analysis from said subject.

92. 92. A method according to claim 90 or claim 91 wherein the sample is a sample of colorectal, kidney, liver, lung, lymphoid, ovarian or pancreatic tissue.

93. 93. A method according to any one of claims 70 to 92 wherein the level that may be detected in the candidate subject who has colorectal cancer, kidney cancer, liver cancer, lung cancer, lymphoid leukaemia (particularly chronic lymphocytic

leukaemia), ovarian cancer or pancreatic cancer is 2 or more fold higher than the level in the healthy subject.

SEQUENCE LISTING

[0299]

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<141> 2008-02-26

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Arg His Met Ala Thr Thr Gln Asp Glu Val His Thr Lys Cys Gln		
1265	1270	1275
Lys Leu Asn Pro Lys Ser His Ile Leu Ser Ile Arg Asp Glu Lys		
1280	1285	1290
Glu Asn Asn Phe Val Leu Glu Gln Leu Leu Tyr Phe Asn Tyr Met		
1295	1300	1305
Ala Ser Trp Val Met Leu Gly Ile Thr Tyr Arg Asn Asn Ser Leu		
1310	1315	1320
Met Trp Phe Asp Lys Thr Pro Leu Ser Tyr Thr His Trp Arg Ala		
1325	1330	1335
Gly Arg Pro Thr Ile Lys Asn Glu Lys Phe Leu Ala Gly Leu Ser		
1340	1345	1350
Thr Asp Gly Phe Trp Asp Ile Gln Thr Phe Lys Val Ile Glu Glu		
1355	1360	1365
Ala Val Tyr Phe His Gln His Ser Ile Leu Ala Cys Lys Ile Glu		
1370	1375	1380

Met Val Asp Tyr Lys Glu Glu His Asn Thr Thr Leu Pro Gln Phe
1385 1390 1395

Met Pro Tyr Glu Asp Gly Ile Tyr Ser Val Ile Gln Lys Lys Val
1400 1405 1410

Thr Trp Tyr Glu Ala Leu Asn Met Cys Ser Gln Ser Gly Gly His
1415 1420 1425

Leu Ala Ser Val His Asn Gln Asn Gly Gln Leu Phe Leu Glu Asp
1430 1435 1440

Ile Val Lys Arg Asp Gly Phe Pro Leu Trp Val Gly Leu Ser Ser
1445 1450 1455

His Asp Gly Ser Glu Ser Ser Phe Glu Trp Ser Asp Gly Ser Thr
1460 1465 1470

Phe Asp Tyr Ile Pro Trp Lys Gly Gln Thr Ser Pro Gly Asn Cys
1475 1480 1485

Val Leu Leu Asp Pro Lys Gly Thr Trp Lys His Glu Lys Cys Asn
1490 1495 1500

Ser Val Lys Asp Gly Ala Ile Cys Tyr Lys Pro Thr Lys Ser Lys
1505 1510 1515

Lys Leu Ser Arg Leu Thr Tyr Ser Ser Arg Cys Pro Ala Ala Lys
1520 1525 1530

Glu Asn Gly Ser Arg Trp Ile Gln Tyr Lys Gly His Cys Tyr Lys
1535 1540 1545

Ser Asp Gln Ala Leu His Ser Phe Ser Glu Ala Lys Lys Leu Cys
1550 1555 1560

Ser Lys His Asp His Ser Ala Thr Ile Val Ser Ile Lys Asp Glu
1565 1570 1575

Asp Glu Asn Lys Phe Val Ser Arg Leu Met Arg Glu Asn Asn Asn
1580 1585 1590

Ile Thr Met Arg Val Trp Leu Gly Leu Ser Gln His Ser Val Asp
1595 1600 1605

Gln Ser Trp Ser Trp Leu Asp Gly Ser Glu Val Thr Phe Val Lys
1610 1615 1620

Trp Glu Asn Lys Ser Lys Ser Gly Val Gly Arg Cys Ser Met Leu
1625 1630 1635

Ile Ala Ser Asn Glu Thr Trp Lys Lys Val Glu Cys Glu His Gly
1640 1645 1650

Phe Gly Arg Val Val Cys Lys Val Pro Leu Gly Pro Asp Tyr Thr
1655 1660 1665

Ala Ile Ala Ile Ile Val Ala Thr Leu Ser Ile Leu Val Leu Met
1670 1675 1680

Gly Gly Leu Ile Trp Phe Leu Phe Gln Arg His Arg Leu His Leu
1685 1690 1695

Ala Gly Phe Ser Ser Val Arg Tyr Ala Gln Gly Val Asn Glu Asp
1700 1705 1710

Glu Ile Met Leu Pro Ser Phe His Asp
1715 1720

<210> 2

<211> 1873

<212> PRT

<213> Homo Sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(1873)

<223> Swissprot Accession No: 060449, Lymphocyte antigen 75 - Designated in application as OGT076 isoform b

<400> 2

Met	Arg	Thr	Gly	Trp	Ala	Thr	Pro	Arg	Arg	Pro	Ala	Gly	Leu	Leu	Met
1				5			10				15				

Leu	Leu	Phe	Trp	Phe	Phe	Asp	Leu	Ala	Glu	Pro	Ser	Gly	Arg	Ala	Ala
							20		25			30			

Asn	Asp	Pro	Phe	Thr	Ile	Val	His	Gly	Asn	Thr	Gly	Lys	Cys	Ile	Lys
							35		40		45				

Pro	Val	Tyr	Gly	Trp	Ile	Val	Ala	Asp	Asp	Cys	Asp	Glu	Thr	Glu	Asp
							50		55		60				

Lys	Leu	Trp	Lys	Trp	Val	Sex	Gln	His	Arg	Leu	Phe	His	Leu	His	Ser
							65		70		75		80		

Gln	Lys	Cys	Leu	Gly	Leu	Asp	Ile	Thr	Lys	Ser	Val	Asn	Glu	Leu	Arg
							85		90			95			

Met	Phe	Ser	Cys	Asp	Ser	Ser	Ala	Met	Leu	Trp	Trp	Lys	Cys	Glu	His
							100		105		110				

His	Ser	Leu	Tyr	Gly	Ala	Ala	Arg	Tyr	Arg	Leu	Ala	Leu	Lys	Asp	Gly
							115		120		125				

His	Gly	Thr	Ala	Ile	Ser	Asn	Ala	Ser	Asp	Val	Trp	Lys	Lys	Gly	Gly
							130		135		140				

Ser	Glu	Glu	Ser	Leu	Cys	Asp	Gln	Pro	Tyr	His	Glu	Ile	Tyr	Thr	Arg
							145		150		155		160		

Asp	Gly	Asn	Ser	Tyr	Gly	Arg	Pro	Cys	Glu	Phe	Pro	Phe	Leu	Ile	Asp
							165		170		175				

Gly	Thr	Trp	His	His	Asp	Cys	Ile	Leu	Asp	Glu	Asp	His	Ser	Gly	Pro
							180		185		190				

Trp	Cys	Ala	Thr	Thr	Leu	Asn	Tyr	Glu	Tyr	Asp	Arg	Lys	Trp	Gly	Ile
							195		200		205				

Cys	Leu	Lys	Pro	Glu	Asn	Gly	Cys	Glu	Asp	Asn	Trp	Glu	Lys	Asn	Glu
							210		215		220				

Gln	Phe	Gly	Ser	Cys	Tyr	Gln	Phe	Asn	Thr	Gln	Thr	Ala	Leu	Ser	Trp
							225		230		235		240		

Lys	Glu	Ala	Tyr	Val	Ser	Cys	Gln	Asn	Gln	Gly	Ala	Asp	Leu	Leu	Ser
							245		250		255				

Ile	Asn	Ser	Ala	Ala	Glu	Leu	Thr	Tyr	Leu	Lys	Glu	Lys	Gly	Ile	
							260		265		270				

Ala	Lys	Ile	Phe	Trp	Ile	Gly	Leu	Asn	Gln	Leu	Tyr	Ser	Ala	Arg	Gly
							275		280		285				

Trp	Glu	Trp	Ser	Asp	His	Lys	Pro	Leu	Asn	Phe	Leu	Asn	Trp	Asp	Pro
							290		295		300				

Asp	Arg	Pro	Ser	Ala	Pro	Thr	Ile	Gly	Gly	Ser	Ser	Cys	Ala	Arg	Met
							305		310		315		320		

Asp	Ala	Glu	Ser	Gly	Leu	Trp	Gln	Ser	Phe	Ser	Cys	Glu	Ala	Gln	Leu
							325		330		335				

Pro	Tyr	Val	Cys	Arg	Lys	Pro	Leu	Asn	Asn	Thr	Val	Glu	Leu	Thr	Asp
							340		345		350				

Val Trp Thr Tyr Ser Asp Thr Arg Cys Asp Ala Gly Trp Leu Pro Asn
 355 360 365

 Asn Gly Phe Cys Tyr Leu Leu Val Asn Glu Ser Asn Ser Trp Asp Lys
 370 375 380

 Ala His Ala Lys Cys Lys Ala Phe Ser Ser Asp Leu Ile Ser Ile His
 385 390 395 400

 Ser Leu Ala Asp Val Glu Val Val Thr Lys Leu His Asn Glu Asp
 405 410 415

 Ile Lys Glu Glu Val Trp Ile Gly Leu Lys Asn Ile Asn Ile Pro Thr
 420 425 430

 Leu Phe Gln Trp Ser Asp Gly Thr Glu Val Thr Leu Thr Tyr Trp Asp
 435 440 445

 Glu Asn Glu Pro Asn Val Pro Tyr Asn Lys Thr Pro Asn Cys Val Ser
 450 455 460

 Tyr Leu Gly Glu Leu Gly Gln Trp Lys Val Gln Ser Cys Glu Glu Lys
 465 470 475 480

 Leu Lys Tyr Val Cys Lys Arg Lys Gly Glu Lys Leu Asn Asp Ala Ser
 485 490 495

 Ser Asp Lys Met Cys Pro Pro Asp Glu Gly Trp Lys Arg His Gly Glu
 500 505 510

 Thr Cys Tyr Lys Ile Tyr Glu Asp Glu Val Pro Phe Gly Thr Asn Cys
 515 520 525

 Asn Leu Thr Ile Thr Ser Arg Phe Glu Gln Glu Tyr Leu Asn Asp Leu
 530 535 540

 Met Lys Lys Tyr Asp Lys Ser Leu Arg Lys Tyr Phe Trp Thr Gly Leu
 545 550 555 560

 Arg Asp Val Asp Ser Cys Gly Glu Tyr Asn Trp Ala Thr Val Gly Gly
 565 570 575

 Arg Arg Arg Ala Val Thr Phe Ser Asn Trp Asn Phe Leu Glu Pro Ala
 580 585 590

 Ser Pro Gly Gly Cys Val Ala Met Ser Thr Gly Lys Ser Val Gly Lys
 595 600 605

Trp Glu Val Lys Asp Cys Arg Ser Phe Lys Ala Leu Ser Ile Cys Lys
 610 615 620

 Lys Met Ser Gly Pro Leu Gly Pro Glu Glu Ala Ser Pro Lys Pro Asp
 625 630 635 640

 Asp Pro Cys Pro Glu Gly Trp Gln Ser Phe Pro Ala Ser Leu Ser Cys
 645 650 655

 Tyr Lys Val Phe His Ala Glu Arg Ile Val Arg Lys Arg Asn Trp Glu
 660 665 670

 Glu Ala Glu Arg Phe Cys Gln Ala Leu Gly Ala His Leu Ser Ser Phe
 675 680 685

 Ser His Val Asp Glu Ile Lys Glu Phe Leu His Phe Leu Thr Asp Gln
 690 695 700

 Phe Ser Gly Gln His Trp Leu Trp Ile Gly Leu Asn Lys Arg Ser Pro
 705 710 715 720

 Asp Leu Gln Gly Ser Trp Gln Trp Ser Asp Arg Thr Pro Val Ser Thr
 725 730 735

 Ile Ile Met Pro Asn Glu Phe Gln Gln Asp Tyr Asp Ile Arg Asp Cys
 740 745 750

 Ala Ala Val Lys Val Phe His Arg Pro Trp Arg Arg Gly Trp His Phe
 755 760 765

 Tyr Asp Asp Arg Glu Phe Ile Tyr Leu Arg Pro Phe Ala Cys Asp Thr
 770 775 780

 Lys Leu Glu Trp Val Cys Gln Ile Pro Lys Gly Arg Thr Pro Lys Thr
 785 790 795 800

 Pro Asp Trp Tyr Asn Pro Asp Arg Ala Gly Ile His Gly Pro Pro Leu
 805 810 815

 Ile Ile Glu Gly Ser Glu Tyr Trp Phe Val Ala Asp Leu His Leu Asn
 820 825 830

 Tyr Glu Glu Ala Val Leu Tyr Cys Ala Ser Asn His Ser Phe Leu Ala
 835 840 845

 Thr Ile Thr Ser Phe Val Gly Leu Lys Ala Ile Lys Asn Lys Ile Ala
 850 855 860

 Asn Ile Ser Gly Asp Gly Gln Lys Trp Trp Ile Arg Ile Ser Glu Trp

865	870	875	880
Pro Ile Asp Asp His Phe Thr Tyr Ser Arg Tyr Pro Trp His Arg Phe			
885	890	895	
Pro Val Thr Phe Gly Glu Glu Cys Leu Tyr Met Ser Ala Lys Thr Trp			
900	905	910	
Leu Ile Asp Leu Gly Lys Pro Thr Asp Cys Ser Thr Lys Leu Pro Phe			
915	920	925	
Ile Cys Glu Lys Tyr Asn Val Ser Ser Leu Glu Lys Tyr Ser Pro Asp			
930	935	940	
Ser Ala Ala Lys Val Gln Cys Ser Glu Gln Trp Ile Pro Phe Gln Asn			
945	950	955	960
Lys Cys Phe Leu Lys Ile Lys Pro Val Ser Leu Thr Phe Ser Gln Ala			
965	970	975	
Ser Asp Thr Cys His Ser Tyr Gly Gly Thr Leu Pro Ser Val Leu Ser			
980	985	990	
Gln Ile Glu Gln Asp Phe Ile Thr Ser Leu Leu Pro Asp Met Glu Ala			
995	1000	1005	
Thr Leu Trp Ile Gly Leu Arg Trp Thr Ala Tyr Glu Lys Ile Asn			
1010	1015	1020	
Lys Trp Thr Asp Asn Arg Glu Leu Thr Tyr Ser Asn Phe His Pro			
1025	1030	1035	
Leu Leu Val Ser Gly Arg Leu Arg Ile Pro Glu Asn Phe Phe Glu			
1040	1045	1050	
Glu Glu Ser Arg Tyr His Cys Ala Leu Ile Leu Asn Leu Gln Lys			
1055	1060	1065	
Ser Pro Phe Thr Gly Thr Trp Asn Phe Thr Ser Cys Ser Glu Arg			
1070	1075	1080	
His Phe Val Ser Leu Cys Gln Lys Tyr Ser Glu Val Lys Ser Arg			
1085	1090	1095	
Gln Thr Leu Gln Asn Ala Ser Glu Thr Val Lys Tyr Leu Asn Asn			
1100	1105	1110	
Leu Tyr Lys Ile Ile Pro Lys Thr Leu Thr Trp His Ser Ala Lys			
1115	1120	1125	

Arg Glu Cys Leu Lys Ser Asn Met Gln Leu Val Ser Ile Thr Asp
 1130 1135 1140

 Pro Tyr Gln Gln Ala Phe Leu Ser Val Gln Ala Leu Leu His Asn
 1145 1150 1155

 Ser Ser Leu Trp Ile Gly Leu Phe Ser Gln Asp Asp Glu Leu Asn
 1160 1165 1170

 Phe Gly Trp Ser Asp Gly Lys Arg Leu His Phe Ser Arg Trp Ala
 1175 1180 1185

 Glu Thr Asn Gly Gln Leu Glu Asp Cys Val Val Leu Asp Thr Asp
 1190 1195 1200

 Gly Phe Trp Lys Thr Val Asp Cys Asn Asp Asn Gln Pro Gly Ala
 1205 1210 1215

 Ile Cys Tyr Tyr Ser Gly Asn Glu Thr Glu Lys Glu Val Lys Pro
 1220 1225 1230

 Val Asp Ser Val Lys Cys Pro Ser Pro Val Leu Asn Thr Pro Trp
 1235 1240 1245

 Ile Pro Phe Gln Asn Cys Cys Tyr Asn Phe Ile Ile Thr Lys Asn
 1250 1255 1260

 Arg His Met Ala Thr Thr Gln Asp Glu Val His Thr Lys Cys Gln
 1265 1270 1275

 Lys Leu Asn Pro Lys Ser His Ile Leu Ser Ile Arg Asp Glu Lys
 1280 1285 1290

 Glu Asn Asn Phe Val Leu Glu Gln Leu Leu Tyr Phe Asn Tyr Met
 1295 1300 1305

 Ala Ser Trp Val Met Leu Gly Ile Thr Tyr Arg Asn Asn Ser Leu
 1310 1315 1320

 Met Trp Phe Asp Lys Thr Pro Leu Ser Tyr Thr His Trp Arg Ala
 1325 1330 1335

 Gly Arg Pro Thr Ile Lys Asn Glu Lys Phe Leu Ala Gly Leu Ser
 1340 1345 1350

 Thr Asp Gly Phe Trp Asp Ile Gln Thr Phe Lys Val Ile Glu Glu
 1355 1360 1365

Ala Val Tyr Phe His Gln His Ser Ile Leu Ala Cys Lys Ile Glu
 1370 1375 1380

 Met Val Asp Tyr Lys Glu Glu His Asn Thr Thr Leu Pro Gln Phe
 1385 1390 1395

 Met Pro Tyr Glu Asp Gly Ile Tyr Ser Val Ile Gln Lys Lys Val
 1400 1405 1410

 Thr Trp Tyr Glu Ala Leu Asn Met Cys Ser Gln Ser Gly Gly His
 1415 1420 1425

 Leu Ala Ser Val His Asn Gln Asn Gly Gln Leu Phe Leu Glu Asp
 1430 1435 1440

 Ile Val Lys Arg Asp Gly Phe Pro Leu Trp Val Gly Leu Ser Ser
 1445 1450 1455

 His Asp Gly Ser Glu Ser Ser Phe Glu Trp Ser Asp Gly Ser Thr
 1460 1465 1470

 Phe Asp Tyr Ile Pro Trp Lys Gly Gln Thr Ser Pro Gly Asn Cys
 1475 1480 1485

 Val Leu Leu Asp Pro Lys Gly Thr Trp Lys His Glu Lys Cys Asn
 1490 1495 1500

 Ser Val Lys Asp Gly Ala Ile Cys Tyr Lys Pro Thr Lys Ser Lys
 1505 1510 1515

 Lys Leu Ser Arg Leu Thr Tyr Ser Ser Arg Cys Pro Ala Ala Lys
 1520 1525 1530

 Glu Asn Gly Ser Arg Trp Ile Gln Tyr Lys Gly His Cys Tyr Lys
 1535 1540 1545

 Ser Asp Gln Ala Leu His Ser Phe Ser Glu Ala Lys Lys Leu Cys
 1550 1555 1560

 Ser Lys His Asp His Ser Ala Thr Ile Val Ser Ile Lys Asp Glu
 1565 1570 1575

 Asp Glu Asn Lys Phe Val Ser Arg Leu Met Arg Glu Asn Asn Asn
 1580 1585 1590

 Ile Thr Met Arg Val Trp Leu Gly Leu Ser Gln His Ser Val Asp
 1595 1600 1605

Gln Ser Trp Ser Trp Leu Asp Gly Ser Glu Val Thr Phe Val Lys
1610 1615 1620

Trp Glu Asn Lys Ser Lys Ser Gly Val Gly Arg Cys Ser Met Leu
1625 1630 1635

Ile Ala Ser Asn Glu Thr Trp Lys Lys Val Glu Cys Glu His Gly
1640 1645 1650

Phe Gly Arg Val Val Cys Lys Val Pro Leu Asp Cys Pro Ser Ser
1655 1660 1665

Thr Trp Ile Gln Phe Gln Asp Ser Cys Tyr Ile Phe Leu Gln Glu
1670 1675 1680

Ala Ile Lys Val Glu Ser Ile Glu Asp Val Arg Asn Gln Cys Thr
1685 1690 1695

Asp His Gly Ala Asp Met Ile Ser Ile His Asn Glu Glu Glu Asn
1700 1705 1710

Ala Phe Ile Leu Asp Thr Leu Lys Lys Gln Trp Lys Gly Pro Asp
1715 1720 1725

Asp Ile Leu Leu Gly Met Phe Tyr Asp Thr Asp Asp Ala Ser Phe
1730 1735 1740

Lys Trp Phe Asp Asn Ser Asn Met Thr Phe Asp Lys Trp Thr Asp
1745 1750 1755

Gln Asp Asp Asp Glu Asp Leu Val Asp Thr Cys Ala Phe Leu His
1760 1765 1770

Ile Lys Thr Gly Glu Trp Lys Lys Gly Asn Cys Glu Val Ser Ser
1775 1780 1785

Val Glu Gly Thr Leu Cys Lys Thr Ala Ile Pro Tyr Lys Arg Lys
1790 1795 1800

Tyr Leu Ser Asp Asn His Ile Leu Ile Ser Ala Leu Val Ile Ala
1805 1810 1815

Ser Thr Val Ile Leu Thr Val Leu Gly Ala Ile Ile Trp Phe Leu
1820 1825 1830

Tyr Lys Lys His Ser Asp Ser Arg Phe Thr Thr Val Phe Ser Thr
1835 1840 1845

Ala Pro Gln Ser Pro Tyr Asn Glu Asp Cys Val Leu Val Val Gly
1850 1855 1860

Glu Glu Asn Glu Tyr Pro Val Gln Phe Asp
1865 1870

<210> 3

<211> 1817

<212> PRT

<213> Homo Sapiens

<220>

<221> MISC_FEATURE

<222> (1).(1817)

<223> Swissprot Accession No: 060449, Lymphocyte antigen 75 - Designated in application as OGT076 isoform c

<400> 3

Met Arg Thr Gly Trp Ala Thr Pro Arg Arg Pro Ala Gly Leu Leu Met
1 5 10 15

Leu Leu Phe Trp Phe Phe Asp Leu Ala Glu Pro Ser Gly Arg Ala Ala
20 25 30

Asn Asp Pro Phe Thr Ile Val His Gly Asn Thr Gly Lys Cys Ile Lys
35 40 45

Pro Val Tyr Gly Trp Ile Val Ala Asp Asp Cys Asp Glu Thr Glu Asp
50 55 60

Lys Leu Trp Lys Trp Val Ser Gln His Arg Leu Phe His Leu His Ser
65 70 75 80

Gln Lys Cys Leu Gly Leu Asp Ile Thr Lys Ser Val Asn Glu Leu Arg
85 90 95

Met Phe Ser Cys Asp Ser Ser Ala Met Leu Trp Trp Lys Cys Glu His
100 105 110

His Ser Leu Tyr Gly Ala Ala Arg Tyr Arg Leu Ala Leu Lys Asp Gly
115 120 125

His Gly Thr Ala Ile Ser Asn Ala Ser Asp Val Trp Lys Lys Gly Gly
130 135 140

Ser Glu Glu Ser Leu Cys Asp Gln Pro Tyr His Glu Ile Tyr Thr Arg
145 150 155 160

Asp Gly Asn Ser Tyr Gly Arg Pro Cys Glu Phe Pro Phe Leu Ile Asp
165 170 175

Gly Thr Trp His His Asp Cys Ile Leu Asp Glu Asp His Ser Gly Pro
 180 185 190

 Trp Cys Ala Thr Thr Leu Asn Tyr Glu Tyr Asp Arg Lys Trp Gly Ile
 195 200 205

 Cys Leu Lys Pro Glu Asn Gly Cys Glu Asp Asn Trp Glu Lys Asn Glu
 210 215 220

 Gln Phe Gly Ser Cys Tyr Gln Phe Asn Thr Gln Thr Ala Leu Ser Trp
 225 230 235 240

 Lys Glu Ala Tyr Val Ser Cys Gln Asn Gln Gly Ala Asp Leu Leu Ser
 245 250 255

 Ile Asn Ser Ala Ala Glu Leu Thr Tyr Leu Lys Glu Lys Glu Gly Ile
 260 265 270

 Ala Lys Ile Phe Trp Ile Gly Leu Asn Gln Leu Tyr Ser Ala Arg Gly
 275 280 285

 Trp Glu Trp Ser Asp His Lys Pro Leu Asn Phe Leu Asn Trp Asp Pro
 290 295 300

 Asp Arg Pro Ser Ala Pro Thr Ile Gly Gly Ser Ser Cys Ala Arg Met
 305 310 315 320

 Asp Ala Glu Ser Gly Leu Trp Gln Ser Phe Ser Cys Glu Ala Gln Leu
 325 330 335

 Pro Tyr Val Cys Arg Lys Pro Leu Asn Asn Thr Val Glu Leu Thr Asp
 340 345 350

 Val Trp Thr Tyr Ser Asp Thr Arg Cys Asp Ala Gly Trp Leu Pro Asn
 355 360 365

 Asn Gly Phe Cys Tyr Leu Leu Val Asn Glu Ser Asn Ser Trp Asp Lys
 370 375 380

 Ala His Ala Lys Cys Lys Ala Phe Ser Ser Asp Leu Ile Ser Ile His
 385 390 395 400

 Ser Leu Ala Asp Val Glu Val Val Thr Lys Leu His Asn Glu Asp
 405 410 415

 Ile Lys Glu Glu Val Trp Ile Gly Leu Lys Asn Ile Asn Ile Pro Thr
 420 425 430

 Leu Phe Gln Trp Ser Asp Gly Thr Glu Val Thr Leu Thr Tyr Trp Asp

435	440	445
Glu Asn Glu Pro Asn Val Pro Tyr Asn Lys Thr Pro Asn Cys Val Ser		
450	455	460
Tyr Leu Gly Glu Leu Gly Gln Trp Lys Val Gln Ser Cys Glu Glu Lys		
465	470	475
Leu Lys Tyr Val Cys Lys Arg Lys Gly Glu Lys Leu Asn Asp Ala Ser		
485	490	495
Ser Asp Lys Met Cys Pro Pro Asp Glu Gly Trp Lys Arg His Gly Glu		
500	505	510
Thr Cys Tyr Lys Ile Tyr Glu Asp Glu Val Pro Phe Gly Thr Asn Cys		
515	520	525
Asn Leu Thr Ile Thr Ser Arg Phe Glu Gln Glu Tyr Leu Asn Asp Leu		
530	535	540
Met Lys Lys Tyr Asp Lys Ser Leu Arg Lys Tyr Phe Trp Thr Gly Leu		
545	550	555
Arg Asp Val Asp Ser Cys Gly Glu Tyr Asn Trp Ala Thr Val Gly Gly		
565	570	575
Arg Arg Arg Ala Val Thr Phe Ser Asn Trp Asn Phe Leu Glu Pro Ala		
580	585	590
Ser Pro Gly Gly Cys Val Ala Met Ser Thr Gly Lys Ser Val Gly Lys		
595	600	605
Trp Glu Val Lys Asp Cys Arg Ser Phe Lys Ala Leu Ser Ile Cys Lys		
610	615	620
Lys Met Ser Gly Pro Leu Gly Pro Glu Glu Ala Ser Pro Lys Pro Asp		
625	630	635
Asp Pro Cys Pro Glu Gly Trp Gln Ser Phe Pro Ala Ser Leu Ser Cys		
645	650	655
Tyr Lys Val Phe His Ala Glu Arg Ile Val Arg Lys Arg Asn Trp Glu		
660	665	670
Glu Ala Glu Arg Phe Cys Gln Ala Leu Gly Ala His Leu Ser Ser Phe		
675	680	685
Ser His Val Asp Glu Ile Lys Glu Phe Leu His Phe Leu Thr Asp Gln		
690	695	700

Phe Ser Gly Gln His Trp Leu Trp Ile Gly Leu Asn Lys Arg Ser Pro
 705 710 715 720

 Asp Leu Gln Gly Ser Trp Gln Trp Ser Asp Arg Thr Pro Val Ser Thr
 725 730 735

 Ile Ile Met Pro Asn Glu Phe Gln Gln Asp Tyr Asp Ile Arg Asp Cys
 740 745 750

 Ala Ala Val Lys Val Phe His Arg Pro Trp Arg Arg Gly Trp His Phe
 755 760 765

 Tyr Asp Asp Arg Glu Phe Ile Tyr Leu Arg Pro Phe Ala Cys Asp Thr
 770 775 780

 Lys Leu Glu Trp Val Cys Gln Ile Pro Lys Gly Arg Thr Pro Lys Thr
 785 790 795 800

 Pro Asp Trp Tyr Asn Pro Asp Arg Ala Gly Ile His Gly Pro Pro Leu
 805 810 815

 Ile Ile Glu Gly Ser Glu Tyr Trp Phe Val Ala Asp Leu His Leu Asn
 820 825 830

 Tyr Glu Glu Ala Val Leu Tyr Cys Ala Ser Asn His Ser Phe Leu Ala
 835 840 845

 Thr Ile Thr Ser Phe Val Gly Leu Lys Ala Ile Lys Asn Lys Ile Ala
 850 855 860

 Asn Ile Ser Gly Asp Gly Gln Lys Trp Trp Ile Arg Ile Ser Glu Trp
 865 870 875 880

 Pro Ile Asp Asp His Phe Thr Tyr Ser Arg Tyr Pro Trp His Arg Phe
 885 890 895

 Pro Val Thr Phe Gly Glu Glu Cys Leu Tyr Met Ser Ala Lys Thr Trp
 900 905 910

 Leu Ile Asp Leu Gly Lys Pro Thr Asp Cys Ser Thr Lys Leu Pro Phe
 915 920 925

 Ile Cys Glu Lys Tyr Asn Val Ser Ser Leu Glu Lys Tyr Ser Pro Asp
 930 935 940

 Ser Ala Ala Lys Val Gln Cys Ser Glu Gln Trp Ile Pro Phe Gln Asn
 945 950 955 960

Lys Cys Phe Leu Lys Ile Lys Pro Val Ser Leu Thr Phe Ser Gln Ala
 965 970 975

 Ser Asp Thr Cys His Ser Tyr Gly Gly Thr Leu Pro Ser Val Leu Ser
 980 985 990

 Gln Ile Glu Gln Asp Phe Ile Thr Ser Leu Leu Pro Asp Met Glu Ala
 995 1000 1005

 Thr Leu Trp Ile Gly Leu Arg Trp Thr Ala Tyr Glu Lys Ile Asn
 1010 1015 1020

 Lys Trp Thr Asp Asn Arg Glu Leu Thr Tyr Ser Asn Phe His Pro
 1025 1030 1035

 Leu Leu Val Ser Gly Arg Leu Arg Ile Pro Glu Asn Phe Phe Glu
 1040 1045 1050

 Glu Glu Ser Arg Tyr His Cys Ala Leu Ile Leu Asn Leu Gln Lys
 1055 1060 1065

 Ser Pro Phe Thr Gly Thr Trp Asn Phe Thr Ser Cys Ser Glu Arg
 1070 1075 1080

 His Phe Val Ser Leu Cys Gln Lys Tyr Ser Glu Val Lys Ser Arg
 1085 1090 1095

 Gln Thr Leu Gln Asn Ala Ser Glu Thr Val Lys Tyr Leu Asn Asn
 1100 1105 1110

 Leu Tyr Lys Ile Ile Pro Lys Thr Leu Thr Trp His Ser Ala Lys
 1115 1120 1125

 Arg Glu Cys Leu Lys Ser Asn Met Gln Leu Val Ser Ile Thr Asp
 1130 1135 1140

 Pro Tyr Gln Gln Ala Phe Leu Ser Val Gln Ala Leu Leu His Asn
 1145 1150 1155

 Ser Ser Leu Trp Ile Gly Leu Phe Ser Gln Asp Asp Glu Leu Asn
 1160 1165 1170

 Phe Gly Trp Ser Asp Gly Lys Arg Leu His Phe Ser Arg Trp Ala
 1175 1180 1185

 Glu Thr Asn Gly Gln Leu Glu Asp Cys Val Val Leu Asp Thr Asp
 1190 1195 1200

Gly Phe Trp Lys Thr Val Asp Cys Asn Asp Asn Gln Pro Gly Ala
 1205 1210 1215

 Ile Cys Tyr Tyr Ser Gly Asn Glu Thr Glu Lys Glu Val Lys Pro
 1220 1225 1230

 Val Asp Ser Val Lys Cys Pro Ser Pro Val Leu Asn Thr Pro Trp
 1235 1240 1245

 Ile Pro Phe Gln Asn Cys Cys Tyr Asn Phe Ile Ile Thr Lys Asn
 1250 1255 1260

 Arg His Met Ala Thr Thr Gln Asp Glu Val His Thr Lys Cys Gln
 1265 1270 1275

 Lys Leu Asn Pro Lys Ser His Ile Leu Ser Ile Arg Asp Glu Lys
 1280 1285 1290

 Glu Asn Asn Phe Val Leu Glu Gln Leu Leu Tyr Phe Asn Tyr Met
 1295 1300 1305

 Ala Ser Trp Val Met Leu Gly Ile Thr Tyr Arg Asn Asn Ser Leu
 1310 1315 1320

 Met Trp Phe Asp Lys Thr Pro Leu Ser Tyr Thr His Trp Arg Ala
 1325 1330 1335

 Gly Arg Pro Thr Ile Lys Asn Glu Lys Phe Leu Ala Gly Leu Ser
 1340 1345 1350

 Thr Asp Gly Phe Trp Asp Ile Gln Thr Phe Lys Val Ile Glu Glu
 1355 1360 1365

 Ala Val Tyr Phe His Gln His Ser Ile Leu Ala Cys Lys Ile Glu
 1370 1375 1380

 Met Val Asp Tyr Lys Glu Glu His Asn Thr Thr Leu Pro Gln Phe
 1385 1390 1395

 Met Pro Tyr Glu Asp Gly Ile Tyr Ser Val Ile Gln Lys Lys Val
 1400 1405 1410

 Thr Trp Tyr Glu Ala Leu Asn Met Cys Ser Gln Ser Gly Gly His
 1415 1420 1425

 Leu Ala Ser Val His Asn Gln Asn Gly Gln Leu Phe Leu Glu Asp
 1430 1435 1440

 Ile Val Lys Arg Asp Gly Phe Pro Leu Trp Val Gly Leu Ser Ser

1445	1450	1455
His Asp Gly Ser Glu Ser Ser Phe Glu Trp Ser Asp Gly Ser Thr		
1460	1465	1470
Phe Asp Tyr Ile Pro Trp Lys Gly Gln Thr Ser Pro Gly Asn Cys		
1475	1480	1485
Val Leu Leu Asp Pro Lys Gly Thr Trp Lys His Glu Lys Cys Asn		
1490	1495	1500
Ser Val Lys Asp Gly Ala Ile Cys Tyr Lys Pro Thr Lys Ser Lys		
1505	1510	1515
Lys Leu Ser Arg Leu Thr Tyr Ser Ser Arg Cys Pro Ala Ala Lys		
1520	1525	1530
Glu Asn Gly Ser Arg Trp Ile Gln Tyr Lys Gly His Cys Tyr Lys		
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PATENTKRAV

1. Antistof, der er i stand til specifikt at bindes til Lymfocytantigen 75, til anvendelse til behandling af pancreascancer, hvor antistoffet indeholder eller er konjugeret til en terapeutisk del.

5 2. Antistof til anvendelse ifølge krav 1, hvilket antistof er et isoleret monoklonalt antistof, eller en antigenbindingsdel deraf eller et antistoffragment.

3. Isoleret monoklonalt antistof til anvendelse ifølge krav 2, hvor:

(A) antistoffet er et fuldlængde antistof af en IgG1-, IgG2-, IgG3- eller IgG4-isotype;

10 (B) antistoffet er udvalgt fra gruppen bestående af et helt antistof, et antistoffragment, et kimærisk antistof, et humant antistof, et humaniseret antistof, et enkelkædet antistof, et immunkonjugat, et defucosyleret antistof og et bispecifikt antistof;

(C) fragmentet er et domæneantistof;

15 (D) antistoffet er et monoklonalt antistof, der er i stand til specifikt at bindes til lymfocytantigen 75, og som har cytotoxicitet mod Lymfocytantigen 75, der udtrykker celler i nærvær af et humant komplement; eller

(E) antistoffet er et monoklonalt antistof, der er i stand til specifikt at bindes til lymfocytantigen 75, og som har cytotoxicitet mod lymfocytantigen 75, der udtrykker celler i nærvær af humane immune effektorceller.

20

4. Antistof til anvendelse ifølge et hvilket som helst af kravene 1-3, hvor den terapeutiske del er en cytotoxisk del eller en radioaktiv isotop, eller hvor affinitetsreagenset er et antistoflægemiddelkonjugat.

25 5. Antistof til anvendelse ifølge et hvilket som helst af kravene 1-3, hvor antistoffet er et fuldt humant eller humaniseret antistof, der er i stand til at inducere antistofdirigeret celle-medieret cytotoxicitet (ADCC)

30 6. Fremgangsmåde in vitro til detektering, diagnosticering og/eller screening for eller overvågning af progressionen af pancreascancer eller overvågningen af effekten af et lægemiddel mod pancreascancer eller terapi hos en person, hvilken fremgangsmåde omfatter detektering af tilstedeværelsen eller niveauet af antistoffer, der er i stand til immunspecifikt at bindes til lymfocytantigen 75, eller et eller flere epitopholdige fragmenter deraf, eller som omfatter detektering af en ændring i niveauet deraf hos personen; eller

35 en fremgangsmåde til detektering, diagnosticering og/eller screening for pancreascancer hos en person, hvilken fremgangsmåde omfatter detektering af tilstedeværelsen af antistoffer, der er i stand til immunspecifikt at bindes til Lymfocytantigen 75, eller et eller flere epitopholdige fragmenter deraf, hos personen, hos hvilken

(a) tilstedeværelsen af et forhøjet niveau af antistoffer, der er i stand til immunspecifikt at bindes til Lymfocytantigen 75, eller det ene eller flere epitopholdige fragmenter deraf, hos personen sammenlignet med niveauet hos en rask person eller

5 (b) tilstedeværelsen af et detekterbart niveau af antistoffer, der er i stand til immunspecifikt at bindes til Lymfocytantigen 75, eller det ene eller flere epitopholdige fragmenter deraf, hos personen sammenlignet med et tilsvarende udetekterbart niveau hos en rask person indikerer tilstedeværelsen af pancreascancer hos personen.

DRAWINGS

Figure 1

OGTA076

Peptide Source: iTRAQ Colorectal cancer

OGTA076b (SEQ ID No: 2) MRTGWATPRRPAGLLMLLFWFFDLAEPSGRAANDPFTI VHGNIGKCIKPVYCWIVADDCD 60
 OGTA076c (SEQ ID No: 3) MRTGWATPRRPAGLLMLLFWFFDLAEPSGRAANDPFTI VHGNIGKCIKPVYCWIVADDCD 60
 OGTA076a (SEQ ID No: 1) MRTGWATPRRPAGLLMLLFWFFDLAEPSGRAANDPFTI VHGNIGKCIKPVYCWIVADDCD 60

 OGTA076b (SEQ ID No: 2) ETEDKLWKKWVSQHRLPHLHSQKCLGLDTIKSVNEELRMFSCDSAMLLWWKCEHHSLYGAAR 120
 OGTA076c (SEQ ID No: 3) ETEDKLWKKWVSQHRLPHLHSQKCLGLDTIKSVNEELRMFSCDSAMLLWWKCEHHSLYGAAR 120
 OGTA076a (SEQ ID No: 1) ETEDKLWKKWVSQHRLPHLHSQKCLGLDTIKSVNEELRMFSCDSAMLLWWKCEHHSLYGAAR 120

 OGTA076b (SEQ ID No: 2) YRKLRLDGHGTAISNASDWWKKGSEESLCDQPYHEIYTRDGSYGRPECFPLIDGTWH 180
 OGTA076c (SEQ ID No: 3) YRKLRLDGHGTAISNASDWWKKGSEESLCDQPYHEIYTRDGSYGRPECFPLIDGTWH 180
 OGTA076a (SEQ ID No: 1) YRKLRLDGHGTAISNASDWWKKGSEESLCDQPYHEIYTRDGSYGRPECFPLIDGTWH 180

 OGTA076b (SEQ ID No: 2) HCDCLDEDHSGPWCATITLYEYDRKKGICLKPBNGCEDDNWERNEQPGSCYQPNITQALSW 240
 OGTA076c (SEQ ID No: 3) HCDCLDEDHSGPWCATITLYEYDRKKGICLKPBNGCEDDNWERNEQPGSCYQPNITQALSW 240
 OGTA076a (SEQ ID No: 1) HCDCLDEDHSGPWCATITLYEYDRKKGICLKPBNGCEDDNWERNEQPGSCYQPNITQALSW 240

 OGTA076b (SEQ ID No: 2) KEAVYSCQNQCAALLSINSAAELTYLKEKEGKAKIWFICLNLQIYSARGEWNSDHKPLNFL 300
 OGTA076c (SEQ ID No: 3) KEAVYSCQNQCAALLSINSAAELTYLKEKEGKAKIWFICLNLQIYSARGEWNSDHKPLNFL 300
 OGTA076a (SEQ ID No: 1) KEAVYSCQNQCAALLSINSAAELTYLKEKEGKAKIWFICLNLQIYSARGEWNSDHKPLNFL 300

 OGTA076b (SEQ ID No: 2) NWDPDRPSAPTIIGGSSCARMDAESCLNQSFSCEAQLEFYCRKPLNNTVELETDWVTTYSDT 360
 OGTA076c (SEQ ID No: 3) NWDPDRPSAPTIIGGSSCARMDAESCLNQSFSCEAQLEFYCRKPLNNTVELETDWVTTYSDT 360
 OGTA076a (SEQ ID No: 1) NWDPDRPSAPTIIGGSSCARMDAESCLNQSFSCEAQLEFYCRKPLNNTVELETDWVTTYSDT 360

 OGTA076b (SEQ ID No: 2) CDAGWLPNNGPCYLLVNESENNSWDAKAKCRAFSSDLISIHSLADVEVVTKLHNEDIKEE 420
 OGTA076c (SEQ ID No: 3) CDAGWLPNNGPCYLLVNESENNSWDAKAKCRAFSSDLISIHSLADVEVVTKLHNEDIKEE 420
 OGTA076a (SEQ ID No: 1) CDAGWLPNNGPCYLLVNESENNSWDAKAKCRAFSSDLISIHSLADVEVVTKLHNEDIKEE 420

 OGTA076b (SEQ ID No: 2) VVIGLKNINIPTLFQNSDGTETVILTYWDENE PNVPYNNKTPNCVSYLCELGQWKVQSCEEK 480
 OGTA076c (SEQ ID No: 3) VVIGLKNINIPTLFQNSDGTETVILTYWDENE PNVPYNNKTPNCVSYLCELGQWKVQSCEEK 480
 OGTA076a (SEQ ID No: 1) VVIGLKNINIPTLFQNSDGTETVILTYWDENE PNVPYNNKTPNCVSYLCELGQWKVQSCEEK 480

 OGTA076b (SEQ ID No: 2) LKVYCKRKGEKLNDASSDKMCPDPCWKRKRCPTCYKLYEDEVPPFCNCNLITTSRFQEQY 540
 OGTA076c (SEQ ID No: 3) LKVYCKRKGEKLNDASSDKMCPDPCWKRKRCPTCYKLYEDEVPPFCNCNLITTSRFQEQY 540
 OGTA076a (SEQ ID No: 1) LKVYCKRKGEKLNDASSDKMCPDPCWKRKRCPTCYKLYEDEVPPFCNCNLITTSRFQEQY 540

 OGTA076b (SEQ ID No: 2) LNDLMKYDKSLRKYFTWGLRDVDSCCGYYNATVGGRRRAVTFNSNNFLEPASPGCCVAM 600
 OGTA076c (SEQ ID No: 3) LNDLMKYDKSLRKYFTWGLRDVDSCCGYYNATVGGRRRAVTFNSNNFLEPASPGCCVAM 600
 OGTA076a (SEQ ID No: 1) LNDLMKYDKSLRKYFTWGLRDVDSCCGYYNATVGGRRRAVTFNSNNFLEPASPGCCVAM 600

 OGTA076b (SEQ ID No: 2) STCKSVGKNEVKDCRSFKALSICKKMSPGLPGEPEASPKFDDPCPEGWQSFPASLSCYKVF 660
 OGTA076c (SEQ ID No: 3) STCKSVGKNEVKDCRSFKALSICKKMSPGLPGEPEASPKFDDPCPEGWQSFPASLSCYKVF 660
 OGTA076a (SEQ ID No: 1) STCKSVGKNEVKDCRSFKALSICKKMSPGLPGEPEASPKFDDPCPEGWQSFPASLSCYKVF 660

 OGTA076b (SEQ ID No: 2) HAERIVRKRNWEEAERFCQALGAHILSSFSHVDEIKEFLHFTDQFSGQHWLWIGINKRSP 720
 OGTA076c (SEQ ID No: 3) HAERIVRKRNWEEAERFCQALGAHILSSFSHVDEIKEFLHFTDQFSGQHWLWIGINKRSP 720
 OGTA076a (SEQ ID No: 1) HAERIVRKRNWEEAERFCQALGAHILSSFSHVDEIKEFLHFTDQFSGQHWLWIGINKRSP 720

 OGTA076b (SEQ ID No: 2) DLQSWQNSDRTPVSTIIMPNFQQDYDIRDCAAVKVPHRFWRNGHWFYDDREFIYLRF 780
 OGTA076c (SEQ ID No: 3) DLQSWQNSDRTPVSTIIMPNFQQDYDIRDCAAVKVPHRFWRNGHWFYDDREFIYLRF 780
 OGTA076a (SEQ ID No: 1) DLQSWQNSDRTPVSTIIMPNFQQDYDIRDCAAVKVPHRFWRNGHWFYDDREFIYLRF 780

OGTA076b (SEQ ID No: 2) ACDTKLEWVQCPKGRTPKTPDWYNPDRAGIHCPLIIEGSEYWFVADLHINYEEAVLYC 840
 OGTA076c (SEQ ID No: 3) ACDTKLEWVQCPKGRTPKTPDWYNPDRAGIHCPLIIEGSEYWFVADLHINYEEAVLYC 840
 OGTA076a (SEQ ID No: 1) ACDTKLEWVQCPKGRTPKTPDWYNPDRAGIHCPLIIEGSEYWFVADLHINYEEAVLYC 840

 OGTA076b (SEQ ID No: 2) ASNHSFLATITSFVGLKAIKNKIANISCDGQKWWIRISEWPICDDHFTYSRYPWHRPFVTF 900
 OGTA076c (SEQ ID No: 3) ASNHSFLATITSFVGLKAIKNKIANISCDGQKWWIRISEWPICDDHFTYSRYPWHRPFVTF 900
 OGTA076a (SEQ ID No: 1) ASNHSFLATITSFVGLKAIKNKIANISCDGQKWWIRISEWPICDDHFTYSRYPWHRPFVTF 900

 OGTA076b (SEQ ID No: 2) GEECLYMSAKTWILDLGKPTDCSTKLPLPICEKYNVSSLEKYSYSPDAAKVCQSEQWIPFQN 960
 OGTA076c (SEQ ID No: 3) GEECLYMSAKTWILDLGKPTDCSTKLPLPICEKYNVSSLEKYSYSPDAAKVCQSEQWIPFQN 960
 OGTA076a (SEQ ID No: 1) GEECLYMSAKTWILDLGKPTDCSTKLPLPICEKYNVSSLEKYSYSPDAAKVCQSEQWIPFQN 960

 OGTA076b (SEQ ID No: 2) KCPFLKIKPVSQASDTCHSYGGTLPSLQS1QDFITSL1LPDMEAATLWIGLRLWTAYE 1020
 OGTA076c (SEQ ID No: 3) KCPFLKIKPVSQASDTCHSYGGTLPSLQS1QDFITSL1LPDMEAATLWIGLRLWTAYE 1020
 OGTA076a (SEQ ID No: 1) KCPFLKIKPVSQASDTCHSYGGTLPSLQS1QDFITSL1LPDMEAATLWIGLRLWTAYE 1020

 OGTA076b (SEQ ID No: 2) KINKWTDNRELTYSNHPILLVSGRLRIPENFEEERSRYHCALILNLQKSPFTGTWNFTSC 1080
 OGTA076c (SEQ ID No: 3) KINKWTDNRELTYSNHPILLVSGRLRIPENFEEERSRYHCALILNLQKSPFTGTWNFTSC 1080
 OGTA076a (SEQ ID No: 1) KINKWTDNRELTYSNHPILLVSGRLRIPENFEEERSRYHCALILNLQKSPFTGTWNFTSC 1080

 OGTA076b (SEQ ID No: 2) SERHFVSLCQKYSSEVKSQRTLQNASETVKYLNNLYK3EKTILTHWSAKRECLKSNMQLVS 1140
 OGTA076c (SEQ ID No: 3) SERHFVSLCQKYSSEVKSQRTLQNASETVKYLNNLYK3EKTILTHWSAKRECLKSNMQLVS 1140
 OGTA076a (SEQ ID No: 1) SERHFVSLCQKYSSEVKSQRTLQNASETVKYLNNLYK3EKTILTHWSAKRECLKSNMQLVS 1140

 OGTA076b (SEQ ID No: 2) ITDPYQQAFLSVQALLHNSL1WIGLFSQDDELNFGWSDCKRLHFSRWAETNGQLEDCVVL 1200
 OGTA076c (SEQ ID No: 3) ITDPYQQAFLSVQALLHNSL1WIGLFSQDDELNFGWSDCKRLHFSRWAETNGQLEDCVVL 1200
 OGTA076a (SEQ ID No: 1) ITDPYQQAFLSVQALLHNSL1WIGLFSQDDELNFGWSDCKRLHFSRWAETNGQLEDCVVL 1200

 OGTA076b (SEQ ID No: 2) DTDCFWKTVDCNDNQPGAIYCYSGNETEKVKPVSVKCPSPVINTFWIPFQNCYNFII 1260
 OGTA076c (SEQ ID No: 3) DTDCFWKTVDCNDNQPGAIYCYSGNETEKVKPVSVKCPSPVINTFWIPFQNCYNFII 1260
 OGTA076a (SEQ ID No: 1) DTDCFWKTVDCNDNQPGAIYCYSGNETEKVKPVSVKCPSPVINTFWIPFQNCYNFII 1260

 OGTA076b (SEQ ID No: 2) TKNRHMMATTQDEVHTKCKQK3EPRSHISL1RDEKENNPFVLECLLYPNYMASVWMLGITYRN 1320
 OGTA076c (SEQ ID No: 3) TKNRHMMATTQDEVHTKCKQK3EPRSHISL1RDEKENNPFVLECLLYPNYMASVWMLGITYRN 1320
 OGTA076a (SEQ ID No: 1) TKNRHMMATTQDEVHTKCKQK3EPRSHISL1RDEKENNPFVLECLLYPNYMASVWMLGITYRN 1320

 OGTA076b (SEQ ID No: 2) NSLMWFDEKTP1PSYTHWRAGRPT1KNEKFLAGLST1GFWD1CTFVKVIEEAVYFHQSILAC 1380
 OGTA076c (SEQ ID No: 3) NSLMWFDEKTP1PSYTHWRAGRPT1KNEKFLAGLST1GFWD1CTFVKVIEEAVYFHQSILAC 1380
 OGTA076a (SEQ ID No: 1) NSLMWFDEKTP1PSYTHWRAGRPT1KNEKFLAGLST1GFWD1CTFVKVIEEAVYFHQSILAC 1380

 OGTA076b (SEQ ID No: 2) KIEMVDYKEEHNNTL1PQFMPYEDGIYSV1QKVTWYEAI1NMC5QSGGHILASVHNQNCQLF 1440
 OGTA076c (SEQ ID No: 3) KIEMVDYKEEHNNTL1PQFMPYEDGIYSV1QKVTWYEAI1NMC5QSGGHILASVHNQNCQLF 1440
 OGTA076a (SEQ ID No: 1) KIEMVDYKEEHNNTL1PQFMPYEDGIYSV1QKVTWYEAI1NMC5QSGGHILASVHNQNCQLF 1440

 OGTA076b (SEQ ID No: 2) LEDIVKRDGFFLWVGILSSHDGSESSFEWSDGSTFDY1PKWQGTPSPNCVLLDPKGTVKHE 1500
 OGTA076c (SEQ ID No: 3) LEDIVKRDGFFLWVGILSSHDGSESSFEWSDGSTFDY1PKWQGTPSPNCVLLDPKGTVKHE 1500
 OGTA076a (SEQ ID No: 1) LEDIVKRDGFFLWVGILSSHDGSESSFEWSDGSTFDY1PKWQGTPSPNCVLLDPKGTVKHE 1500

 OGTA076b (SEQ ID No: 2) KCNSVKDGAICYKPTKSKKLSRLTYSSRCPAAKENGSRW1QYKCHCYKSDQALHSFSEAK 1560
 OGTA076c (SEQ ID No: 3) KCNSVKDGAICYKPTKSKKLSRLTYSSRCPAAKENGSRW1QYKCHCYKSDQALHSFSEAK 1560
 OGTA076a (SEQ ID No: 1) KCNSVKDGAICYKPTKSKKLSRLTYSSRCPAAKENGSRW1QYKCHCYKSDQALHSFSEAK 1560

 OGTA076b (SEQ ID No: 2) KLCSDKHDSATIVS1K1DDEDENKFVSRIMRNN1TMRVWLGLS9HSDGWSWLDGSEVT 1620
 OGTA076c (SEQ ID No: 3) KLCSDKHDSATIVS1K1DDEDENKFVSRIMRNN1TMRVWLGLS9HSDGWSWLDGSEVT 1608
 OGTA076a (SEQ ID No: 1) KLCSDKHDSATIVS1K1DDEDENKFVSRIMRNN1TMRVWLGLS9HSDGWSWLDGSEVT 1608

 OGTA076b (SEQ ID No: 2) FVKWENKSCKSGVGRCSMLIASNETWKKVECEHGPGRVVKVPI1DCPSS1W1QFQDSCYIF 1680
 OGTA076c (SEQ ID No: 3) FVKWENKSCKSGVGRCSMLIASNETWKKVECEHGPGRVVKVPI1DCPSS1W1QFQDSCYIF 1624
 OGTA076a (SEQ ID No: 1) FVKWENKSCKSGVGRCSMLIASNETWKKVECEHGPGRVVKVPI1DCPSS1W1QFQDSCYIF 1624

OCTA076b (SEQ ID No: 2) LQEAIIKVESIEDVRNQCTDHGADMISIHNEEENAFILEDILKKQWKGPDIDLLGMFYDTDD 1740
 OCTA076c (SEQ ID No: 3) LQEAIIKVESIEDVRNQCTDHGADMISIHNEEENAFILEDILKKQWKGPDIDLLGMFYDTDD 1684
 OCTA076a (SEQ ID No: 1) VLMGGLIWFLFQRHRLLAGPSSVRYAQGVNEDETIMLYSPHD-----

OCTA076b (SEQ ID No: 2) ASPKWFEDNSNMTEFDKWTQDDDEDLVDTCAPLHIRTGEWKKGNCEVSSVEGLCKTAIPY 1800
 OCTA076c (SEQ ID No: 3) ASFKWFEDNSNMTEFDKWTQDDDEDLVDTCAPLHIRTGEWKKGNCEVSSVEGLCKTAIPY 1744
 OCTA076a (SEQ ID No: 1) -----

OCTA076b (SEQ ID No: 2) KRYKILSDNHTLISALVIASTVILITVLGAIWIFLYKKHSRFTTVFSTAPQSPYNEDCVL 1860
 OCTA076c (SEQ ID No: 3) KRYKILSDNHTLISALVIASTVILITVLGAIWIFLYKKHSRFTTVFSTAPQSPYNEDCVL 1804
 OCTA076a (SEQ ID No: 1) -----

OCTA076b (SEQ ID No: 2) VVGEENNEYPVQFD 1873
 OCTA076c (SEQ ID No: 3) VVGEENNEYPVQFD 1817
 OCTA076a (SEQ ID No: 1) -----

Mass Match Peptides (bold):

EGIAK [14]
 IIPK [28]
 LALK [35]
 LNPK [40]

Tandem Peptides (underline):

EGIAK [14]
 IIPK [28]
 LALK [35]
 LNPK [40]

Peptide Source: iTRAQ Kidney cancer

OCTA076b (SEQ ID No: 2) MRTGWATPQQPAGLILMILWFFDLAPPSGRAANDPFTIVHGNITCKC1KPVVGWIVADDCD 60
 OCTA076c (SEQ ID No: 3) MRTGWATPQQPAGLILMILWFFDLAPPSGRAANDPFTIVHGNITCKC1KPVVGWIVADDCD 60
 OCTA076a (SEQ ID No: 1) MRTGWATPQQPAGLILMILWFFDLAPPSGRAANDPFTIVHGNITCKC1KPVVGWIVADDCD 60

OCTA076b (SEQ ID No: 2) ETEDKLWKKVVSQHRLFHILSOKCLGLDITTKSVNELRMFSCDSSAMLWKKCERHHS1LYGAR 120
 OCTA076c (SEQ ID No: 3) ETEDKLWKKVVSQHRLFHILSOKCLGLDITTKSVNELRMFSCDSSAMLWKKCERHHS1LYGAR 120
 OCTA076a (SEQ ID No: 1) ETEDKLWKKVVSQHRLFHILSOKCLGLDITTKSVNELRMFSCDSSAMLWKKCERHHS1LYGAR 120

OCTA076b (SEQ ID No: 2) YPAWDKGHTAISNASDWKKGGSEESLCDQPYHEIYIRDGNSYCRPCEFPFLIDGTWH 180
 OCTA076c (SEQ ID No: 3) YPAWDKGHTAISNASDWKKGGSEESLCDQPYHEIYIRDGNSYCRPCEFPFLIDGTWH 180
 OCTA076a (SEQ ID No: 1) YPAWDKGHTAISNASDWKKGGSEESLCDQPYHEIYIRDGNSYCRPCEFPFLIDGTWH 180

OCTA076b (SEQ ID No: 2) HCDLILDEDHSGFWCATTLNYEYDRKGICLKPENCECDNWEKNEQFCGSCYQFNNTQTAWS 240
 OCTA076c (SEQ ID No: 3) HCDLILDEDHSGFWCATTLNYEYDRKGICLKPENCECDNWEKNEQFCGSCYQFNNTQTAWS 240
 OCTA076a (SEQ ID No: 1) HCDLILDEDHSGFWCATTLNYEYDRKGICLKPENCECDNWEKNEQFCGSCYQFNNTQTAWS 240

OCTA076b (SEQ ID No: 2) KFAVYSCQNGADLISINSAAEITLYKEWQFWWIGLNQLYSARGWEWSDHKPLNFL 300
 OCTA076c (SEQ ID No: 3) KFAVYSCQNGADLISINSAAEITLYKEWQFWWIGLNQLYSARGWEWSDHKPLNFL 300
 OCTA076a (SEQ ID No: 1) KFAVYSCQNGADLISINSAAEITLYKEWQFWWIGLNQLYSARGWEWSDHKPLNFL 300

OCTA076b (SEQ ID No: 2) NWDPDRPSAPTIGGSSCARMDAESGLWQSFSCAEQLPYCRKPLNNTVELTDWWTYSDTR 360
 OCTA076c (SEQ ID No: 3) NWDPDRPSAPTIGGSSCARMDAESGLWQSFSCAEQLPYCRKPLNNTVELTDWWTYSDTR 360
 OCTA076a (SEQ ID No: 1) NWDPDRPSAPTIGGSSCARMDAESGLWQSFSCAEQLPYCRKPLNNTVELTDWWTYSDTR 360

OCTA076b (SEQ ID No: 2) CDAGWLPNNGFCYLLVNVESNSWDKAHAKCKAFSSDLISIHSLADVEVVVTKLHNNEDEIKE 420
 OCTA076c (SEQ ID No: 3) CDAGWLPNNGFCYLLVNVESNSWDKAHAKCKAFSSDLISIHSLADVEVVVTKLHNNEDEIKE 420
 OCTA076a (SEQ ID No: 1) CDAGWLPNNGFCYLLVNVESNSWDKAHAKCKAFSSDLISIHSLADVEVVVTKLHNNEDEIKE 420

OGTA076b (SEQ ID No: 2) VVIGLKNINIPTLFOWSDGTEVTLTYWDENEPNVVPYNTKPNCVS1YLGQWKRQSCBEK 480
 OGTA076c (SEQ ID No: 3) VVIGLKNINIPTLFOWSDGTEVTLTYWDENEPNVVPYNTKPNCVS1YLGQWKRQSCBEK 480
 OGTA076a (SEQ ID No: 1) VVIGLKNINIPTLFOWSDGTEVTLTYWDENEPNVVPYNTKPNCVS1YLGQWKRQSCBEK 480

 OGTA076b (SEQ ID No: 2) LKVVCKRKGKEKLNDASSDKMCPD2GKRHGCTCYK1YEDEVPPGTCNCNLIT1SRFEGY 540
 OGTA076c (SEQ ID No: 3) LKVVCKRKGKEKLNDASSDKMCPD2GKRHGCTCYK1YEDEVPPGTCNCNLIT1SRFEGY 540
 OGTA076a (SEQ ID No: 1) LKVVCKRKGKEKLNDASSDKMCPD2GKRHGCTCYK1YEDEVPPGTCNCNLIT1SRFEGY 540

 OGTA076b (SEQ ID No: 2) LNDLMMKDYDKSLRKYFWTGLRDVDSGCGEYNWAVTGGRRAVTFSNNNFLEPASPGCCVAM 600
 OGTA076c (SEQ ID No: 3) LNDLMMKDYDKSLRKYFWTGLRDVDSGCGEYNWAVTGGRRAVTFSNNNFLEPASPGCCVAM 600
 OGTA076a (SEQ ID No: 1) LNDLMMKDYDKSLRKYFWTGLRDVDSGCGEYNWAVTGGRRAVTFSNNNFLEPASPGCCVAM 600

 OGTA076b (SEQ ID No: 2) STGKSVGKMEVKDCRSEKALSIKKMSGPLGPEEASPKPDCPCPEGWQSFPA5LSCYKVF 660
 OGTA076c (SEQ ID No: 3) STGKSVGKMEVKDCRSEKALSIKKMSGPLGPEEASPKPDCPCPEGWQSFPA5LSCYKVF 660
 OGTA076a (SEQ ID No: 1) STGKSVGKMEVKDCRSEKALSIKKMSGPLGPEEASPKPDCPCPEGWQSFPA5LSCYKVF 660

 OGTA076b (SEQ ID No: 2) HAR1VRKRNWEEAERFCQALGAHLSSTFSHVHD1KEFLH1T1DQFSCQHNLWIGLNKSP /20
 OGTA076c (SEQ ID No: 3) HAR1VRKRNWEEAERFCQALGAHLSSTFSHVHD1KEFLH1T1DQFSCQHNLWIGLNKSP 720
 OGTA076a (SEQ ID No: 1) HAR1VRKRNWEEAERFCQALGAHLSSTFSHVHD1KEFLH1T1DQFSCQHNLWIGLNKSP 720

 OGTA076b (SEQ ID No: 2) DLQGSWQNSDRTPVST1IMPNBPFQDQDIDRCAAVKVPHRPWRGRWHPYDREFIYLRF 780
 OGTA076c (SEQ ID No: 3) DLQGSWQNSDRTPVST1IMPNBPFQDQDIDRCAAVKVPHRPWRGRWHPYDREFIYLRF 780
 OGTA076a (SEQ ID No: 1) DLQGSWQNSDRTPVST1IMPNBPFQDQDIDRCAAVKVPHRPWRGRWHPYDREFIYLRF 780

 OGTA076b (SEQ ID No: 2) ACDTKLEWVCQ1PKGRTPKTPDWNPDRAGIHGPPLIIFGSEYWFVADLH1NYYEAVLYC 840
 OGTA076c (SEQ ID No: 3) ACDTKLEWVCQ1PKGRTPKTPDWNPDRAGIHGPPLIIFGSEYWFVADLH1NYYEAVLYC 840
 OGTA076a (SEQ ID No: 1) ACDTKLEWVCQ1PKGRTPKTPDWNPDRAGIHGPPLIIFGSEYWFVADLH1NYYEAVLYC 840

 OGTA076b (SEQ ID No: 2) ASNHSFLATITSFVGLKA1KNNKIAN1SGDGKWWIRISEWPIDCHFTYSRXPWHRPWF 900
 OGTA076c (SEQ ID No: 3) ASNHSFLATITSFVGLKA1KNNKIAN1SGDGKWWIRISEWPIDCHFTYSRXPWHRPWF 900
 OGTA076a (SEQ ID No: 1) ASNHSFLATITSFVGLKA1KNNKIAN1SGDGKWWIRISEWPIDCHFTYSRXPWHRPWF 900

 OGTA076b (SEQ ID No: 2) GEECLYMSAKTWLIDLGKPTDCSTKLPCF1CKYVNSSLKYSFDSAAKQCSQWIPFQN 960
 OGTA076c (SEQ ID No: 3) GEECLYMSAKTWLIDLGKPTDCSTKLPCF1CKYVNSSLKYSFDSAAKQCSQWIPFQN 960
 OGTA076a (SEQ ID No: 1) GEECLYMSAKTWLIDLGKPTDCSTKLPCF1CKYVNSSLKYSFDSAAKQCSQWIPFQN 960

 OGTA076b (SEQ ID No: 2) KCFLK1KPVSLTFSQASDTCHSYGG1PSVLS1QIEQDFITSSL1PMDMA1LWIGLRWTAYE 1020
 OGTA076c (SEQ ID No: 3) KCFLK1KPVSLTFSQASDTCHSYGG1PSVLS1QIEQDFITSSL1PMDMA1LWIGLRWTAYE 1020
 OGTA076a (SEQ ID No: 1) KCFLK1KPVSLTFSQASDTCHSYGG1PSVLS1QIEQDFITSSL1PMDMA1LWIGLRWTAYE 1020

 OGTA076b (SEQ ID No: 2) KINKWT0NRELITYSNFPHPLVSGRLR1PENFEEESRYHCALILN1QKSPPTGTWNFTSC 1080
 OGTA076c (SEQ ID No: 3) KINKWT0NRELITYSNFPHPLVSGRLR1PENFEEESRYHCALILN1QKSPPTGTWNFTSC 1080
 OGTA076a (SEQ ID No: 1) KINKWT0NRELITYSNFPHPLVSGRLR1PENFEEESRYHCALILN1QKSPPTGTWNFTSC 1080

 OGTA076b (SEQ ID No: 2) SERHFVSLCQKYSEVKSRQTLQNASETVEK1PKTLTWHESAKRECLKSNNQQLVS 1140
 OGTA076c (SEQ ID No: 3) SERHFVSLCQKYSEVKSRQTLQNASETVEK1PKTLTWHESAKRECLKSNNQQLVS 1140
 OGTA076a (SEQ ID No: 1) SERHFVSLCQKYSEVKSRQTLQNASETVEK1PKTLTWHESAKRECLKSNNQQLVS 1140

 OGTA076b (SEQ ID No: 2) ITDPYQQAFLSVQALLHNSLWIGLFSQDDELNFCWSDGKRLHFSWARTNGQ1EDCVVL 1200
 OGTA076c (SEQ ID No: 3) ITDPYQQAFLSVQALLHNSLWIGLFSQDDELNFCWSDGKRLHFSWARTNGQ1EDCVVL 1200
 OGTA076a (SEQ ID No: 1) ITDPYQQAFLSVQALLHNSLWIGLFSQDDELNFCWSDGKRLHFSWARTNGQ1EDCVVL 1200

 OGTA076b (SEQ ID No: 2) DTGFWHTVDCNDNQPG1CYSGNETEKEVKPVDSVKCPSPV1NTPW1PPQNCCYNII 1260
 OGTA076c (SEQ ID No: 3) DTGFWHTVDCNDNQPG1CYSGNETEKEVKPVDSVKCPSPV1NTPW1PPQNCCYNII 1260
 OGTA076a (SEQ ID No: 1) DTGFWHTVDCNDNQPG1CYSGNETEKEVKPVDSVKCPSPV1NTPW1PPQNCCYNII 1260

 OGTA076b (SEQ ID No: 2) TKNRHMATTQDEVHTKCKQ1PKPSHILSIRDEKENNPVLEQ1LYFNYMASVWMLGITYRN 1320
 OGTA076c (SEQ ID No: 3) TKNRHMATTQDEVHTKCKQ1PKPSHILSIRDEKENNPVLEQ1LYFNYMASVWMLGITYRN 1320
 OGTA076a (SEQ ID No: 1) TKNRHMATTQDEVHTKCKQ1PKPSHILSIRDEKENNPVLEQ1LYFNYMASVWMLGITYRN 1320

OGTA076b (SEQ ID No: 2) NSLMWFDKTPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDIQTFKVIEEAVYFHQHSILAC 1380
 OGTA076c (SEQ ID No: 3) NSLMWFDKTPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDIQTFKVIEEAVYFHQHSILAC 1380
 OGTA076a (SEQ ID No: 1) NSLMWFDKTPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDIQTFKVIEEAVYFHQHSILAC 1380

 OGTA076b (SEQ ID No: 2) KIEMVYKEEHNNTLQPQMPYEDGIVSVIQQKVWYEALNNCSQGGHILASVHNONGQLP 1440
 OGTA076c (SEQ ID No: 3) KIEMVYKEEHNNTLQPQMPYEDGIVSVIQQKVWYEALNNCSQGGHILASVHNONGQLP 1440
 OGTA076a (SEQ ID No: 1) KIEMVYKEEHNNTLQPQMPYEDGIVSVIQQKVWYEALNNCSQGGHILASVHNONGQLP 1440

 OGTA076b (SEQ ID No: 2) LEDIVKRQDGFLWVGLSSHDGSESSFEWSDGSTFDYIPWKQQTSPGNCVLLDPKGTVKHE 1500
 OGTA076c (SEQ ID No: 3) LEDIVKRQDGFLWVGLSSHDGSESSFEWSDGSTFDYIPWKQQTSPGNCVLLDPKGTVKHE 1500
 OGTA076a (SEQ ID No: 1) LEDIVKRQDGFLWVGLSSHDGSESSFEWSDGSTFDYIPWKQQTSPGNCVLLDPKGTVKHE 1500

 OGTA076b (SEQ ID No: 2) KCNSVKDGAICYPPKSKKLRSRILTYSSRCPAAKENGSRWICYKCHYCKSQQALHSFSEAK 1560
 OGTA076c (SEQ ID No: 3) KCNSVKDGAICYPPKSKKLRSRILTYSSRCPAAKENGSRWICYKCHYCKSQQALHSFSEAK 1560
 OGTA076a (SEQ ID No: 1) KCNSVKDGAICYPPKSKKLRSRILTYSSRCPAAKENGSRWICYKCHYCKSQQALHSFSEAK 1560

 OGTA076b (SEQ ID No: 2) KLCSDKHDHSATIVS1KDEDENKFVSRMLRMRNNNTIMRVWVGLSCHSVDQSWSLDGSEVT 1620
 OGTA076c (SEQ ID No: 2) KLCSDKHDHSATIVS1KDEDENKFVSRMLRMRNNNTIMRVWVGLSCHSVDQSWSLDGSEVT 1620
 OGTA076c (SEQ ID No: 3) KLCSDKHDHSATIVS1KDEDENKFVSRMLRMRNNNTIMRVWVGLSCHSVDQSWSLDGSEVT 1608
 OGTA076a (SEQ ID No: 1) KLCSDKHDHSATIVS1KDEDENKFVSRMLRMRNNNTIMRVWVGLSCHSVDQSWSLDGSEVT 1608

 OGTA076b (SEQ ID No: 2) FVKWENKSKSGVGRCRCSMLIASNETWKKVECEHGPGRVCKVPLDCPSSTWQFQDSCYIF 1680
 OGTA076c (SEQ ID No: 3) FVKWENKSKSGVGRCRCSMLIASNETWKKVECEHGPGRVCKVPLDCPSSTWQFQDSCYIF 1624
 OGTA076a (SEQ ID No: 1) FVKWENKSKSGVGRCRCSMLIASNETWKKVECEHGPGRVCKVPLDCPSSTWQFQDSCYIF 1624

 OGTA076b (SEQ ID No: 2) LQEAIIKVESIEDVRNQCTDHGADMISIHNNEEENAFILDILKKQWKGPDILLQMFYDTDD 1740
 OGTA076c (SEQ ID No: 3) LQEAIIKVESIEDVRNQCTDHGADMISIHNNEEENAFILDILKKQWKGPDILLQMFYDTDD 1694
 OGTA076a (SEQ ID No: 1) VIMGGLIWPLFQRHLHLAGFSSVRYAQGVNEDEIMLPSFHD-----

 OGTA076b (SEQ ID No: 2) ASFKWFDNSNMIFDKWTDQDDDEDLVDTCAPLHIKTCEWKKGNCEVSSVECTLCKTAIPY 1800
 OGTA076c (SEQ ID No: 3) ASFKWFDNSNMIFDKWTDQDDDEDLVDTCAPLHIKTCEWKKGNCEVSSVECTLCKTAIPY 1744
 OGTA076a (SEQ ID No: 1) ASFKWFDNSNMIFDKWTDQDDDEDLVDTCAPLHIKTCEWKKGNCEVSSVECTLCKTAIPY 1804

 OGTA076b (SEQ ID No: 2) VVGEENEYPVQFD 1873
 OGTA076c (SEQ ID No: 3) VVGEENEYPVQFD 1817
 OGTA076a (SEQ ID No: 1) -----

Mass Match Peptides (bold):

EGIAK [14]
 LALK [35]
 LNPK [40]
 YLNNIYK [66]

Tandem Peptides (underline):

EGIAK [14]
 LALK [35]
 LNPK [40]
 YLNNIYK [66]

Peptide Source: iTRAQ Liver cancer

OGTA076b (SEQ ID No: 2) MRTGNATPRRPAGLIMLFWFDFDLAEPSGRAANDPFTIVHGNNTGKCIKPVYGVIVADDDC 60
 OGTA076c (SEQ ID No: 3) MRTGNATPRRPAGLIMLFWFDFDLAEPSGRAANDPFTIVHGNNTGKCIKPVYGVIVADDDC 60
 OGTA076a (SEQ ID No: 1) MRTGNATPRRPAGLIMLFWFDFDLAEPSGRAANDPFTIVHGNNTGKCIKPVYGVIVADDDC 60

 OCTA076b (SEQ ID No: 2) ETEDKILWKWVSOHRLPHLHSQXCLGLDITKSVN2LRMPSCESSAMLWWKCPEEHSLYGAAR 120
 OCTA076c (SEQ ID No: 3) ETEDKILWKWVSOHRLPHLHSQXCLGLDITKSVN2LRMPSCESSAMLWWKCPEEHSLYGAAR 120
 OCTA076a (SEQ ID No: 1) ETEDKILWKWVSOHRLPHLHSQXCLGLDITKSVN2LRMPSCESSAMLWWKCPEEHSLYGAAR 120

 OCTA076b (SEQ ID No: 2) YRLAKDGHGTAISNAQDWKKKGSEB51C10PQYHEIY10RGN5YGRCEPFPFLIDGWH 180
 OCTA076c (SEQ ID No: 3) YRLAKDGHGTAISNAQDWKKKGSEB51C10PQYHEIY10RGN5YGRCEPFPFLIDGWH 180
 OCTA076a (SEQ ID No: 1) YRLAKDGHGTAISNAQDWKKKGSEB51C10PQYHEIY10RGN5YGRCEPFPFLIDGWH 180

 OCTA076b (SEQ ID No: 2) HECILDEDHSGPWCATTINBYDQRKGICLKPENGCEIDNNEKNEQFGSCYQFN1Q1ALSW 240
 OCTA076c (SEQ ID No: 3) HECILDEDHSGPWCATTINBYDQRKGICLKPENGCEIDNNEKNEQFGSCYQFN1Q1ALSW 240
 OCTA076a (SEQ ID No: 1) HECILDEDHSGPWCATTINBYDQRKGICLKPENGCEIDNNEKNEQFGSCYQFN1Q1ALSW 240

 OCTA076b (SEQ ID No: 2) KEAVVSCQNOQADLLSINSAAELTYLKEEKIKA1IFWTG1LNQLYSAGRMENSDHKPLNLF 300
 OCTA076c (SEQ ID No: 3) KEAVVSCQNOQADLLSINSAAELTYLKEEKIKA1IFWTG1LNQLYSAGRMENSDHKPLNLF 300
 OCTA076a (SEQ ID No: 1) KEAVVSCQNOQADLLSINSAAELTYLKEEKIKA1IFWTG1LNQLYSAGRMENSDHKPLNLF 300

 OCTA076b (SEQ ID No: 2) NWDPDRPSAPТИGSSCARMDAE5GLWQSFSCSEAQLPYVCRKPLNNV8LTDV8TYSOTR 360
 OCTA076c (SEQ ID No: 3) NWDPDRPSAPТИGSSCARMDAE5GLWQSFSCSEAQLPYVCRKPLNNV8LTDV8TYSOTR 360
 OCTA076a (SEQ ID No: 1) NWDPDRPSAPТИGSSCARMDAE5GLWQSFSCSEAQLPYVCRKPLNNV8LTDV8TYSOTR 360

 OCTA076b (SEQ ID No: 2) CDAGWLPPNNGFCYLLVNE3NSWCKAHAKCKAFSSDL1IS1SLADVEVVVTKLHNEDIKEE 420
 OCTA076c (SEQ ID No: 3) CDAGWLPPNNGFCYLLVNE3NSWCKAHAKCKAFSSDL1IS1SLADVEVVVTKLHNEDIKEE 420
 OCTA076a (SEQ ID No: 1) CDAGWLPPNNGFCYLLVNE3NSWCKAHAKCKAFSSDL1IS1SLADVEVVVTKLHNEDIKEE 420

 OCTA076b (SEQ ID No: 2) VV1GLRNIN1IPT1FQWQSDC1EV1LTYWDENEPVPV8Y8NKT1NCVSYLCE1CQKVQSCEEF 480
 OCTA076c (SEQ ID No: 3) VV1GLRNIN1IPT1FQWQSDC1EV1LTYWDENEPVPV8Y8NKT1NCVSYLCE1CQKVQSCEEF 480
 OCTA076a (SEQ ID No: 1) VV1GLRNIN1IPT1FQWQSDC1EV1LTYWDENEPVPV8Y8NKT1NCVSYLCE1CQKVQSCEEF 480

 OCTA076b (SEQ ID No: 2) LKVYCKRGEKELINDASSDMC1PDEGKRRHGETCYK1YEDEVPEFGTNCL1ITSRFEQEY 540
 OCTA076c (SEQ ID No: 3) LKVYCKRGEKELINDASSDMC1PDEGKRRHGETCYK1YEDEVPEFGTNCL1ITSRFEQEY 540
 OCTA076a (SEQ ID No: 1) LKVYCKRGEKELINDASSDMC1PDEGKRRHGETCYK1YEDEVPEFGTNCL1ITSRFEQEY 540

 OCTA076b (SEQ ID No: 2) LNDLMMKDYK51RKYFWTGLRVD3CGEYNWATVGRERRAVTFSNWNFILEPAS2GGCVAM 600
 OCTA076c (SEQ ID No: 3) LNDLMMKDYK51RKYFWTGLRVD3CGEYNWATVGRERRAVTFSNWNFILEPAS2GGCVAM 600
 OCTA076a (SEQ ID No: 1) LNDLMMKDYK51RKYFWTGLRVD3CGEYNWATVGRERRAVTFSNWNFILEPAS2GGCVAM 600

 OCTA076b (SEQ ID No: 2) STGKSVGKWEVKDCRSFKAL1S1KMSGP1CPEEASFKPDP1CPCPFGQSCFASLSCYKVF 660
 OCTA076c (SEQ ID No: 3) STGKSVGKWEVKDCRSFKAL1S1KMSGP1CPEEASFKPDP1CPCPFGQSCFASLSCYKVF 660
 OCTA076a (SEQ ID No: 1) STGKSVGKWEVKDCRSFKAL1S1KMSGP1CPEEASFKPDP1CPCPFGQSCFASLSCYKVF 660

 OCTA076b (SEQ ID No: 2) HAERIVKRKNWEEAERPFCQALCAHLSSEFSHVDE1KEF1HPLT1CQFSQH1W1G1LNKKSP 720
 OCTA076c (SEQ ID No: 3) HAERIVKRKNWEEAERPFCQALCAHLSSEFSHVDE1KEF1HPLT1CQFSQH1W1G1LNKKSP 720
 OCTA076a (SEQ ID No: 1) HAERIVKRKNWEEAERPFCQALCAHLSSEFSHVDE1KEF1HPLT1CQFSQH1W1G1LNKKSP 720

 OCTA076b (SEQ ID No: 2) DLQGSWQMSDRTPVSTI1MPNNEFQDQYD1RDCAAVVFHWPWRRGHWYDREFTYLRPF 780
 OCTA076c (SEQ ID No: 3) DLQGSWQMSDRTPVSTI1MPNNEFQDQYD1RDCAAVVFHWPWRRGHWYDREFTYLRPF 780
 OCTA076a (SEQ ID No: 1) DLQGSWQMSDRTPVSTI1MPNNEFQDQYD1RDCAAVVFHWPWRRGHWYDREFTYLRPF 780

 OCTA076b (SEQ ID No: 2) ACDT1KLEWVQC1PKGRTPKTPDWYNPDRAG1IHGPPLI1EGSEYWFVADLH1N1YEA1LYC 840
 OCTA076c (SEQ ID No: 3) ACDT1KLEWVQC1PKGRTPKTPDWYNPDRAG1IHGPPLI1EGSEYWFVADLH1N1YEA1LYC 840
 OCTA076a (SEQ ID No: 1) ACDT1KLEWVQC1PKGRTPKTPDWYNPDRAG1IHGPPLI1EGSEYWFVADLH1N1YEA1LYC 840

 OCTA076b (SEQ ID No: 2) ASNHSP1AT1TSFVCLKA1K1KIAN1SCDCQKWN1R1SEWP1D1DHTY8RYPWHRFPVTF 900
 OCTA076c (SEQ ID No: 3) ASNHSP1AT1TSFVCLKA1K1KIAN1SDGQKWN1R1SEWP1D1DHTY8RYPWHRFPVTF 900
 OCTA076a (SEQ ID No: 1) ASNHSP1AT1TSFVCLKA1K1KIAN1SDGQKWN1R1SEWP1D1DHTY8RYPWHRFPVTF 900

 OCTA076b (SEQ ID No: 2) GEECLYMSAKTWN1L1LGKPTDCSTKLPPF1CEKYNVSSLEKYS1PDSAAKVQCSEQWIPFQN 960
 OCTA076c (SEQ ID No: 3) GEECLYMSAKTWN1L1LGKPTDCSTKLPPF1CEKYNVSSLEKYS1PDSAAKVQCSEQWIPFQN 960
 OCTA076a (SEQ ID No: 1) GEECLYMSAKTWN1L1LGKPTDCSTKLPPF1CEKYNVSSLEKYS1PDSAAKVQCSEQWIPFQN 960

 OCTA076b (SEQ ID No: 2) KCFPLIKPVSLSLPSO&STCHSYGGTILPSVLSLGIODRITSLLPDMRATLWIGLRWTAYE 1020
 OCTA076c (SEQ ID No: 3) KCFPLIKPVSLSLPSO&STCHSYGGTILPSVLSLGIODRITSLLPDMRATLWIGLRWTAYE 1020
 OCTA076a (SEQ ID No: 1) KCFPLIKPVSLSLPSO&STCHSYGGTILPSVLSLGIODRITSLLPDMRATLWIGLRWTAYE 1020

 OCTA076b (SEQ ID No: 2) KINKWTDNEELTY3NFHPLLV3G3RIR1PENPFFEE3SRVHCALIINLQKSPTFTGWNFTSC 1083
 OCTA076c (SEQ ID No: 3) KINKWTDNEELTY3NFHPLLV3G3RIR1PENPFFEE3SRVHCALIINLQKSPTFTGWNFTSC 1083
 OCTA076a (SEQ ID No: 1) KINKWTDNEELTY3NFHPLLV3G3RIR1PENPFFEE3SRVHCALIINLQKSPTFTGWNFTSC 1083

 OCTA076b (SEQ ID No: 2) SERHFVSLCQKYSEVKSRSQTLQNASETV~~X~~1140
 OCTA076c (SEQ ID No: 3) SERHFVSLCQKYSEVKSRSQTLQNASETV~~X~~1140
 OCTA076a (SEQ ID No: 1) SERHFVSLCQKYSEVKSRSQTLQNASETV~~X~~1140

 OCTA076b (SEQ ID No: 2) ITDPYQQAELSVQALLHNNSSLWIGLFSQDDELFNFGWSDCRLLHFSRWAETNGQLEDGVUL 1200
 OCTA076c (SEQ ID No: 3) ITDPYQQAELSVQALLHNNSSLWIGLFSQDDELFNFGWSDCRLLHFSRWAETNGQLEDGVUL 1200
 OCTA076a (SEQ ID No: 1) ITDPYQQAELSVQALLHNNSSLWIGLFSQDDELFNFGWSDCRLLHFSRWAETNGQLEDGVUL 1200

 OCTA076b (SEQ ID No: 2) DTDCFWKTVDCNINOPGACIYCYSGNTEKEVVKPVDVKCPSVPLNTW1PFCNCYCNFII 1260
 OCTA076c (SEQ ID No: 3) DTDCFWKTVDCNINOPGACIYCYSGNTEKEVVKPVDVKCPSVPLNTW1PFCNCYCNFII 1260
 OCTA076a (SEQ ID No: 1) DTDCFWKTVDCNINOPGACIYCYSGNTEKEVVKPVDVKCPSVPLNTW1PFCNCYCNFII 1260

 OCTA076b (SEQ ID No: 2) TKNRHMMATTOBEVH~~X~~1320
 OCTA076c (SEQ ID No: 3) TKNRHMMATTOBEVH~~X~~1320
 OCTA076a (SEQ ID No: 1) TKNRHMMATTOBEVH~~X~~1320

 OCTA076b (SEQ ID No: 2) NSLMWFDKTPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDICLCTFKVIEAVYFHQSILAC 1380
 OCTA076c (SEQ ID No: 3) NSLMWFDKTPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDICLCTFKVIEAVYFHQSILAC 1380
 OCTA076a (SEQ ID No: 1) NSLMWFDKTPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDICLCTFKVIEAVYFHQSILAC 1380

 OCTA076b (SEQ ID No: 2) KIENVYDKEEINNTLPOFMPYEDGIYSVICKVTKYFEALINCSQSGGHLASVHNQNGOLF 1440
 OCTA076c (SEQ ID No: 3) KIENVYDKEEINNTLPOFMPYEDGIYSVICKVTKYFEALINCSQSGGHLASVHNQNGOLF 1440
 OCTA076a (SEQ ID No: 1) KIENVYDKEEINNTLPOFMPYEDGIYSVICKVTKYFEALINCSQSGGHLASVHNQNGOLF 1440

 OCTA076b (SEQ ID No: 2) LEDIVKRDGFPILWGLSSHDGSESSFEWSDGSEFTDYIPWKGQTSPGNCVLLDPKGTVWHE 1500
 OCTA076c (SEQ ID No: 3) LEDIVKRDGFPILWGLSSHDGSESSFEWSDGSEFTDYIPWKGQTSPGNCVLLDPKGTVWHE 1500
 OCTA076a (SEQ ID No: 1) LEDIVKRDGFPILWGLSSHDGSESSFEWSDGSEFTDYIPWKGQTSPGNCVLLDPKGTVWHE 1500

 OCTA076b (SEQ ID No: 2) KCNSVKDGAICYKPTKSKLRSR1TYSRCPAAKEENGSRW1QYKHCYKSDQALHSFSEAK 1560
 OCTA076c (SEQ ID No: 3) KCNSVKDGAICYKPTKSKLRSR1TYSRCPAAKEENGSRW1QYKHCYKSDQALHSFSEAK 1560
 OCTA076a (SEQ ID No: 1) KCNSVKDGAICYKPTKSKLRSR1TYSRCPAAKEENGSRW1QYKHCYKSDQALHSFSEAK 1560

 OCTA076b (SEQ ID No: 2) KLC SKHDESATIVS1KDEDENKFVSLRMLRENNNT1TMRW1GLSCHSVDSWSMLDGSSEVT 1620
 OCTA076c (SEQ ID No: 3) KLC SKHDESATIVS1KDEDENKFVSLRMLRENNNT1TMRW1GLSCHSOHSE----- 1608
 OCTA076a (SEQ ID No: 1) KLC SKHDESATIVS1KDEDENKFVSLRMLRENNNT1TMRW1GLSCHSVDSWSMLDGSSEVT 1608

 OCTA076b (SEQ ID No: 2) FVKWENKSKCGVGRCSMLIASN2TKVCEHGFGRVVCKV1DCPSTTW1QFQDSCYIF 1680
 OCTA076c (SEQ ID No: 3) FVKWENKSKCGVGRCSMLIASN2TKVCEHGFGRVVCKV1DCPSTTW1QFQDSCYIF 1624
 OCTA076a (SEQ ID No: 1) FVKWENKSKCGVGRCSMLIASN2TKVCEHGFGRVVCKV1DCPSTTW1QFQDSCYIF 1624

 OCTA076b (SEQ ID No: 2) LQEAIKIVSIEDVRNQCTTHGADMISIHNEEENAFILDILKKQWKGPDILLGMFYDID 1740
 OCTA076c (SEQ ID No: 3) LQEAIKIVSIEDVRNQCTTHGADMISIHNEEENAFILDILKKQWKGPDILLGMFYDID 1684
 OCTA076a (SEQ ID No: 1) VLMGGLINFLFQRHLHLAGFSSVYRQGVNEDEIMLPSPH-----

 OCTA076b (SEQ ID No: 2) ASFKWEDNSNNTFDKWTDQDDEDLVDTCAFLHIKTCEAKKNCCEVSSVECTLCKTAIPY 1800
 OCTA076c (SEQ ID No: 3) ASFKWEDNSNNTFDKWTDQDDEDLVDTCAFLHIKTCEAKKNCCEVSSVECTLCKTAIPY 1744
 OCTA076a (SEQ ID No: 1) -----

 OCTA076b (SEQ ID No: 2) KRKYLSDNHILISALVIASIASTVLTIVGAI1WFLYKKHSDSRFTTVPSTAPQSPYNEDCWL 1860
 OCTA076c (SEQ ID No: 3) KRKYLSDNHILISALVIASIASTVLTIVGAI1WFLYKKHSDSRFTTVPSTAPQSPYNEDCWL 1804
 OCTA076a (SEQ ID No: 1) -----

OGTA076b (SEQ ID No: 2) VVCEENENYPVQFD 1873
 OGTA076c (SEQ ID No: 3) VVCEENENYPVQFD 1817
 OGTA076a (SEQ ID No: 1) -----

Mass Match Peptides (bold):
YLNLYK [66]

Tandem Peptides (underline):
YLNLYK [66]

Peptide Source: iTRAQ Non-small cell lung cancer

OGTA076b (SEQ ID No: 2) MRTGKATPRRPAGLMLLWFFDLAETSGRAANDPFTIVHGNNTGKCIKPVYGNIVADDCC 60
 OGTA076c (SEQ ID No: 3) MRTGKATPRRPAGLMLLWFFDLAETSGRAANDPFTIVHGNNTGKCIKPVYGNIVADDCC 60
 OGTA076a (SEQ ID No: 1) MRTGKATPRRPAGLMLLWFFDLAETSGRAANDPFTIVHGNNTGKCIKPVYGNIVADDCC 60

 OGTA076b (SEQ ID No: 2) PTEDKLWKV5QHRLFHHSQKCLG1LITKSVNEIRMPSCCSSAMLWKCCEHHSLYGAR 120
 OGTA076c (SEQ ID No: 3) PTEDKLWKV5QHRLFHHSQKCLG1LITKSVNEIRMPSCCSSAMLWKCCEHHSLYGAR 120
 OGTA076a (SEQ ID No: 1) PTEDKLWKV5QHRLFHHSQKCLG1LITKSVNEIRMPSCCSSAMLWKCCEHHSLYGAR 120

 OGTA076b (SEQ ID No: 2) YR~~LL~~EDGHTGTAISASDWKKGSEESLCPQPYHEIYIRDGNSYGRCPCEFFPLIDGTW 180
 OGTA076c (SEQ ID No: 3) YR~~LL~~EDGHTGTAISASDWKKGSEESLCPQPYHEIYIRDGNSYGRCPCEFFPLIDGTW 180
 OGTA076a (SEQ ID No: 1) YR~~LL~~EDGHTGTAISASDWKKGSEESLCPQPYHEIYIRDGNSYGRCPCEFFPLIDGTW 180

 OGTA076b (SEQ ID No: 2) HDC1LDEDHSCPWCATTLNVEYCRKGWICLKPENGCEDNWEKNCQFQNSYGRCPCEFFPLIDGTW 240
 OGTA076c (SEQ ID No: 3) HDC1LDEDHSCPWCATTLNVEYCRKGWICLKPENGCEDNWEKNCQFQNSYGRCPCEFFPLIDGTW 240
 OGTA076a (SEQ ID No: 1) HDC1LDEDHSCPWCATTLNVEYCRKGWICLKPENGCEDNWEKNCQFQNSYGRCPCEFFPLIDGTW 240

 OGTA076b (SEQ ID No: 2) KEAVSCQMQGADLISINSAAEITYLKEKEGIAK1TWWG1LNQLYSARGWWSDHKPLNEL 300
 OGTA076c (SEQ ID No: 3) KEAVSCQMQGADLISINSAAEITYLKEKEGIAK1TWWG1LNQLYSARGWWSDHKPLNEL 300
 OGTA076a (SEQ ID No: 1) KEAVSCQMQGADLISINSAAEITYLKEKEGIAK1TWWG1LNQLYSARGWWSDHKPLNEL 300

 OGTA076b (SEQ ID No: 2) NWDDDRPSAPТИGGSСААРМДАЕССЛWQSFСЕАQIЛPYVСRКРPLNNTVELTWVWTTSDTR 360
 OGTA076c (SEQ ID No: 3) NWDDDRPSAPТИGGSСААРМДАЕССЛWQSFСЕАQIЛPYVСRКРPLNNTVELTWVWTTSDTR 360
 OGTA076a (SEQ ID No: 1) NWDDDRPSAPТИGGSСААРМДАЕССЛWQSFСЕАQIЛPYVСRКРPLNNTVELTWVWTTSDTR 360

 OGTA076b (SEQ ID No: 2) CDAGWLPNNGFCYLLVNESNSDKAIAHKCKAFSSDLI5IHS1AD0VEVVTKLHNEDIKEE 420
 OGTA076c (SEQ ID No: 3) CDAGWLPNNGFCYLLVNESNSDKAIAHKCKAFSSDLI5IHS1AD0VEVVTKLHNEDIKEE 420
 OGTA076a (SEQ ID No: 1) CDAGWLPNNGFCYLLVNESNSDKAIAHKCKAFSSDLI5IHS1AD0VEVVTKLHNEDIKEE 420

 OGTA076b (SEQ ID No: 2) VVIGLKRNNIIP1LFWQSDGTEVLTWVDBNEPWPVYPKTNCVSYLGELGQWVQSCCEK 480
 OGTA076c (SEQ ID No: 3) VVIGLKRNNIIP1LFWQSDGTEVLTWVDBNEPWPVYPKTNCVSYLGELGQWVQSCCEK 480
 OGTA076a (SEQ ID No: 1) VVIGLKRNNIIP1LFWQSDGTEVLTWVDBNEPWPVYPKTNCVSYLGELGQWVQSCCEK 480

 OGTA076b (SEQ ID No: 2) LKYVCKRKGEKLNDASSDKMCPDEGWKRHGETCYK1YIYEDEV/2EGTNCNLTTSRFPEQY 540
 OGTA076c (SEQ ID No: 3) LKYVCKRKGEKLNDASSDKMCPDEGWKRHGETCYK1YIYEDEV/2EGTNCNLTTSRFPEQY 540
 OGTA076a (SEQ ID No: 1) LKYVCKRKGEKLNDASSDKMCPDEGWKRHGETCYK1YIYEDEV/2EGTNCNLTTSRFPEQY 540

 OGTA076b (SEQ ID No: 2) LNDLMMKDYDSLRKYFWTGLRWDVSDCGEYNWAWTVGRRRAVTFSNNWNELEPASPGCCVAM 600
 OGTA076c (SEQ ID No: 3) LNDLMMKDYDSLRKYFWTGLRWDVSDCGEYNWAWTVGRRRAVTFSNNWNELEPASPGCCVAM 600
 OGTA076a (SEQ ID No: 1) LNDLMMKDYDSLRKYFWTGLRWDVSDCGEYNWAWTVGRRRAVTFSNNWNELEPASPGCCVAM 600

 OGTA076b (SEQ ID No: 2) STGKSVGRW2VKDCRSFKL51CKKMSGPGLPPEASPKEDPCPGWQSEPPASLSCYKVF 660
 OGTA076c (SEQ ID No: 3) STGKSVGRW2VKDCRSFKL51CKKMSGPGLPPEASPKEDPCPGWQSEPPASLSCYKVF 660
 OGTA076a (SEQ ID No: 1) STGKSVGRW2VKDCRSFKL51CKKMSGPGLPPEASPKDCPCPGWQSEPPASLSCYKVF 660

OGTA076b (SEQ ID No: 2) KLCSKHDHSATIVSIKDEDENKFVSRILMRENNNIIMRVWLGLSQHSVDQSWLWGSEVT 1620
 OGTA076c (SEQ ID No: 3) KLCSKHDHSATIVSIKDEDENKFVSRILMRENNNIIMRVWLGLSQHSVD----- 1608
 OGTA076a (SEQ ID No: 1) KLCSKHDHSATIVSIKDEDENKFVSRILMRENNNIIMRVWLGLSQHSVDQSWLWGSEVT 1608

 OGTA076b (SEQ ID No: 2) FVKWENKSKSGVGRCMSLIAASNETWKKVECEHGFGRVVCKVPLDCPSTWIQFQDSCYIF 1680
 OGTA076c (SEQ ID No: 3) -----CPSSSTWIQFQDSCYIF 1624
 OGTA076a (SEQ ID No: 1) FVKWENKSKSGVGRCMSLIAASNETWKKVECEHGFGRVVCKVPLGFDITAIAIIVATLSIL

 OGTA076b (SEQ ID No: 2) IQEAIKVESIEDVNRNQCTDHGADMISIHNNEEENAFILDILKKQWKGPDDILLGMFYDID 1740
 OGTA076c (SEQ ID No: 3) IQEAIKVESIEDVNRNQCTDHGADMISIHNNEEENAFILDILKKQWKGPDDILLGMFYDID 1684
 OGTA076a (SEQ ID No: 1) VLMGGLIWFQRLRHLLAGFSSVRYAQGVNEDEIMLPSFH-----

 OGTA076b (SEQ ID No: 2) KRKYLSNDNHILISALVIASIASTVILTVLGAIIMFLYKKHSDSRFTTVFSTAPQSPYNEDCVL 1860
 OGTA076c (SEQ ID No: 3) KRKYLSNDNHILISALVIASIASTVILTVLGAIIMFLYKKHSDSRFTTVFSTAPQSPYNEDCVL 1804
 OGTA076a (SEQ ID No: 1) -----

 OGTA076b (SEQ ID No: 2) VVGEENEYFPVQFD 1873
 OGTA076c (SEQ ID No: 3) VVGEENEYFPVQFD 1817
 OGTA076a (SEQ ID No: 1) -----

Mass Match Peptides (bold):

IIPK [28]
 LALK [35]
 YLNNLYK [66]

Tandem Peptides (underline):

IIPK [28]
 LALK [35]
 YLNNLYK [66]

Peptide Source: iTRAQ Small cell lung cancer

OGTA076b (SEQ ID No: 2) MRTGWATPPIPAPCLMLLPFWFFDIAEPSCRAANDPFTIVHNCNTCKCIKPVCYCNIVADDCD 60
 OGTA076c (SEQ ID No: 3) MRTGWATPPIPAPCLMLLPFWFFDIAEPSCRAANDPFTIVHNCNTCKCIKPVCYCNIVADDCD 60
 OGTA076a (SEQ ID No: 1) MRTGWATPPIPAPCLMLLPFWFFDIAEPSCRAANDPFTIVHNCNTCKCIKPVCYCNIVADDCD 60

 OGTA076b (SEQ ID No: 2) ETEDKLWKKWVSQHRLIFHLSQKCLGLDITKSVNELRMPSCCSSAMLWWRKCPHHSLYGAAR 120
 OGTA076c (SEQ ID No: 3) ETEDKLWKKWVSQHRLIFHLSQKCLGLDITKSVNELRMPSCCSSAMLWWRKCEHHSLYGAAR 120
 OGTA076a (SEQ ID No: 1) ETEDKLWKKWVSQHRLIFHLSQKCLGLDITKSVNELRMPSCCSSAMLWWRKCEHHSLYGAAR 120

 OGTA076b (SEQ ID No: 2) YRLALKDGHGTAISNAISWVKKGSEESLCDQPYHEIYTRDCNSYGRPCCEPFLIDGTVH 180
 OGTA076c (SEQ ID No: 3) YRLALKDGHGTAISNAISWVKKGSEESLCDQPYHEIYTRDCNSYGRPCCEPFLIDGTVH 180
 OGTA076a (SEQ ID No: 1) YRLALKDGHGTAISNAISWVKKGSEESLCDQPYHEIYTRDCNSYGRPCCEPFLIDGTVH 180

 OGTA076b (SEQ ID No: 2) HDCILDEDHSGPWCATTLYEYDRKKGICLKPENGCDDNWEKNEQFGSCYQFNTQTAWS 240
 OGTA076c (SEQ ID No: 3) HDCILDEDHSGPWCATTLYEYDRKKGICLKPENGCDDNWEKNEQFGSCYQFNTQTAWS 240
 OGTA076a (SEQ ID No: 1) HDCILDEDHSGPWCATTLYEYDRKKGICLKPENGCDDNWEKNEQFGSCYQFNTQTAWS 240

 OGTA076b (SEQ ID No: 2) KEAYVSCQNGQADLLSINSAAEITLKEKEGIAKIFWICLNLQIYSARGNEWSDHKPLNPL 300
 OGTA076c (SEQ ID No: 3) KEAYVSCQNGQADLLSINSAAEITLKEKEGIAKIFWICLNLQIYSARGNEWSDHKPLNPL 300
 OGTA076a (SEQ ID No: 1) KEAYVSCQNGQADLLSINSAAEITLKEKEGIAKIFWICLNLQIYSARGNEWSDHKPLNPL 300

OGTA076b (SEQ ID No: 2) NWDPDRPSAPTIQGSSCARMDAESGLWQSFCBAGLPHYVCRKPLNNNTVELTDWWTYSSTR 360
 OGTA076c (SEQ ID No: 3) NWDPDRPSAPTIQGSSCARMDAESGLWQSFCBAGLPHYVCRKPLNNNTVELTDWWTYSSTR 360
 OGTA076a (SEQ ID No: 1) NWDPDRPSAPTIQGSSCARMDAESGLWQSFCBAGLPHYVCRKPLNNNTVELTDWWTYSSTR 360

 OGTA076b (SEQ ID No: 2) CDAGWLPNNGFCYLLVNEGSNSWDKAHAKCKAFSSDLISIHSALADVEVVVTKLHNEDIKEE 420
 OGTA076c (SEQ ID No: 3) CDAGWLPNNGFCYLLVNEGSNSWDKAHAKCKAFSSDLISIHSALADVEVVVTKLHNEDIKEE 420
 OGTA076a (SEQ ID No: 1) CDAGWLPNNGFCYLLVNEGSNSWDKAHAKCKAFSSDLISIHSALADVEVVVTKLHNEDIKEE 420

 OGTA076b (SEQ ID No: 2) VVIGLKNINIPITLFWSDGTEVILTYDNEPEPNVYNKIPNQCVSYLGEIGCQMKVQSCERK 480
 OGTA076c (SEQ ID No: 3) VVIGLKNINIPITLFWSDGTEVILTYDNEPEPNVYNKIPNQCVSYLGEIGCQMKVQSCERK 480
 OGTA076a (SEQ ID No: 1) VVIGLKNINIPITLFWSDGTEVILTYDNEPEPNVYNKIPNQCVSYLGEIGCQMKVQSCERK 480

 OGTA076b (SEQ ID No: 2) LKVKCRKRGEEKLNDAKSDKMCPPDEGWKEBREGTCYKIXEDEVEFGTNCNLTTSRFEQERY 540
 OGTA076c (SEQ ID No: 3) LKVKCRKRGEEKLNDAKSDKMCPPDEGWKEBREGTCYKIXEDEVEFGTNCNLTTSRFEQERY 540
 OGTA076a (SEQ ID No: 1) LKVKCRKRGEEKLNDAKSDKMCPPDEGWKEBREGTCYKIXEDEVEFGTNCNLTTSRFEQERY 540

 OGTA076b (SEQ ID No: 2) LNLMKRYDKSLRKYFTGLRDVSCGEYNATVGRRRAVTSNNNNFLEPASPGCVM 600
 OGTA076c (SEQ ID No: 3) LNLMKRYDKSLRKYFTGLRDVSCGEYNATVGRRRAVTSNNNNFLEPASPGCVM 600
 OGTA076a (SEQ ID No: 1) LNLMKRYDKSLRKYFTGLRDVSCGEYNATVGRRRAVTSNNNNFLEPASPGCVM 600

 OGTA076b (SEQ ID No: 2) STGKSVGKWEVDCRSFKALSIKKMSGFLGPEEEASPKPDDPCPEGQSFPAISLSCYKVF 660
 OGTA076c (SEQ ID No: 3) STGKSVGKWEVDCRSFKALSIKKMSGFLGPEEEASPKPDDPCPEGQSFPAISLSCYKVF 660
 OGTA076a (SEQ ID No: 1) STGKSVGKWEVDCRSFKALSIKKMSGFLGPEEEASPKPDDPCPEGQSFPAISLSCYKVF 660

 OGTA076b (SEQ ID No: 2) HARIRVRKRNWEEAERFCQALGAHLSFFSHVDEIKEPLHIFTDQFSGQHWWLIGIINKRSP 720
 OGTA076c (SEQ ID No: 3) HARIRVRKRNWEEAERFCQALGAHLSFFSHVDEIKEPLHIFTDQFSGQHWWLIGIINKRSP 720
 OGTA076a (SEQ ID No: 1) HARIRVRKRNWEEAERFCQALGAHLSFFSHVDEIKEPLHIFTDQFSGQHWWLIGIINKRSP 720

 OGTA076b (SEQ ID No: 2) DLQGSWQNSDRTPVSTIIMPNEFQDQDYDIRDCAVKVFHRPWRGWHFYDDREPIIYLRRPP 780
 OGTA076c (SEQ ID No: 3) DLQGSWQNSDRTPVSTIIMPNEFQDQDYDIRDCAVKVFHRPWRGWHFYDDREPIIYLRRPP 780
 OGTA076a (SEQ ID No: 1) DLQGSWQNSDRTPVSTIIMPNEFQDQDYDIRDCAVKVFHRPWRGWHFYDDREPIIYLRRPP 780

 OGTA076b (SEQ ID No: 2) ACTDKLEWVCQ1PKGRTPK1PDWNPDRAGIHPPL11EGSEYHWFADLHINYEEAVLYC 840
 OGTA076c (SEQ ID No: 3) ACTDKLEWVCQ1PKGRTPK1PDWNPDRAGIHPPL11EGSEYHWFADLHINYEEAVLYC 840
 OGTA076a (SEQ ID No: 1) ACTDKLEWVCQ1PKGRTPK1PDWNPDRAGIHPPL11EGSEYHWFADLHINYEEAVLYC 840

 OGTA076b (SEQ ID No: 2) ASHNSFLATITSFVGLRAIKNKHIANISGDGKWWIRISEWPIDDIHTYSRYWHRFPVTF 900
 OGTA076c (SEQ ID No: 3) ASHNSFLATITSFVGLRAIKNKHIANISGDGKWWIRISEWPIDDIHTYSRYWHRFPVTF 900
 OGTA076a (SEQ ID No: 1) ASHNSFLATITSFVGLRAIKNKHIANISGDGKWWIRISEWPIDDIHTYSRYWHRFPVTF 900

 OGTA076b (SEQ ID No: 2) GEECLYMSAKTNLIDLCKPTDCSTKLPPFCKVNVSSELEYKSFDSAAKVQCSQEWIPFCN 960
 OGTA076c (SEQ ID No: 3) GEECLYMSAKTNLIDLCKPTDCSTKLPPFCKVNVSSELEYKSFDSAAKVQCSQEWIPFCN 960
 OGTA076a (SEQ ID No: 1) GEECLYMSAKTNLIDLCKPTDCSTKLPPFCKVNVSSELEYKSFDSAAKVQCSQEWIPFCN 960

 OGTA076b (SEQ ID No: 2) KCFLKIKPVSILTSQASDTCHSYGGTLPVSLSQIEQDFITSLPPMPEATLWIGLRWTAYE 1020
 OGTA076c (SEQ ID No: 3) KCFLKIKPVSILTSQASDTCHSYGGTLPVSLSQIEQDFITSLPPMPEATLWIGLRWTAYE 1020
 OGTA076a (SEQ ID No: 1) KCFLKIKPVSILTSQASDTCHSYGGTLPVSLSQIEQDFITSLPPMPEATLWIGLRWTAYE 1020

 OGTA076b (SEQ ID No: 2) KINKTNDNREIYTSNFHPLLVSGLRILIPENFPEEESRYHCALILNLQSPFPTGTWNPTSC 1080
 OGTA076c (SEQ ID No: 3) KINKTNDNREIYTSNFHPLLVSGLRILIPENFPEEESRYHCALILNLQSPFPTGTWNPTSC 1080
 OGTA076a (SEQ ID No: 1) KINKTNDNREIYTSNFHPLLVSGLRILIPENFPEEESRYHCALILNLQSPFPTGTWNPTSC 1080

 OGTA076b (SEQ ID No: 2) SERHVFVSLCQKYSEVKSRQTLQNASETVKYIINNNLYKIIPTKLTWHSAKRECLKSNNMQLVS 1140
 OGTA076c (SEQ ID No: 3) SERHVFVSLCQKYSEVKSRQTLQNASETVKYIINNNLYKIIPTKLTWHSAKRECLKSNNMQLVS 1140
 OGTA076a (SEQ ID No: 1) SERHVFVSLCQKYSEVKSRQTLQNASETVKYIINNNLYKIIPTKLTWHSAKRECLKSNNMQLVS 1140

 OGTA076b (SEQ ID No: 2) ITDPYQQAFLSVQALLHNSSLWICLFSODDELFNFCWSDGKRLHFSRWAETNQLEDCVVL 1200
 OGTA076c (SEQ ID No: 3) ITDPYQQAFLSVQALLHNSSLWICLFSODDELFNFCWSDGKRLHFSRWAETNQLEDCVVL 1200
 OGTA076a (SEQ ID No: 1) ITDPYQQAFLSVQALLHNSSLWICLFSODDELFNFCWSDGKRLHFSRWAETNQLEDCVVL 1200

OGTA076b (SEQ ID No: 2) DTDCFWKTVDCNDNQPGAI^CYYSGNTEKEVKPVDSVKCPSPVLM^TWIPFQNCCYNFII 1260
 OGTA076c (SEQ ID No: 3) DTDCFWKTVDCNDNQPGAI^CYYSGNTEKEVKPVDSVKCPSPVLM^TWIPFQNCCYNFII 1260
 OGTA076a (SEQ ID No: 1) DTDCFWKTVDCNDNQPGAI^CYYSGNTEKEVKPVDSVKCPSPVLM^TWIPFQNCCYNFII 1260

 OGTA076b (SEQ ID No: 2) TGNRHMTTQDEVHTKCQKLMPKSHILSIRDEKENNFVLE^CLLYFNYMASWVMLGITYRN 1320
 OGTA076c (SEQ ID No: 3) TGNRHMTTQDEVHTKCQKLMPKSHILSIRDEKENNFVLE^CLLYFNYMASWVMLGITYRN 1320
 OGTA076a (SEQ ID No: 1) TGNRHMTTQDEVHTKCQKLMPKSHILSIRDEKENNFVLE^CLLYFNYMASWVMLGITYRN 1320

 OGTA076b (SEQ ID No: 2) NSIWMFDKTPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDTC^ICKVIEAVYFHQSILAC 1380
 OGTA076c (SEQ ID No: 3) NSIWMFDKTPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDTC^ICKVIEAVYFHQSILAC 1380
 OGTA076a (SEQ ID No: 1) NSIWMFDKTPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDTC^ICKVIEAVYFHQSILAC 1380

 OGTA076b (SEQ ID No: 2) KIENVYDKEEINNTLPQPNPYED^IYSVICKVYTKYEA^ILMCSOSGGHLASVHNNGOLF 1440
 OGTA076c (SEQ ID No: 3) KIENVYDKEEINNTLPQPNPYED^IYSVICKVYTKYEA^ILMCSOSGGHLASVHNNGOLF 1440
 OGTA076a (SEQ ID No: 1) KIENVYDKEEINNTLPQPNPYED^IYSVICKVYTKYEA^ILMCSOSGGHLASVHNNGOLF 1440

 OGTA076b (SEQ ID No: 2) LEDIVKRDGFFI^IWVGLSSHDGSESSPBWGDGSTFYIPWKQQTSPGNCVL^ILDPKG^IWKH^I 1500
 OGTA076c (SEQ ID No: 3) LEDIVKRDGFFI^IWVGLSSHDGSESSPBWGDGSTFYIPWKQQTSPGNCVL^ILDPKG^IWKH^I 1500
 OGTA076a (SEQ ID No: 1) LEDIVKRDGFFI^IWVGLSSHDGSESSPBWGDGSTFYIPWKQQTSPGNCVL^ILDPKG^IWKH^I 1500

 OGTA076b (SEQ ID No: 2) KCNSVKDGAICYKPTKSKL^ISLR^ITYSSRCPAAKENGSR^ICYKHCYKSDQALHSFSEAK 1560
 OGTA076c (SEQ ID No: 3) KCNSVKDGAICYKPTKSKL^ISLR^ITYSSRCPAAKENGSR^ICYKHCYKSDQALHSFSEAK 1560
 OGTA076a (SEQ ID No: 1) KCNSVKDGAICYKPTKSKL^ISLR^ITYSSRCPAAKENGSR^ICYKHCYKSDQALHSFSEAK 1560

 OGTA076b (SEQ ID No: 2) KLC SKHDHSATI^IVKDEDENK^IEVSLR^IMEENNNT^ITM^IRV^IWL^IGS^IHS^IVS^ID^IGS^IMLGSEVT 1620
 OGTA076c (SEQ ID No: 3) KLC SKHDHSATI^IVKDEDENK^IEVSLR^IMEENNNT^ITM^IRV^IWL^IGS^IHS^IVS^ID^IGS^IMLGSEVT 1608
 OGTA076a (SEQ ID No: 1) KLC SKHDHSATI^IVKDEDENK^IEVSLR^IMEENNNT^ITM^IRV^IWL^IGS^IHS^IVS^ID^IGS^IMLGSEVT 1608

 OGTA076b (SEQ ID No: 2) FV^IKWENK^ISKC^IVGRC^ICSMLIASNETWKKVECEHGF^ICRV^ICKV^IPLDCP^ISST^IW^IQ^IPFQ^IDS^ICY^IF 1680
 OGTA076c (SEQ ID No: 3) -----CFPS^ISTW^IQ^IPFQ^IDS^ICF^I 1624
 OGTA076a (SEQ ID No: 1) FV^IKWENK^ISKC^IVGRC^ICSMLIASNETWKKVECEHGF^ICRV^ICKV^IPLDCP^IY^ITATA^IIVAT^ISIL

 OGTA076b (SEQ ID No: 2) 1QEA^IIKVEGIED^IDRV^IQCT^ID^IAD^IMS^II^INEE^IEN^IAF^IIL^ILKKQ^IW^IKG^IPD^ILLG^ICM^IFY^ID^IDD 1740
 OGTA076c (SEQ ID No: 3) 1QEA^IIKVEGIED^IDRV^IQCT^ID^IAD^IMS^II^INEE^IEN^IAF^IIL^ILKKQ^IW^IKG^IPD^ILLG^ICM^IFY^ID^IDD 1684
 OGTA076a (SEQ ID No: 1) VLM^ICG^ILI^IW^IFL^IQ^IR^IH^IL^IAC^IFS^ISV^IY^IAQ^IGV^INE^IDE^IIM^IL^IP^ISHD^I-----

 OGTA076b (SEQ ID No: 2) KRKYLSDNHILISALVIASIA^ISTV^ITLV^IGAI^IINFLYKKHSDSRFTT^IV^IF^IST^IAP^IQ^ISP^IY^INE^IDC^IVL 1860
 OGTA076c (SEQ ID No: 3) KRKYLSDNHILISALVIASIA^ISTV^ITLV^IGAI^IINFLYKKHSDSRFTT^IV^IF^IST^IAP^IQ^ISP^IY^INE^IDC^IVL 1804
 OGTA076a (SEQ ID No: 1) -----
 OGTA076b (SEQ ID No: 2) VVGEEN^IEPVQ^IFD 1873
 OGTA076c (SEQ ID No: 3) VVGEEN^IEPVQ^IFD 1817
 OGTA076a (SEQ ID No: 1) -----

Mass Match Peptides (bold):EG^IAK [14]**Tandem Peptides (underline):**

EGIAK [14]

Peptide Source: iTRAQ Ovarian cancer

OGTA076b (SEQ ID No: 2) MRTGWATP^IRRPAGL^IMLL^IW^IF^IDL^IA^IP^ISCRA^IANDP^IFT^IV^IH^IG^IN^IG^IK^IC^IK^IP^IV^IY^IG^IW^IV^IADD^ICD 60
 OGTA076c (SEQ ID No: 3) MRTGWATP^IRRPAGL^IMLL^IW^IF^IDL^IA^IP^ISCRA^IANDP^IFT^IV^IH^IG^IN^IG^IK^IC^IK^IP^IV^IY^IG^IW^IV^IADD^ICD 60

OGTA076a (SEQ ID No: 1) MRTGWT PRRPAGLMLILEWFFDLAEPGR&ANDPPTIVHGNNTGKCKIKPVYGNIVADD 60

OGTA076c (SEQ ID No: 2) ETEOKLWKVSQHRLPHLHSOKC1GLDITKSVNLRMESCDSAMLMWKCERHSLYGAA 120
OGTA076c (SEQ ID No: 3) ETEOKLWKVSQHRLPHLHSOKC1GLDITKSVNLRMESCDSAMLMWKCERHSLYGAA 120
ETEOKLWKVSQHRLPHLHSOKC1GLDITKSVNLRMESCDSAMLMWKCERHSLYGAA 120

OGTA076b (SEQ ID No: 2) YTKLDRGHTAISNASDWKKGGSEESLCQCQPYHEIYTRDGNNSYGRCEPPFLIDGTW 180
OGTA076c (SEQ ID No: 3) YTKLDRGHTAISNASDWKKGGSEESLCQCQPYHEIYTRDGNNSYGRCEPPFLIDGTW 180
OGTA076a (SEQ ID No: 1) YTKLDRGHTAISNASDWKKGGSEESLCQCQPYHEIYTRDGNNSYGRCEPPFLIDGTW 180

OGTA076b (SEQ ID No: 2) HDCILDEDHSGPWCATTLNEYEDRKWCICLKPENGCEENWEKNECFCGSCYQPNIQALSW 240
OGTA076c (SEQ ID No: 3) HDCILDEDHSGPWCATTLNEYEDRKWCICLKPENGCEENWEKNECFCGSCYQPNIQALSW 240
OGTA076a (SEQ ID No: 1) HDCILDEDHSGPWCATTLNEYEDRKWCICLKPENGCEENWEKNECFCGSCYQPNIQALSW 240

OGTA076b (SEQ ID No: 2) KERAVSCQNAQADLISNSAELITYLKEECAIKAIPWICLNGQIYSAQHGWNSDHKPLN 300
OGTA076c (SEQ ID No: 3) KERAVSCQNAQADLISNSAELITYLKEECAIKAIPWICLNGQIYSAQHGWNSDHKPLN 300
OGTA076a (SEQ ID No: 1) KERAVSCQNAQADLISNSAELITYLKEECAIKAIPWICLNGQIYSAQHGWNSDHKPLN 300

OGTA076b (SEQ ID No: 2) NWDPDPRPSAPTICCSSCARMDAE5CLWQSFSCEAQLPYVCRKPLNNTVELTDWVTTSDT 360
OGTA076c (SEQ ID No: 3) NWDPDPRPSAPTIIGGSSCARMDAE5CLWQSFSCEAQLPYVCRKPLNNTVELTDWVTTSDT 360
OGTA076a (SEQ ID No: 1) NWDPDPRPSAPTIIGGSSCARMDAE5CLWQSFSCEAQLPYVCRKPLNNTVELTDWVTTSDT 360

OGTA076b (SEQ ID No: 2) CDAGWLPNNGPCVLLVNESSNSUCAHAKCKAFSSLISIHSIADVEVVVTKLHNEDIKEE 420
OGTA076c (SEQ ID No: 3) CDAGWLPNNGPCVLLVNESSNSUCAHAKCKAFSSLISIHSIADVEVVVTKLHNEDIKEE 420
OGTA076a (SEQ ID No: 1) CDAGWLPNNGPCVLLVNESSNSUCAHAKCKAFSSLISIHSIADVEVVVTKLHNEDIKEE 420

OGTA076b (SEQ ID No: 2) VVIGLKNINIPTLFWQASDGTEVTLTYWDENSPNVFVNKTFCVSYLGELGQWKVQSCEEK 480
OGTA076c (SEQ ID No: 3) VVIGLKNINIPTLFWQASDGTEVTLTYWDENSPNVFVNKTFCVSYLGELGQWKVQSCEEK 480
OGTA076a (SEQ ID No: 1) VVIGLKNINIPTLFWQASDGTEVTLTYWDENSPNVFVNKTFCVSYLGELGQWKVQSCEEK 480

OGTA076b (SEQ ID No: 2) IKYVCRKGEKINLADASSDRMCPPDEGWRKRHGETCYK1YEDDEVPGTNCNLITTSRFEQEV 540
OGTA076c (SEQ ID No: 3) IKYVCRKGEKINLADASSDRMCPPDEGWRKRHGETCYK1YEDDEVPGTNCNLITTSRFEQEV 540
OGTA076a (SEQ ID No: 1) IKYVCRKGEKINLADASSDRMCPPDEGWRKRHGETCYK1YEDDEVPGTNCNLITTSRFEQEV 540

OGTA076b (SEQ ID No: 2) LNDLMKYDK5L8K1YFWTGLRDVDSGCGEYNWATVCGRRRAVTFSNNWFLP2PASPGGCVAM 600
OGTA076c (SEQ ID No: 3) LNDLMKYDK5L8K1YFWTGLRDVDSGCGEYNWATVCGRRRAVTFSNNWFLP2PASPGGCVAM 600
OGTA076a (SEQ ID No: 1) LNDLMKYDK5L8K1YFWTGLRDVDSGCGEYNWATVCGRRRAVTFSNNWFLP2PASPGGCVAM 600

OGTA076b (SEQ ID No: 2) STGKSVGKWEVKCRSFKA1SLCKKMSGPGLPREEASPKPDEPCPGWQSFPA5LSCYKF 660
OGTA076c (SEQ ID No: 3) STGKSVGKWEVKCRSFKA1SLCKKMSGPGLPREEASPKPDEPCPGWQSFPA5LSCYKF 660
OGTA076a (SEQ ID No: 1) STGKSVGKWEVKCRSFKA1SLCKKMSGPGLPREEASPKPDCPCPGWQSFPA5LSCYKF 660

OGTA076b (SEQ ID No: 2) HAERIVRKRNWEEAERFCQALGAHLSFSHVDEIKEFLHPLTQFSGQHNLWIGLNKRSP 720
OGTA076c (SEQ ID No: 3) HAERIVRKRNWEEAERFCQALGAHLSFSHVDEIKEFLHPLTQFSGQHNLWIGLNKRSP 720
OGTA076a (SEQ ID No: 1) HAERIVRKRNWEEAERFCQALGAHLSFSHVDEIKEFLHPLTQFSGQHNLWIGLNKRSP 720

OGTA076b (SEQ ID No: 2) DLQGSWQWSDRTPVSTIIMPNFQOQYDIDRCAAVKVPHRPWRRGWHFYDDRFEFYLRPF 780
OGTA076c (SEQ ID No: 3) DLQGSWQWSDRTPVSTIIMPNFQOQYDIDRCAAVKVPHRPWRRGWHFYDDRFEFYLRPF 780
OGTA076a (SEQ ID No: 1) DLQGSWQWSDRTPVSTIIMPNFQOQYDIDRCAAVKVPHRPWRRGWHFYDDRFEFYLRPF 780

OGTA076b (SEQ ID No: 2) ACDTKLENVQOIPKGRTPKTDPWNPDRAIGHCPPLIIEGSEYFVADLHLYEEAVLYC 840
OGTA076c (SEQ ID No: 3) ACDTKLENVQOIPKGRTPKTDPWNPDRAIGHCPPLIIEGSEYFVADLHLYEEAVLYC 840
OGTA076a (SEQ ID No: 1) ACDTKLENVQOIPKGRTPKTDPWNPDRAIGHCPPLIIEGSEYFVADLHLYEEAVLYC 840

OGTA076b (SEQ ID No: 2) ASHHSFLATITISFVGLKAIKNIKIANISGDGQKWWIRIS2WPIIDHFTYSRYPWHRFPVTF 900
OGTA076c (SEQ ID No: 3) ASHHSFLATITISFVGLKAIKNIKIANISGDGQKWWIRIS2WPIIDHFTYSRYPWHRFPVTF 900
OGTA076a (SEQ ID No: 1) ASHHSFLATITISFVGLKAIKNIKIANISGDGQKWWIRIS2WPIIDHFTYSRYPWHRFPVTF 900

OGTA076b (SEQ ID No: 2) GEECLYMSAKTWLIDLKPTDCSTKLPF1CEKYNVSSLKYSYPDAAKVQCSEQWIPFQN 960
OGTA076c (SEQ ID No: 3) GEECLYMSAKTWLIDLKPTDCSTKLPF1CEKYNVSSLKYSYPDAAKVQCSEQWIPFQN 960
OGTA076a (SEQ ID No: 1) GEECLYMSAKTWLIDLKPTDCSTKLPF1CEKYNVSSLKYSYPDAAKVQCSEQWIPFQN 960

OGTA076a (SEQ ID No: 1) GEECLYMSAKTNWLIDLGKPTDCSTKLPPFICEKVNVSSEKYS PDSA AKVQCS EOWI PFCN 960 *****
 OGTA076b (SEQ ID No: 2) KCFPLK(KPVSLTFSQASDTCHSYGGTLPVSLSQIPQDFITSLLPDMEATLWIGLRWTAYE 1020
 OGTA076c (SEQ ID No: 3) KCFPLK(KPVSLTFSQASDTCHSYGGTLPVSLSQIPQDFITSLLPDMEATLWIGLRWTAYE 1020
 OGTA076a (SEQ ID No: 1) KCFPLK(KPVSLTFSQASDTCHSYGGTLPVSLSQIPQDFITSLLPDMEATLWIGLRWTAYE 1020 *****
 OGTA076b (SEQ ID No: 2) KINKWDTNRELTYSHFHPLLVSGLRLIPENPFEPEESRYHCALILNLOKSPPTGTWNFTSC 1080
 OGTA076c (SEQ ID No: 3) KINKWDTNRELTYSHFHPLLVSGLRLIPENPFEPEESRYHCALILNLOKSPPTGTWNFTSC 1080
 OGTA076a (SEQ ID No: 1) KINKWDTNRELTYSHFHPLLVSGLRLIPENPFEPEESRYHCALILNLOKSPPTGTWNFTSC 1080 *****
 OGTA076b (SEQ ID No: 2) SERHFVSLCQKYSRVKSRQTLQNASETVX~~YNNNNNN~~IIIPKILLWHSAKR3CLSKNNQQLVS 1140
 OGTA076c (SEQ ID No: 3) SERHFVSLCQKYSRVKSRQTLQNASETVX~~YNNNNNN~~IIIPKILLWHSAKR3CLSKNNQQLVS 1140
 OGTA076a (SEQ ID No: 1) SERHFVSLCQKYSRVKSRQTLQNASETVX~~YNNNNNN~~IIIPKILLWHSAKR3CLSKNNQQLVS 1140 *****
 OGTA076b (SEQ ID No: 2) ITDPYQQAFLSVQALLHNSSLWIGLFSQDLSLNGMSDGKRLHPSRNASTNGQ-EDCVVL 1200
 OGTA076c (SEQ ID No: 3) ITDPYQQAFLSVQALLHNSSLWIGLFSQDLSLNGMSDGKRLHPSRNASTNGQ-EDCVVL 1200
 OGTA076a (SEQ ID No: 1) ITDPYQQAFLSVQALLHNSSLWIGLFSQDLSLNGMSDGKRLHPSRNASTNGQ-EDCVVL 1200 *****
 OGTA076b (SEQ ID No: 2) DTGCFWKTVDNDNQPGCAIYYYSGN2TEKEVKVPSVKCPSVPLNT~~Y~~WIPFQNCCYNFII 1260
 OGTA076c (SEQ ID No: 3) DTGCFWKTVDNDNQPGCAIYYYSGN2TEKEVKVPSVKCPSVPLNT~~Y~~WIPFQNCCYNFII 1260
 OGTA076a (SEQ ID No: 1) DTGCFWKTVDNDNQPGCAIYYYSGN2TEKEVKVPSVKCPSVPLNT~~Y~~WIPFQNCCYNFII 1260 *****
 OGTA076b (SEQ ID No: 2) TKNRHMATTQDEVHTKQCLNPKSHLISIRDEKENNFVLECLLYENYMASVWMLGITYRN 1320
 OGTA076c (SEQ ID No: 3) TKNRHMATTQDEVHTKQCLNPKSHLISIRDEKENNFVLECLLYENYMASVWMLGITYRN 1320
 OGTA076a (SEQ ID No: 1) TKNRHMATTQDEVHTKQCLNPKSHLISIRDEKENNFVLECLLYENYMASVWMLGITYRN 1320 *****
 OGTA076b (SEQ ID No: 2) NSLMWFDTKPLSYTHWRAGRPTIKNEKFLAGLSTDCPWD1C7PKV12EAVYFHQSILAC 1380
 OGTA076c (SEQ ID No: 3) NSLMWFDTKPLSYTHWRAGRPTIKNEKFLAGLSTDCPWD1C7PKV12EAVYFHQSILAC 1380
 OGTA076a (SEQ ID No: 1) NSLMWFDTKPLSYTHWRAGRPTIKNEKFLAGLSTDCPWD1C7PKV12EAVYFHQSILAC 1380 *****
 OGTA076b (SEQ ID No: 2) KIENVDYKEEENNTLQFQNPYEDGIYSV1QRKVTWYFA1NMC3QSGGHLASVHNQNGQLE 1440
 OGTA076c (SEQ ID No: 3) KIENVDYKEEENNTLQFQNPYEDGIYSV1QRKVTWYFA1NMC3QSGGHLASVHNQNGQLE 1440
 OGTA076a (SEQ ID No: 1) KIENVDYKEEENNTLQFQNPYEDGIYSV1QRKVTWYFA1NMC3QSGGHLASVHNQNGQLE 1440 *****
 OGTA076b (SEQ ID No: 2) LEDIVKRQDFPLWVGLSHDGSESSFEWSDGSTFDYIPWKGQTSPGNCVLLDPKGTVKHE 1500
 OGTA076c (SEQ ID No: 3) LEDIVKRQDFPLWVGLSHDGSESSFEWSDGSTFDYIPWKGQTSPGNCVLLDPKGTVKHE 1500
 OGTA076a (SEQ ID No: 1) LEDIVKRQDFPLWVGLSHDGSESSFEWSDGSTFDYIPWKGQTSPGNCVLLDPKGTVKHE 1500 *****
 OGTA076b (SEQ ID No: 2) KCSVKDGAICYKPTKSKLRLITYSSRCPAAKENGSPW1C7KGHCYK5DQALHSFSEAK 1560
 OGTA076c (SEQ ID No: 3) KCSVKDGAICYKPTKSKLRLITYSSRCPAAKENGSPW1C7KGHCYK5DQALHSFSEAK 1560
 OGTA076a (SEQ ID No: 1) KCSVKDGAICYKPTKSKLRLITYSSRCPAAKENGSPW1C7KGHCYK5DQALHSFSEAK 1560 *****
 OGTA076b (SEQ ID No: 2) KLCSDKHDSATIVS1KDEDENKEVSR1MPEENNNT1MRWVLGLSQHSVDSWLDGSEVT 1620
 OGTA076c (SEQ ID No: 3) KLCSDKHDSATIVS1KDEDENKEVSR1MPEENNNT1MRWVLGLSQHSVDSWLDGSEVT 1608
 OGTA076a (SEQ ID No: 1) KLCSDKHDSATIVS1KDEDENKEVSR1MPEENNNT1MRWVLGLSQHSVDSWLDGSEVT 1608 *****
 OGTA076b (SEQ ID No: 2) FVKWENKSKSGVCRCSMLIASNETWKKVECEHGFGRVVCKVPLDCPSSTWQFDSCYIF 1680
 OGTA076c (SEQ ID No: 3) FVKWENKSKSGVCRCSMLIASNETWKKVECEHGFGRVVCKVPLGPDY1A1A1VATLSIL 1624
 OGTA076a (SEQ ID No: 1) FVKWENKSKSGVCRCSMLIASNETWKKVECEHGFGRVVCKVPLGPDY1A1A1VATLSIL 1624
 OGTA076b (SEQ ID No: 2) LQZAIKVESIEDVRNQTDHGADMISIHNNEENAFILDILKKCWNKGPDIDLLGMFYDIDD 1740
 OGTA076c (SEQ ID No: 3) LQZAIKVESIEDVRNQTDHGADMISIHNNEENAFILDILKKCWNKGPDIDLLGMFYDIDD 1684
 OGTA076a (SEQ ID No: 1) VLMGGIWF1FQHRLHLAGFSSVRYAQGVNEDIMLPSFHD-----
 OGTA076b (SEQ ID No: 2) ASFWFPDNSNMNTFDKWTDDDBDILWDTCAPLHKTGEWKKGNCEVSSVEGLCKTAIPY 1800
 OGTA076c (SEQ ID No: 3) ASFWFPDNSNMNTFDKWTDDDBDILWDTCAPLHKTGEWKKGNCEVSSVEGLCKTAIPY 1744
 OGTA076a (SEQ ID No: 1) KRXYLSDNHILITSLVIASTV1TVLGAI1WFLYKKHSDSRFTTVFSTAPQSPYNEDCVL 1860
 OGTA076c (SEQ ID No: 2) KRXYLSDNHILITSLVIASTV1TVLGAI1WFLYKKHSDSRFTTVFSTAPQSPYNEDCVL 1804

OGTA076a (SEQ ID No: 1) -----
 OGTA076b (SEQ ID No: 2) VVGEENEYFVQFD 1873
 OGTA076c (SEQ ID No: 3) VVGEENEYFVQFD 1817
 OGTA076a (SEQ ID No: 1) -----

Mass Match Peptides (bold):

LALK [35]
 YLNNLYK [66]

Tandem Peptides (underline):

LALK [35]
 YLNNLYK [66]

Peptide Source: 1D GE Chronic lymphocytic leukaemia

OGTA076b (SEQ ID No: 2) MRTGWATPWRPAGLIMLILWFFDIAEEDSGR **MRTGWATPWRPAGLIMLILWFFDIAEEDSGR** 60
 OGTA076c (SEQ ID No: 3) MRTGWATPWRPAGLIMLILWFFDIAEEDSGR **MRTGWATPWRPAGLIMLILWFFDIAEEDSGR** 60
 OGTA076a (SEQ ID No: 1) MRTGWATPWRPAGLIMLILWFFDIAEEDSGR **MRTGWATPWRPAGLIMLILWFFDIAEEDSGR** 60

 OGTA076b (SEQ ID No: 2) FTEDKLWIKVSGQRIEHHHSORKCQGNDITKSVNFLRMPSCCSSAMILWWCERHSLYGRAR 120
 OGTA076c (SEQ ID No: 3) FTEDKLWIKVSGQRIEHHHSORKCQGNDITKSVNFLRMPSCCSSAMILWWCERHSLYGRAR 120
 OGTA076a (SEQ ID No: 1) FTEDKLWIKVSGQRIEHHHSORKCQGNDITKSVNFLRMPSCCSSAMILWWCERHSLYGRAR 120

 OGTA076b (SEQ ID No: 2) YRLAIKDGHGWAISASDVKKGSEESLICQDQYH8IYTRCGNSYGRPCPEFFLIDGTW8 180
 OGTA076c (SEQ ID No: 3) YRLAIKDGHGWAISASDVKKGSEESLICQDQYH8IYTRCGNSYGRPCPEFFLIDGTW8 180
 OGTA076a (SEQ ID No: 1) YRLAIKDGHGWAISASDVKKGSEESLICQDQYH8IYTRCGNSYGRPCPEFFLIDGTW8 180

 OGTA076b (SEQ ID No: 2) HDC1LDE2DHSGPWCAPIVLLNVEYCDRKWGIKLKPENGCEDNWEKNEQFGSCYCQFNITQIALSW 240
 OGTA076c (SEQ ID No: 3) HDC1LDE2DHSGPWCAPIVLLNVEYCDRKWGIKLKPENGCEDNWEKNEQFGSCYCQFNITQIALSW 240
 OGTA076a (SEQ ID No: 1) HDC1LDE2DHSGPWCAPIVLLNVEYCDRKWGIKLKPENGCEDNWEKNEQFGSCYCQFNITQIALSW 240

 OGTA076b (SEQ ID No: 2) KEAVVSCQNQGADLLSINSAAEITYLKEKEGIAKIFWIGLNQIYSGRGWENSDHKPLNFL 300
 OGTA076c (SEQ ID No: 3) KEAVVSCQNQGADLLSINSAAEITYLKEKEGIAKIFWIGLNQIYSGRGWENSDHKPLNFL 300
 OGTA076a (SEQ ID No: 1) KEAVVSCQNQGADLLSINSAAEITYLKEKEGIAKIFWIGLNQIYSGRGWENSDHKPLNFL 300

 OGTA076b (SEQ ID No: 2) NWDPDRPASPTIGGSSCARMDAESGLWQSFPCCAEQLPVVCRKEFLNNTVVELTDVWTYSDTR 360
 OGTA076c (SEQ ID No: 3) NWDPDRPASPTIGGSSCARMDAESGLWQSFPCCAEQLPVVCRKEFLNNTVVELTDVWTYSDTR 360
 OGTA076a (SEQ ID No: 1) NWDPDRPASPTIGGSSCARMDAESGLWQSFPCCAEQLPVVCRKEFLNNTVVELTDVWTYSDTR 360

 OGTA076b (SEQ ID No: 2) CDAGNLPNNGFCYLLVNESSWDRAHAKCKAFSSDLISIHS1ADWVVVTTKIHNEDEKEE 420
 OGTA076c (SEQ ID No: 3) CDAGNLPNNGFCYLLVNESSWDRAHAKCKAFSSDLISIHS1ADWVVVTTKIHNEDEKEE 420
 OGTA076a (SEQ ID No: 1) CDAGNLPNNGFCYLLVNESSWDRAHAKCKAFSSDLISIHS1ADWVVVTTKIHNEDEKEE 420

 OGTA076b (SEQ ID No: 2) VVIGLKNINIPITLFWQSDDGTEVTLIYWDNEPNVNVYNTIPNCVSYLGEELGQNKVQSCCEEK 480
 OGTA076c (SEQ ID No: 3) VVIGLKNINIPITLFWQSDDGTEVTLIYWDNEPNVNVYNTIPNCVSYLGEELGQNKVQSCCEEK 480
 OGTA076a (SEQ ID No: 1) VVIGLKNINIPITLFWQSDDGTEVTLIYWDNEPNVNVYNTIPNCVSYLGEELGQNKVQSCCEEK 480

 OGTA076b (SEQ ID No: 2) LKYVCKRKGEKLNDASSDRMCPPDEGWKRRHGETCYK1YEDEVFEGTNCLNTITSREKEE 540
 OGTA076c (SEQ ID No: 3) LKYVCKRKGEKLNDASSDRMCPPDEGWKRRHGETCYK1YEDEVFEGTNCLNTITSREKEE 540
 OGTA076a (SEQ ID No: 1) LKYVCKRKGEKLNDASSDRMCPPDEGWKRRHGETCYK1YEDEVFEGTNCLNTITSREKEE 540

 OGTA076b (SEQ ID No: 2) ~~IKDDEKKYD~~KS1KPKKEPTTCERDVS~~CG~~EYINWATVGRERRAVTFSNWNFLEPASPGCCVAM 600
 OGTA076c (SEQ ID No: 3) ~~IKDDEKKYD~~KS1KPKKEPTTCERDVS~~CG~~EYINWATVGRERRAVTFSNWNFLEPASPGCCVAM 600
 OGTA076a (SEQ ID No: 1) ~~IKDDEKKYD~~KS1KPKKEPTTCERDVS~~CG~~EYINWATVGRERRAVTFSNWNFLEPASPGCCVAM 600

 OGTA076b (SEQ ID No: 2) STGKSVGKNEVKDCRSFKALSCKKMSGPIGPKRASPKPDCPCPEGWQSFPASLSCYKVF 660
 OGTA076c (SEQ ID No: 3) STGKSVGKNEVKDCRSFKALSCKKMSGPIGPKRASPKPDCPCPEGWQSFPASLSCYKVF 660

OGTA076a (SEQ ID No: 1) KCNSVKDQGQKQXEPKTSKKLSRITYSSRCPAAKENGSRWIQYKCHCYKSPQATRHFSEAK 1560

OGTA076b (SEQ ID No: 2) KLCSKHRSATIVSIRDDENKFVSRIMRENNNTMRVNLLSQHSVDQSWSLDGSEVT 1620

OGTA076c (SEQ ID No: 3) KLCSKHRSATIVSIRDDENKFVSRIMRENNNTMRVNLLSQHSV----- 1608

OGTA076a (SEQ ID No: 1) KLCSKHRSATIVSIRDDENKFVSRIMRENNNTMRVNLLSQHSVDQSWSLDGSEVT 1608

OGTA076b (SEQ ID No: 2) FVWENKSKSGVGRCSMIYASNEPTWIKVCEPICGRVVKVPLDCPSSTWIQKQDSCYIF 1680

OGTA076c (SEQ ID No: 3) FVWENKSKSGVGRCSMIYASNEPTWIKVCEPICGRVVKVPLDCPSSTWIQKQDSCYIF 1624

OGTA076a (SEQ ID No: 1) FVWENKSKSGVGRCSMIYASNEPTWIKVCEPICGRVVKVPLGPDYTAIAIIIVATLSL

OGTA076b (SEQ ID No: 2) IQLAIKVESIEDVRNOCTDHGADMISIHNEEENAPILILKKQWKGPDLLLGMPYDTD 1740

OGTA076c (SEQ ID No: 3) IQLAIKVESIEDVRNQCTDHGADMISIHNEEENAPILILKKQWKGPDLLLGMPYDTD 1684

OGTA076a (SEQ ID No: 1) VLMGGLIWFLFQRHRLHAGFSSVRYAQGVNEDIEMPSFHD-----

OGTA076b (SEQ ID No: 2) ASFKWFDNNSNMPDKWTDODDDEDLVICAFLHIKTGEWKKGNCEVSSVEGLCKTAIPY 1800

OGTA076c (SEQ ID No: 3) ASFKWFDNNSNMPDKWTDODDDEDLVICAFLHIKTGEWKKGNCEVSSVEGLCKTAIPY 1744

OGTA076a (SEQ ID No: 1) -----

OGTA076b (SEQ ID No: 2) VVGEENEYPVQFD 1873

OGTA076c (SEQ ID No: 3) VVGEENEYPVQFD 1817

OGTA076a (SEQ ID No: 1) -----

Mass Match Peptides (bold):

AANDEFTIVHGNTK [4]
 CEHSLYGAA [6]
 CLGLDIK [7]
 CSMLIASNETWKK [8]
 DGAICYKPTK [9]
 ELTYSNFHPLLVSGR [15]
 FEQEYLNDIMK [18]
 GHHFYDDR [21]
 HDHSATIVSIK [22]
 HFVSILCQR [23]
 HMATTQDEVIIK [25]
 IEMVDIK [27]
 IPENFEEEESR [29]
 ISEWMPIDDHFTYSR [30]
 KGNCEVSSVEGLCK [31]
 KVECEHGFGR [33]
 KYFWTGLR [34]
 LFHLSQK [36]
 LHNEDIK [38]
 NWEEAER [46]
 RGWHFYDDR [48]
 KLHFSR [50]
 RNWEEDAER [51]
 SDQALESFSEAK [52]
 SHILSIR [53]
 SPDLOGSQWQSDR [55]
 TPLSYTHWR [58]
 TPVSIIMPNEFQQDYDIR [59]
 VCECHGFGGR [60]
 VFHRPWR [61]
 VQCSEQWIPFQNK [63]
 WVSQHR [64]

YFWTGLR [65]
YLNLYK [66]

Tandem Peptides (underline):

DGAICYKPTK [9]
IFENFFEEESR [29]
ISETWPIDDHFTYSR [30]
SDQALHSFSEAK [52]
SNFHPLLVSGR [54]
SPDIQGSWQWSDR [55]
TPLSYTHWR [58]

Peptide Source: 1D GE Colorectal cancer

OGTA076b (SEQ ID No: 2) MRTGWATPRRPAGLMLFLWFFDLAEPGSGRAANDPFTIVHGNNTGKCKIKPVYGVIVADCD 60
OGTA076c (SEQ ID No: 3) MRTGWATPRRPAGLMLFLWFFDLAEPGSGRAANDPFTIVHGNNTGKCKIKPVYGVIVADCD 60
OGTA076a (SEQ ID No: 1) MRTGWATPRRPAGLMLFLWFFDLAEPGSGRAANDPFTIVHGNNTGKCKIKPVYGVIVADCD 60

OGTA076b (SEQ ID No: 2) ETEDKLWKVVSQHRLPHLHSOKCGIDIDITSVNELMHC1CSAAMMWCEHHHSILYGAAR 120
OGTA076c (SEQ ID No: 3) ETEDKLWKVVSQHRLPHLHSOKCGIDIDITSVNELMHC1CSAAMMWCEHHHSILYGAAR 120
OGTA076a (SEQ ID No: 1) ETEDKLWKVVSQHRLPHLHSOKCGIDIDITSVNELMHC1CSAAMMWCEHHHSILYGAAR 120

OGTA076b (SEQ ID No: 2) YRLAIKDGHGTIAISNASDWMKKGSEESLCDQPYHEIYIREGNSYGRPCBFPFLIDCTWII 180
OGTA076c (SEQ ID No: 3) YRLAIKDGHGTIAISNASDWMKKGSEESLCDQPYHEIYIREGNSYGRPCBFPFLIDCTWII 180
OGTA076a (SEQ ID No: 1) YRLAIKDGHGTIAISNASDWMKKGSEESLCDQPYHEIYIREGNSYGRPCBFPFLIDCTWII 180

OGTA076b (SEQ ID No: 2) HDC II DEDHSGPWCATTINVEYDREWG1CLKPENGCEDNWKEKNEQFGSCYQFNTQTALSW 240
OGTA076c (SEQ ID No: 3) HDC II DEDHSGPWCATTINVEYDREWG1CLKPENGCEDNWKEKNEQFGSCYQFNTQTALSW 240
OGTA076a (SEQ ID No: 1) HDC II DEDHSGPWCATTINVEYDREWG1CLKPENGCEDNWKEKNEQFGSCYQFNTQTALSW 240

OGTA076b (SEQ ID No: 2) KEAVYSCQNOGADLLISNSAAEELTYLKEKEG1AKIFWICLNLQLYSARGWEWSDHKPLNLF 300
OGTA076c (SEQ ID No: 3) KEAVYSCQNOGADLLISNSAAEELTYLKEKEG1AKIFWICLNLQLYSARGWEWSDHKPLNLF 300
OGTA076a (SEQ ID No: 1) KEAVYSCQNOGADLLISNSAAEELTYLKEKEG1AKIFWICLNLQLYSARGWEWSDHKPLNLF 300

OGTA076b (SEQ ID No: 2) NWDPDRPSAPTIIGGSSCARMDAESGLWQSFSCSEAQLPYVCRKFLNNIVELTDWVWYSDTR 360
OGTA076c (SEQ ID No: 3) NWDPDRPSAPTIIGGSSCARMDAESGLWQSFSCSEAQLPYVCRKFLNNIVELTDWVWYSDTR 360
OGTA076a (SEQ ID No: 1) NWDPDRPSAPTIIGGSSCARMDAESGLWQSFSCSEAQLPYVCRKFLNNIVELTDWVWYSDTR 360

OGTA076b (SEQ ID No: 2) CDAGWLPNNGPCYLLVNESNWDKAHAKCKAFSSDLISIHSLADVEVVVTKLHNEDIKE 420
OGTA076c (SEQ ID No: 3) CDAGWLPNNGPCYLLVNESNWDKAHAKCKAFSSDLISIHSLADVEVVVTKLHNEDIKE 420
OGTA076a (SEQ ID No: 1) CDAGWLPNNGPCYLLVNESNWDKAHAKCKAFSSDLISIHSLADVEVVVTKLHNEDIKE 420

OGTA076b (SEQ ID No: 2) YVYELKININIPITPLFOWSDETEVITYDENEENPVYNYKTPMCVSYLGLICQNKVOSCEEK 480
OGTA076c (SEQ ID No: 3) YVYELKININIPITPLFOWSDETEVITYDENEENPVYNYKTPMCVSYLGLICQNKVOSCEEK 480
OGTA076a (SEQ ID No: 1) YVYELKININIPITPLFOWSDETEVITYDENEENPVYNYKTPMCVSYLGLICQNKVOSCEEK 480

OGTA076b (SEQ ID No: 2) LKIVCKRKKG2EDPDEGMRGKREGCTYIYD5VPGTNCNLITITSRFFQZY 540
OGTA076c (SEQ ID No: 3) LKIVCKRKKG2EDPDEGMRGKREGCTYIYD5VPGTNCNLITITSRFFQZY 540
OGTA076a (SEQ ID No: 1) LKIVCKRKKG2EDPDEGMRGKREGCTYIYD5VPGTNCNLITITSRFFQZY 540

OGTA076b (SEQ ID No: 2) LNDLMKKYDKSLRYKTYTGLRDVDSCGEYNNWATVGGRRRAVTSNNNLEPASPAGCVM 600
OGTA076c (SEQ ID No: 3) LNDLMKKYDKSLRYKTYTGLRDVDSCGEYNNWATVGGRRRAVTSNNNLEPASPAGCVM 600
OGTA076a (SEQ ID No: 1) LNDLMKKYDKSLRYKTYTGLRDVDSCGEYNNWATVGGRRRAVTSNNNLEPASPAGCVM 600

OGTA076b (SEQ ID No: 2) STGKSVGKWEVKDCRSFKALSICKMNSGPIGPEEASPKPDDPCPREGWQSPASLSCYKVF 660
OGTA076c (SEQ ID No: 3) STGKSVGKWEVKDCRSFKALSICKMNSGPIGPEEASPKPDDPCPREGWQSPASLSCYKVF 660
OGTA076a (SEQ ID No: 1) STGKSVGKWEVKDCRSFKALSICKMNSGPIGPEEASPKPDDPCPREGWQSPASLSCYKVF 660

OGTA076b (SEQ ID No: 2) HAEIRVRKRNWEEAERPCQALGAHLSSFSHVDEIKEFLHPLITEQPSGQHNLWIGLNKRSP 720
 OGTA076c (SEQ ID No: 3) HAEIRVRKRNWEEAERPCQALGAHLSSFSHVDEIKEFLHPLITEQPSGQHNLWIGLNKRSP 720
 OGTA076a (SEQ ID No: 1) HAEIRVRKRNWEEAERPCQALGAHLSSFSHVDEIKEFLHPLITEQPSGQHNLWIGLNKRSP 720

 OGTA076b (SEQ ID No: 2) DLQGSWQWSDRTPVSTIIMPHEPQQDYDIRDCAAVKVFHRPWRPGRHIFYDPRFIFIYLRF 780
 OGTA076c (SEQ ID No: 3) DLQGSWQWSDRTPVSTIIMPHEPQQDYDIRDCAAVKVFHRPWRPGRHIFYDPRFIFIYLRF 780
 OGTA076a (SEQ ID No: 1) DLQGSWQWSDRTPVSTIIMPHEPQQDYDIRDCAAVKVFHRPWRPGRHIFYDPRFIFIYLRF 780

 OGTA076b (SEQ ID No: 2) ACDTKLE8WVCQIPKGRTPKTHEDWVNRDAGIHGPPLIIEGSEYWFADLHILNYYEAVLYC 840
 OGTA076c (SEQ ID No: 3) ACDTKLE8WVCQIPKGRTPKTHEDWVNRDAGIHGPPLIIEGSEYWFADLHILNYYEAVLYC 840
 OGTA076a (SEQ ID No: 1) ACDTKLE8WVCQIPKGRTPKTHEDWVNRDAGIHGPPLIIEGSEYWFADLHILNYYEAVLYC 840

 OGTA076b (SEQ ID No: 2) ASNHSPLATITSFVGLKAIRNKIANISGDDQKWWIRISEWPIDDDHTYSRYPWHRFPVTF 900
 OGTA076c (SEQ ID No: 3) ASNHSPLATITSFVGLKAIRNKIANISGDDQKWWIRISEWPIDDDHTYSRYPWHRFPVTF 900
 OGTA076a (SEQ ID No: 1) ASNHSPLATITSFVGLKAIRNKIANISGDDQKWWIRISEWPIDDDHTYSRYPWHRFPVTF 900

 OGTA076b (SEQ ID No: 2) GEECLYMSAKTWLIDLGKPTDCSTKTHEDWVNRDAGIHGPPLIIEGSEYWFADLHILNYYEAVLYC 960
 OGTA076c (SEQ ID No: 3) GEECLYMSAKTWLIDLGKPTDCSTKTHEDWVNRDAGIHGPPLIIEGSEYWFADLHILNYYEAVLYC 960
 OGTA076a (SEQ ID No: 1) GEECLYMSAKTWLIDLGKPTDCSTKTHEDWVNRDAGIHGPPLIIEGSEYWFADLHILNYYEAVLYC 960

 OGTA076b (SEQ ID No: 2) KCFLIKIKPVSLSFQASDTCHSYGGTLPSVLSQIFQDFITSLLPIMEATLWIGLWRAYE 1020
 OGTA076c (SEQ ID No: 3) KCFLIKIKPVSLSFQASDTCHSYGGTLPSVLSQIFQDFITSLLPIMEATLWIGLWRAYE 1020
 OGTA076a (SEQ ID No: 1) KCFLIKIKPVSLSFQASDTCHSYGGTLPSVLSQIFQDFITSLLPIMEATLWIGLWRAYE 1020

 OGTA076b (SEQ ID No: 2) KINKWTDNRELITYSNFPHPLLVSGRLRTHEDWVNRDAGIHGPPLIIEGSEYHCALILNQKSPPTCTWNFTSC 1080
 OGTA076c (SEQ ID No: 3) KINKWTDNRELITYSNFPHPLLVSGRLRTHEDWVNRDAGIHGPPLIIEGSEYHCALILNQKSPPTCTWNFTSC 1080
 OGTA076a (SEQ ID No: 1) KINKWTDNRELITYSNFPHPLLVSGRLRTHEDWVNRDAGIHGPPLIIEGSEYHCALILNQKSPPTCTWNFTSC 1080

 OGTA076b (SEQ ID No: 2) SERHEVSIICORYSEVKRSQTYLQASSETVYKLNNYYKIIPKTHEDWVNRDAGIHGPPLIIEGSEYRECLKSNNQI VS 1140
 OGTA076c (SEQ ID No: 3) SERHEVSIICORYSEVKRSQTYLQASSETVYKLNNYYKIIPKTHEDWVNRDAGIHGPPLIIEGSEYRECLKSNNQI VS 1140
 OGTA076a (SEQ ID No: 1) SERHEVSIICORYSEVKRSQTYLQASSETVYKLNNYYKIIPKTHEDWVNRDAGIHGPPLIIEGSEYRECLKSNNQI VS 1140

 OGTA076b (SEQ ID No: 2) ITDPYQQAFLSVQALLHNSLWIGLFSQDDELNFGWDCKRLHPSRWAETNGQLEDCVVI 1200
 OGTA076c (SEQ ID No: 3) ITDPYQQAFLSVQALLHNSLWIGLFSQDDELNFGWDCKRLHPSRWAETNGQLEDCVVI 1200
 OGTA076a (SEQ ID No: 1) ITDPYQQAFLSVQALLHNSLWIGLFSQDDELNFGWDCKRLHPSRWAETNGQLEDCVVI 1200

 OGTA076b (SEQ ID No: 2) CTDGFWKTVDCNDNQPGAIACYSGNETEKTHEDWVNRDAGIHGPPLIIEGSEYRECLKSNNQI VS 1260
 OGTA076c (SEQ ID No: 3) CTDGFWKTVDCNDNQPGAIACYSGNETEKTHEDWVNRDAGIHGPPLIIEGSEYRECLKSNNQI VS 1260
 OGTA076a (SEQ ID No: 1) CTDGFWKTVDCNDNQPGAIACYSGNETEKTHEDWVNRDAGIHGPPLIIEGSEYRECLKSNNQI VS 1260

 OGTA076b (SEQ ID No: 2) TKNRHIMATTQDEVHTKCKQKLNPKSHIILSIRDKENNFVLEGGLYFNYMASWVMLGITYR 1320
 OGTA076c (SEQ ID No: 3) TKNRHIMATTQDEVHTKCKQKLNPKSHIILSIRDKENNFVLEGGLYFNYMASWVMLGITYR 1320
 OGTA076a (SEQ ID No: 1) TKNRHIMATTQDEVHTKCKQKLNPKSHIILSIRDKENNFVLEGGLYFNYMASWVMLGITYR 1320

 OGTA076b (SEQ ID No: 2) *****TPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDICQTFKVIEAVYFHQHSILAC 1380
 OGTA076c (SEQ ID No: 3) *****TPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDICQTFKVIEAVYFHQHSILAC 1380
 OGTA076a (SEQ ID No: 1) *****TPLSYTHWRAGRPTIKNEKFLAGLSTDGFWDICQTFKVIEAVYFHQHSILAC 1380

 OGTA076b (SEQ ID No: 2) KIEMVDYKEEHTTILQPMYEDGIVS10KRVTWYALNMCSSGHLASVHNONGOLF 1440
 OGTA076c (SEQ ID No: 3) KIEMVDYKEEHTTILQPMYEDGIVS10KRVTWYALNMCSSGHLASVHNONGOLF 1440
 OGTA076a (SEQ ID No: 1) KIEMVDYKEEHTTILQPMYEDGIVS10KRVTWYALNMCSSGHLASVHNONGOLF 1440

 OGTA076b (SEQ ID No: 2) LEDIVKRDCGPFLWVGLSHGDSESSFEWNSDGSFTFDYIPWKGQTSPGNCVLLDPKGCTWKHE 1500
 OGTA076c (SEQ ID No: 3) LEDIVKRDCGPFLWVGLSHGDSESSFEWNSDGSFTFDYIPWKGQTSPGNCVLLDPKGCTWKHE 1500
 OGTA076a (SEQ ID No: 1) LEDIVKRDCGPFLWVGLSHDCSESSFEWSDCSTFDYIPWKGQTSPGNCVLLDPKGCTWKHE 1500

 OGTA076b (SEQ ID No: 2) KCNSVKDCAICXKPTTSKKLRSRITYSSRCPAAKENGSRWICYKGCHYKSDQALHSFSEAK 1560
 OGTA076c (SEQ ID No: 3) KCNSVKDCAICXKPTTSKKLRSRITYSSRCPAAKENGSRWICYKGCHYKSDQALHSFSEAK 1560
 OGTA076a (SEQ ID No: 1) KCNSVKDCAICXKPTTSKKLRSRITYSSRCPAAKENGSRWICYKGCHYKSDQALHSFSEAK 1560

OGTA076b (SEQ ID No: 2) KLCSKHDEHSATIVSIXDEDENKPVSRIMRENNNTM-----WKLGLSQHSVVDQSWNLDGSEVT 1620
 OGTA076c (SEQ ID No: 3) KLCSKHDEHSATIVSIXDEDENKPVSRIMRENNNTM-----WKLGLSQHSVVDQSWNLDGSEVT 1608
 OGTA076a (SEQ ID No: 1) KLCSKHDEHSATIVSIXDEDENKPVSRIMRENNNTM-----WKLGLSQHSVVDQSWNLDGSEVT 1608

 OGTA076b (SEQ ID No: 2) FVKWENKSESGVGRCMSMLIASNETWKKVECEHGFGRVVCKVPIOCPSSTWIQFQDSCYIF 1680
 OGTA076c (SEQ ID No: 3) FVKWENKSESGVGRCMSMLIASNETWKKVECEHGFGRVVCKVPIOCPSSTWIQFQDSCYIF 1624
 OGTA076a (SEQ ID No: 1) FVKWENKSESGVGRCMSMLIASNETWKKVECEHGFGRVVCKVPIOCPSSTWIQFQDSCYIF 1608

 OGTA076b (SEQ ID No: 2) IQRAIKVESTEDVVRNQCTDHADMISTHNRRENAP-----ILKKQWKGPDDTLGMFYDDTDD 1740
 OGTA076c (SEQ ID No: 3) IQRAIKVESTEDVVRNQCTDHADMISTHNRRENAP-----ILKKQWKGPDDTLGMFYDDTDD 1684
 OGTA076a (SEQ ID No: 1) IQRAIKVESTEDVVRNQCTDHADMISTHNRRENAP-----ILKKQWKGPDDTLGMFYDDTDD 1608

 OGTA076b (SEQ ID No: 2) ÅSEKWFDNSNMTFDKWTDQDDDEDLVDTCAFLHIKVGEWKKGNCEVSSVEGTLCKTAIPY 1800
 OGTA076c (SEQ ID No: 3) ÅSEKWFDNSNMTFDKWTDQDDDEDLVDTCAFLHIKVGEWKKGNCEVSSVEGTLCKTAIPY 1744
 OGTA076a (SEQ ID No: 1) ÅSEKWFDNSNMTFDKWTDQDDDEDLVDTCAFLHIKVGEWKKGNCEVSSVEGTLCKTAIPY 1804

 OGTA076b (SEQ ID No: 2) VVGEENEYPVQPD 1873
 OGTA076c (SEQ ID No: 3) VVGEENEYPVQPD 1817
 OGTA076a (SEQ ID No: 1) VVGEENEYPVQPD 1804

Mass Match Peptides (bold):

CLGLDITK [7]
 DGAICYKPK [9]
 DVDSCEYNWATGGRRR [11]
 EEWVIGLK [12]
 ENNNITMR [16]
 EVKPVDSVK [17]
 GHWFYDDR [21]
 HDHSATIVSIK [22]
 HFVSLCQK [23]
 HGETCYK [24]
 IPENFEEEEESR [29]
 KYFWTGLR [34]
 LNDASSDK [39]
 LPFICEK [41]
 MCPPDEGWKR [42]
 MFSCDSSAMLWW [43]
 NNSLWFDK [45]
 QTLQNASETVK [47]
 RHGETCYK [49]
 TLTWISAK [56]
 TPDWYNPDR [57]

Tandem Peptides (underline):

IPENFEEEEESR [29]

Peptide Source: 1D GE Pancreatic cancer

OGTA076b (SEQ ID No: 2) MRTGWATPRRPAGLIMLLEWFFDLAEPSGRAANDPFTIVHGNTGKCIKPKVYGWIVADDCD 60
 OGTA076c (SEQ ID No: 3) MRTGWATPRRPAGLIMLLEWFFDLAEPSGRAANDPFTIVHGNTGKCIKPKVYGWIVADDCD 60
 OGTA076a (SEQ ID No: 1) MRTGWATPRRPAGLIMLLEWFFDLAEPSGRAANDPFTIVHGNTGKCIKPKVYGWIVADDCD 60

 OGTA076b (SEQ ID No: 2) ETEDKLWKWVSCQKSELFLHLQKEIGLDITKSVNELRMFCCSSAMLWHKCEHHGSLYGAAR 120
 OGTA076c (SEQ ID No: 3) ETEDKLWKWVSCQKSELFLHLQKEIGLDITKSVNELRMFCCSSAMLWHKCEHHGSLYGAAR 120

OGTA076a (SEQ ID No: 1)
 ETEDKILWIKVYSGRLPHLHSOKCEGEDITKVSNEIRMFSCDSSAMILNNKCEHHSLYGRAR 120

OGTA076b (SEQ ID No: 2)
 YRLAIDGCGTATGNSASDQHGGSEESLICDQPHIEIYTRDGNSTGRPCCEPPFLIDGTWH 180
 OGTA076c (SEQ ID No: 3)
 YRLAIDGCGTATGNSASDQHGGSEESLICDQPHIEIYTRDGNSTGRPCCEPPFLIDGTWH 180
 OGTA076a (SEQ ID No: 1)
 YRLAIDGCGTATGNSASDQHGGSEESLICDQPHIEIYTRDGNSTGRPCCEPPFLIDGTWH 180

OGTA076b (SEQ ID No: 2)
 HICLIDDEDISGPWCATINNEYDRKGICLKPENGCEDNNEKNEQFSCSYQFNTQALSW 240
 OGTA076c (SEQ ID No: 3)
 HICLIDDEDISGPWCATINNEYDRKGICLKPENGCEDNWEKNEQFSCSYQFNTQALSW 240
 OGTA076a (SEQ ID No: 1)
 HICLIDDEDISGPWCATINNEYDRKGICLKPENGCEDNWEKNEQFSCSYQFNTQALSW 240

OGTA076b (SEQ ID No: 2)
 KEAVVSCQNQGADLLINSAAEITYLKEKEGIAKIPIWIGLNLQLYSARGNEWDHKFLNLF 300
 OGTA076c (SEQ ID No: 3)
 KEAVVSCQNQGADLLINSAAEITYLKEKEGIAKIPIWIGLNLQLYSARGNEWDHKFLNLF 300
 OGTA076a (SEQ ID No: 1)
 KEAVVSCQNQGADLLINSAAEITYLKEKEGIAKIPIWIGLNLQLYSARGNEWDHKFLNLF 300

OGTA076b (SEQ ID No: 2)
 NWDPDRPSAPTIIGGSSCARMdASGLMQSPSCBQLPVCRKFLNNTVLTDTWVTSDFR 360
 OGTA076c (SEQ ID No: 3)
 NWDPDRPSAPTIIGGSSCARMdASGLMQSPSCBQLPVCRKFLNNTVLTDTWVTSDFR 360
 OGTA076a (SEQ ID No: 1)
 NWDPDRPSAPTIIGGSSCARMdASGLMQSPSCBQLPVCRKFLNNTVLTDTWVTSDFR 360

OGTA076b (SEQ ID No: 2)
 CDACWLPNNGPCYIIVNFSNSNDKAHAKC2KESDHTIHSIADVEVYVTKHNNEDIKEK 420
 OGTA076c (SEQ ID No: 3)
 CDAGWLPNNGPCYIIVNFSNSNDKAHAKC2KESDHTIHSIADVEVYVTKHNNEDIKEK 420
 OGTA076a (SEQ ID No: 1)
 CDAGWLPNNGPCYIIVNFSNSNDKAHAKC2KESDHTIHSIADVEVYVTKHNNEDIKEK 420

OGTA076b (SEQ ID No: 2)
 VVTCIENINIPITLFWQNSDGTETVLTLYDENEPIVNPYNTIPNCVSYLGEIGQWKVQSCKEK 480
 OGTA076c (SEQ ID No: 3)
 VVTCIENINIPITLFWQNSDGTETVLTLYDENEPIVNPYNTIPNCVSYLGEIGQWKVQSCKEK 480
 OGTA076a (SEQ ID No: 1)
 VVTCIENINIPITLFWQNSDGTETVLTLYDENEPIVNPYNTIPNCVSYLGEIGQWKVQSCKEK 480

OGTA076b (SEQ ID No: 2)
 LKVCKRKGEELNDACGDKMCPPDBGKHEKESDHTIHSIADVEVYDVEFGTMCNLITTSDFEY 540
 OGTA076c (SEQ ID No: 3)
 LKVCKRKGEELNDACGDKMCPPDBGKHEKESDHTIHSIADVEVYDVEFGTMCNLITTSDFEY 540
 OGTA076a (SEQ ID No: 1)
 LKVCKRKGEELNDACGDKMCPPDBGKHEKESDHTIHSIADVEVYDVEFGTMCNLITTSDFEY 540

OGTA076b (SEQ ID No: 2)
 LNDLNRKYLNSLRLKESDHTIHSIADVEVYDVEFGTMCNLITTSDFEY 600
 OGTA076c (SEQ ID No: 3)
 LNDLNRKYLNSLRLKESDHTIHSIADVEVYDVEFGTMCNLITTSDFEY 600
 OGTA076a (SEQ ID No: 1)
 LNDLNRKYLNSLRLKESDHTIHSIADVEVYDVEFGTMCNLITTSDFEY 600

OGTA076b (SEQ ID No: 2)
 STGKSVGKWEVKDCRSFKALSIICKMNSGEPFEEERASPKPDCPCPGWQGQSPFASLSCYKV 660
 OGTA076c (SEQ ID No: 3)
 STGKSVGKWEVKDCRSFKALSIICKMNSGEPFEEERASPKPDCPCPGWQGQSPFASLSCYKV 660
 OGTA076a (SEQ ID No: 1)
 STGKSVGKWEVKDCRSFKALSIICKMNSGEPFEEERASPKPDCPCPGWQGQSPFASLSCYKV 660

OGTA076b (SEQ ID No: 2)
 HARERIVRKRNNEEAEKFQDQALGAHLSSEFSHVDEIKEFELHFIITQFSCQHMLWIGLNLKRSF 720
 OGTA076c (SEQ ID No: 3)
 HARERIVRKRNNEEAEKFQDQALGAHLSSEFSHVDEIKEFELHFIITQFSCQHMLWIGLNLKRSF 720
 OGTA076a (SEQ ID No: 1)
 HARERIVRKRNNEEAEKFQDQALGAHLSSEFSHVDEIKEFELHFIITQFSCQHMLWIGLNLKRSF 720

OGTA076b (SEQ ID No: 2)
 DIOGSNQSDTETVLTLYDENEPIVNPYNTIPNCVSYLGEIGQWKVQSCKEK 780
 OGTA076c (SEQ ID No: 3)
 DIOGSNQSDTETVLTLYDENEPIVNPYNTIPNCVSYLGEIGQWKVQSCKEK 780
 OGTA076a (SEQ ID No: 1)
 DIOGSNQSDTETVLTLYDENEPIVNPYNTIPNCVSYLGEIGQWKVQSCKEK 780

OGTA076b (SEQ ID No: 2)
 ACDDLEWVWCQ1PKGRTPKIDPDTWVNPYNTIPNCVSYLGEIGQWKVQSCKEK 840
 OGTA076c (SEQ ID No: 3)
 ACDDLEWVWCQ1PKGRTPKIDPDTWVNPYNTIPNCVSYLGEIGQWKVQSCKEK 840
 OGTA076a (SEQ ID No: 1)
 ACDDLEWVWCQ1PKGRTPKIDPDTWVNPYNTIPNCVSYLGEIGQWKVQSCKEK 840

OGTA076b (SEQ ID No: 2)
 ASNHSFLATITTSFVG1KAIKNN1AN1SGDCKWVIR1SENEIDDEPTYSRYPWREPEK 900
 OGTA076c (SEQ ID No: 3)
 ASNHSFLATITTSFVG1KAIKNN1AN1SGDCKWVIR1SENEIDDEPTYSRYPWREPEK 900
 OGTA076a (SEQ ID No: 1)
 ASNHSFLATITTSFVG1KAIKNN1AN1SGDCKWVIR1SENEIDDEPTYSRYPWREPEK 900

OGTA076b (SEQ ID No: 2)
 KCFLKIKPVSLTFSQASDTCHSYGGTLPVSLSQIFQDFITSLLPDMEARLWIGLRTAYE 1020
 OGTA076c (SEQ ID No: 3)
 KCFLKIKPVSLTFSQASDTCHSYGGTLPVSLSQIFQDFITSLLPDMEARLWIGLRTAYE 1020

OGTA076a (SEQ ID No: 1)	KCFLKIKPVSLTFSQASDTCHSYGGTLPVLSQIIPQDFITSLLPDMEATIWIIGLRLWTAYE 1020
OGTA076b (SEQ ID No: 2)	KINKWTNDRELTYSHEPFLVSYSLERLIPENFEERESRHYCALILNLQKSPPTGTWNFTSC 1080
OGTA076c (SEQ ID No: 3)	KINKWTNDRELTYSHEPFLVSYSLERLIPENFEERESRHYCALILNLQKSPPTGTWNFTSC 1080
OGTA076a (SEQ ID No: 1)	KINKWTNDRELTYSHEPFLVSYSLERLIPENFEERESRHYCALILNLQKSPPTGTWNFTSC 1080
OGTA076b (SEQ ID No: 2)	SERPHFVSYLQKSYSEVKRSRQTLQNASETVKYLINNNLYKIPKTTTREHSAECLIKSNQMLVS 1140
OGTA076c (SEQ ID No: 3)	SERPHFVSYLQKSYSEVKRSRQTLQNASETVKYLINNNLYKIPKTTTREHSAECLIKSNQMLVS 1140
OGTA076a (SEQ ID No: 1)	SERPHFVSYLQKSYSEVKRSRQTLQNASETVKYLINNNLYKIPKTTTREHSAECLIKSNQMLVS 1140
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	ITDDQYQAFQALSVQALLNNNSLWIGLFSQDDEFLNGFWEDGCRHLHESRWAETNGQLEDCVVL 1200
OGTA076c (SEQ ID No: 3)	ITDDQYQAFQALSVQALLNNNSLWIGLFSQDDEFLNGFWEDGCRHLHESRWAETNGQLEDCVVL 1200
OGTA076a (SEQ ID No: 1)	ITDDQYQAFQALSVQALLNNNSLWIGLFSQDDEFLNGFWEDGCRHLHESRWAETNGQLEDCVVL 1200
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	DTDGFWKTVDCDNQPGAIICYSGNTEKBEVKPVDSVKCPSFVNTPVLFQNCQCCYNFII 1260
OGTA076c (SEQ ID No: 3)	DTDGFWKTVDCDNQPGAIICYSGNTEKBEVKPVDSVKCPSFVNTPVLFQNCQCCYNFII 1260
OGTA076a (SEQ ID No: 1)	DTDGFWKTVDCDNQPGAIICYSGNTEKBEVKPVDSVKCPSFVNTPVLFQNCQCCYNFII 1260
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	TKRHRHDTTTCVTCQKLNPKHSEHSEDEPKNPNNPVLFGIYTFYNYMARSWVMILGTYRN 1320
OGTA076c (SEQ ID No: 3)	TKRHRHDTTTCVTCQKLNPKHSEHSEDEPKNPNNPVLFGIYTFYNYMARSWVMILGTYRN 1320
OGTA076a (SEQ ID No: 1)	TKRHRHDTTTCVTCQKLNPKHSEHSEDEPKNPNNPVLFGIYTFYNYMARSWVMILGTYRN 1320
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	NSLIMMEDPTLYSYTHNSLWIGRPTIINKSKFLPLAGLSTDGFWDICTEYKTYEAEVHESBILAC 1380
OGTA076c (SEQ ID No: 3)	NSLIMMEDPTLYSYTHNSLWIGRPTIINKSKFLPLAGLSTDGFWDICTEYKTYEAEVHESBILAC 1380
OGTA076a (SEQ ID No: 1)	NSLIMMEDPTLYSYTHNSLWIGRPTIINKSKFLPLAGLSTDGFWDICTEYKTYEAEVHESBILAC 1380
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	KIEMVDYKEEHNNTLPQFMPYEDGIVYQKVKTYGFLALNMCQSGGHHLASVHNQNCQLF 1440
OGTA076c (SEQ ID No: 3)	KIEMVDYKEEHNNTLPQFMPYEDGIVYQKVKTYGFLALNMCQSGGHHLASVHNQNCQLF 1440
OGTA076a (SEQ ID No: 1)	KIEMVDYKEEHNNTLPQFMPYEDGIVYQKVKTYGFLALNMCQSGGHHLASVHNQNCQLF 1440
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	LEDIVKRDFGFLWLGVLSHSDGESSFEWSGDSTFDYIIPWKGQTSPGNCVLLDPKGTKWHE 1500
OGTA076c (SEQ ID No: 3)	LEDIVKRDFGFLWLGVLSHSDGESSFEWSGDSTFDYIIPWKGQTSPGNCVLLDPKGTKWHE 1500
OGTA076a (SEQ ID No: 1)	LEDIVKRDFGFLWLGVLSHSDGESSFEWSGDSTFDYIIPWKGQTSPGNCVLLDPKGTKWHE 1500
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	KCNSVKDGTCTCTKPSKSLRSLITYSSRCPAAKENGSRNLWYKCHGHCYNSDQALNRESEK 1560
OGTA076c (SEQ ID No: 3)	KCNSVKDGTCTCTKPSKSLRSLITYSSRCPAAKENGSRNLWYKCHGHCYNSDQALNRESEK 1560
OGTA076a (SEQ ID No: 1)	KCNSVKDGTCTCTKPSKSLRSLITYSSRCPAAKENGSRNLWYKCHGHCYNSDQALNRESEK 1560
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	KLCSDKHDHSATIVS1K1DE0DENDKPVSRSLMRNNNT1MVRWLGLSOSHVSQWSWLDGSEVT 1620
OGTA076c (SEQ ID No: 3)	KLCSDKHDHSATIVS1K1DE0DENDKPVSRSLMRNNNT1MVRWLGLSOSHVSQWSWLDGSEVT 1608
OGTA076a (SEQ ID No: 1)	KLCSDKHDHSATIVS1K1DE0DENDKPVSRSLMRNNNT1MVRWLGLSOSHVSQWSWLDGSEVT 1608
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	PVKWENKSKSKGVRCSMLIASNSETWRKLYCETECEHGFGRVVKVFLDCPSSTWIIPQFQDCSYI 1680
OGTA076c (SEQ ID No: 3)	-----CPSSTWIIPQFQDCSYI 1624
OGTA076a (SEQ ID No: 1)	PVKWENKSKSKGVRCSMLIASNSETWRKLYCETECEHGFGRVVKVFLDCPSSTWIIPQFQDCSYI 1680
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	LQEAIIKVESTEDVRNQCTDHGADMISIHNEEENAFIDLILKKWQKGPDQDILGMYFTD 1740
OGTA076c (SEQ ID No: 3)	LQEAIIKVESTEDVRNQCTDHGADMISIHNEEENAFIDLILKKWQKGPDQDILGMYFTD 1684
OGTA076a (SEQ ID No: 1)	VIMGGLIWFLPQRHLHILAGESSVYTAQGVNEDBIMLPSFHD-----
OGTA076b (SEQ ID No: 2)	*****
OGTA076c (SEQ ID No: 3)	*****
OGTA076a (SEQ ID No: 1)	*****
OGTA076b (SEQ ID No: 2)	ASFKWEDNQNMFTDKWTQDDEDDLVDLTCAPLHIKTGEWKKYGNCEVSSVSECTLQTAIPY 1800
OGTA076c (SEQ ID No: 3)	ASFKWEDNQNMFTDKWTQDDEDDLVDLTCAPLHIKTGEWKKYGNCEVSSVSECTLQTAIPY 1744
OGTA076a (SEQ ID No: 1)	-----
OGTA076b (SEQ ID No: 2)	KRKY1SDNHLILSALVIASIASTVILVGLAIIFWYKHKHSDSRFTTVFSTAPQSPYNECDVL 1860
OGTA076c (SEQ ID No: 3)	KRKY1SDNHLILSALVIASIASTVILVGLAIIFWYKHKHSDSRFTTVFSTAPQSPYNECDVL 1860
OGTA076a (SEQ ID No: 1)	-----
OGTA076b (SEQ ID No: 2)	VVGEENEPYVQFQD 1873
OGTA076c (SEQ ID No: 3)	VVGEENEPYVQFQD 1817
OGTA076a (SEQ ID No: 1)	-----

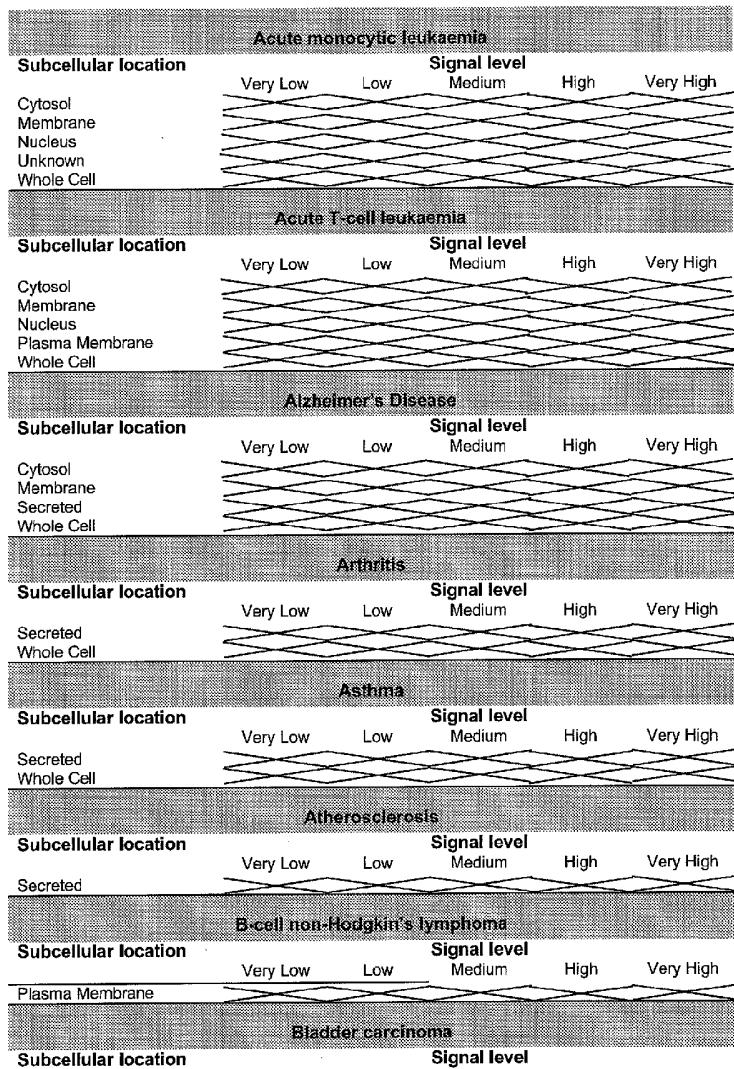
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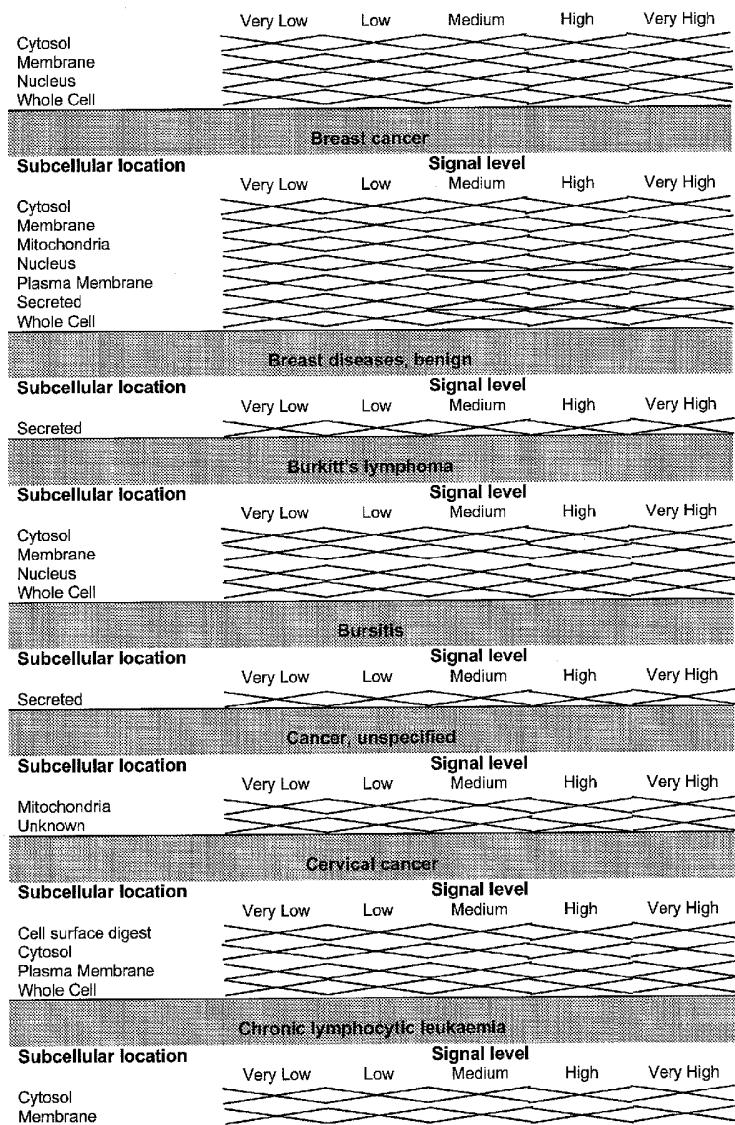
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CLGLDITK [7]
DGAICYKPTK [9]
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DVDSCGEYNWATGGRRR [11]
EEVWIGLK [12]
EFIYLRPACDTK [13]
FEQEYLNLDLK [18]
FPVTGFEECLYMSAK [19]
GNCEVSVSVEGTLCK [20]
GWHFYDDR [21]
HFVSLCQK [23]
HMATTQDEVHTK [25]
IANISGDGQK [26]
IPENFFEEESR [29]
ISEWPIDDHFTYSR [30]
KRNWEEAER [32]
KYFWTGLR [34]
LHLAGFSSVR [37]
NNSLMWFDK [45]
RGWHPYDDR [48]
RHGETCYK [49]
RNWESEAER [51]
SDQALHSFSEAK [52]
SHILSIR [53]
SPDLQGSWQWSDR [55]
TLTWHSAK [56]
TPDWYNPDR [57]
TPLSYTHWR [58]
TPVSTIIMPNEFQQDYDIR [59]
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VFHRPWR [61]
VQCSBQWIPFQNK [63]
WVSQRR [64]
YFWTGGLR [65]

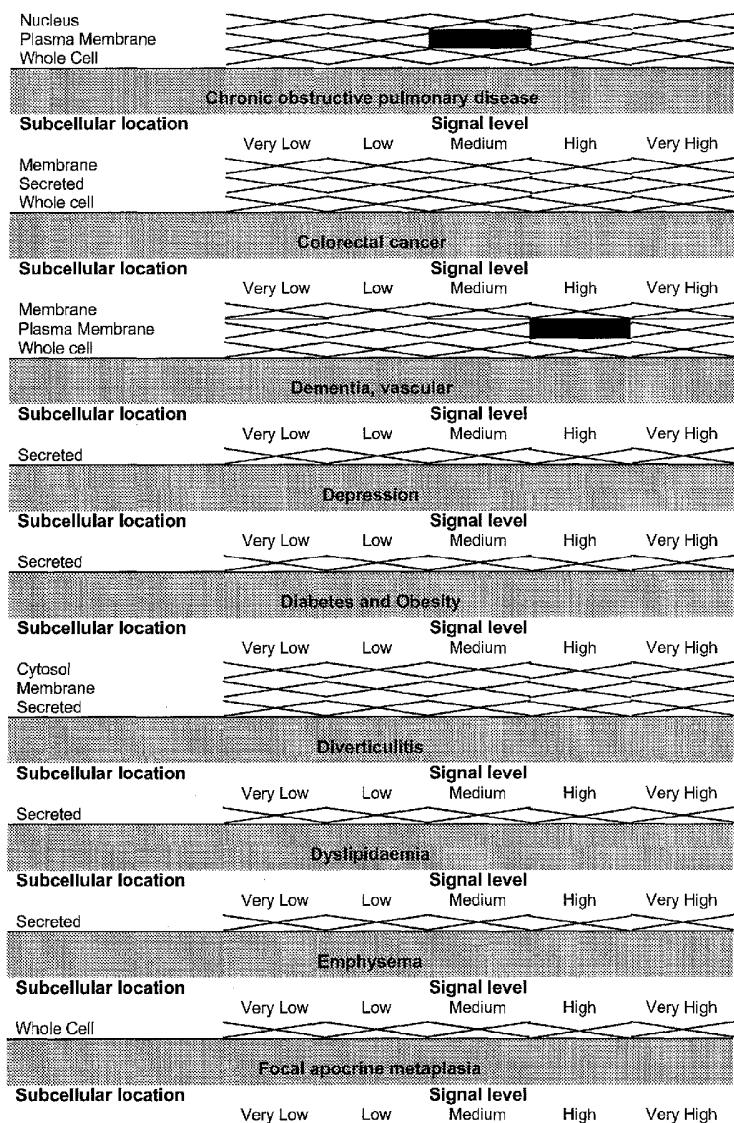
Tandem Peptides (underline):

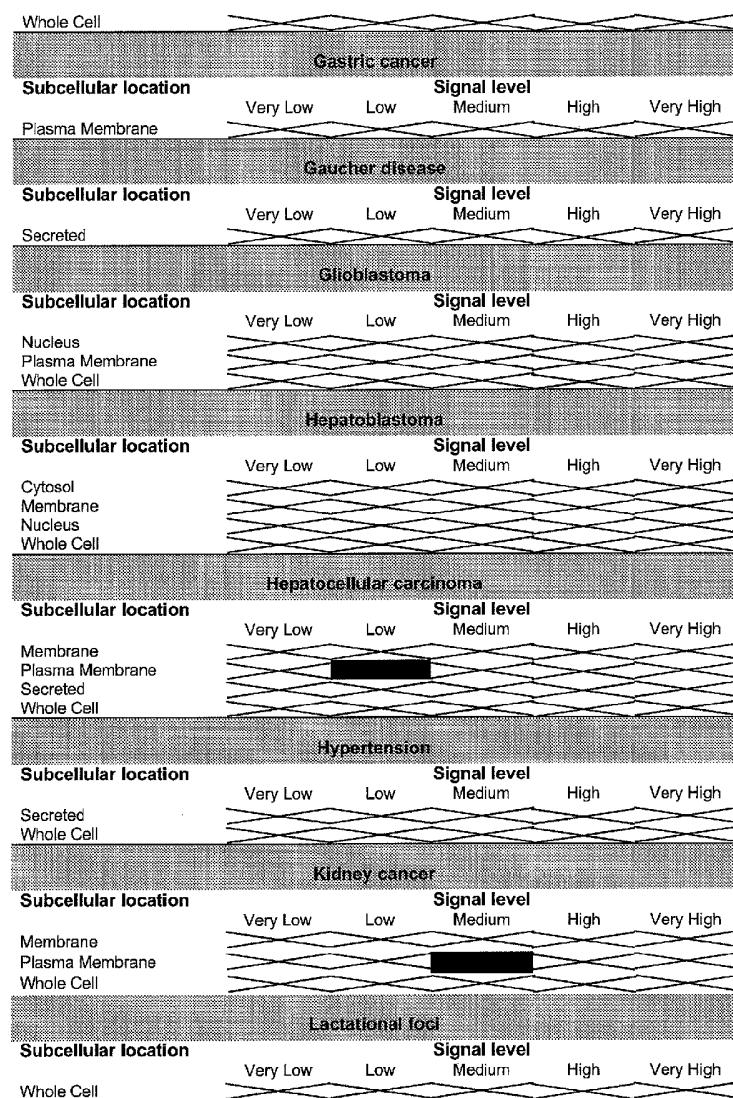
FEQEYLNLDLK [18]
IPENFFEEESR [29]
MSGPLGPEEASPK [44]
SNFHPLLVSGR [54]
SPDLQGSWQWSDR [55]
VIEEAVYFHQH [62]

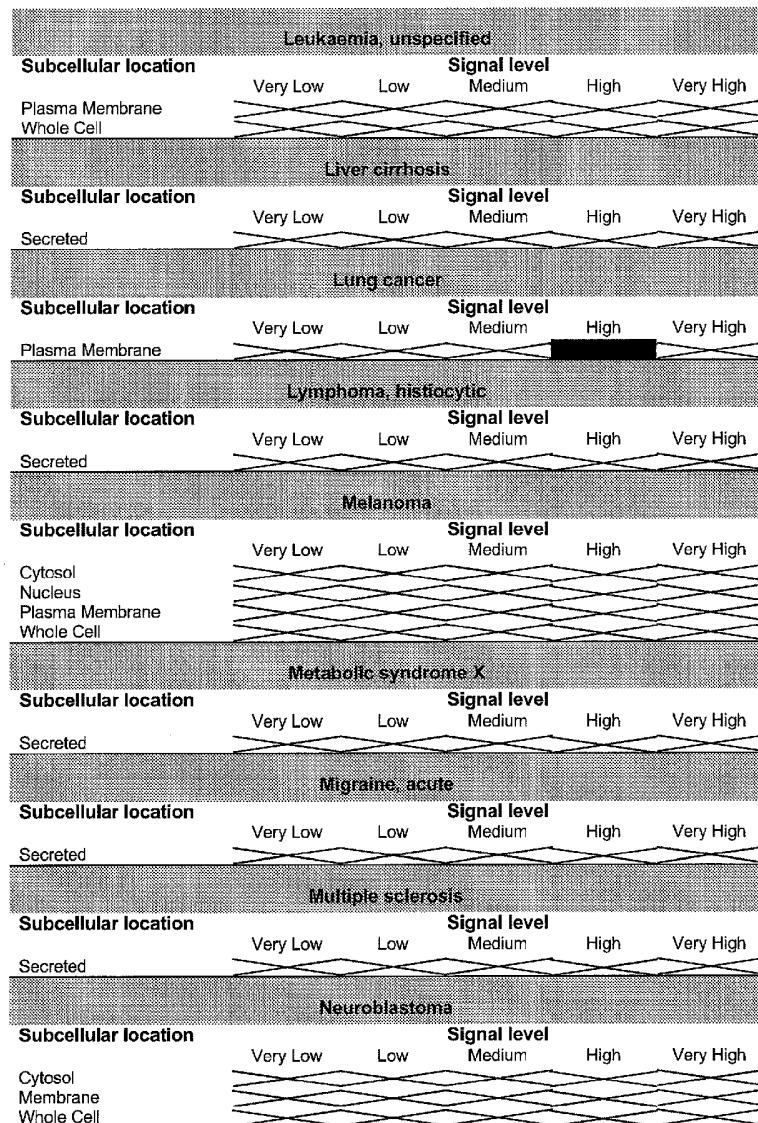
Figure 2

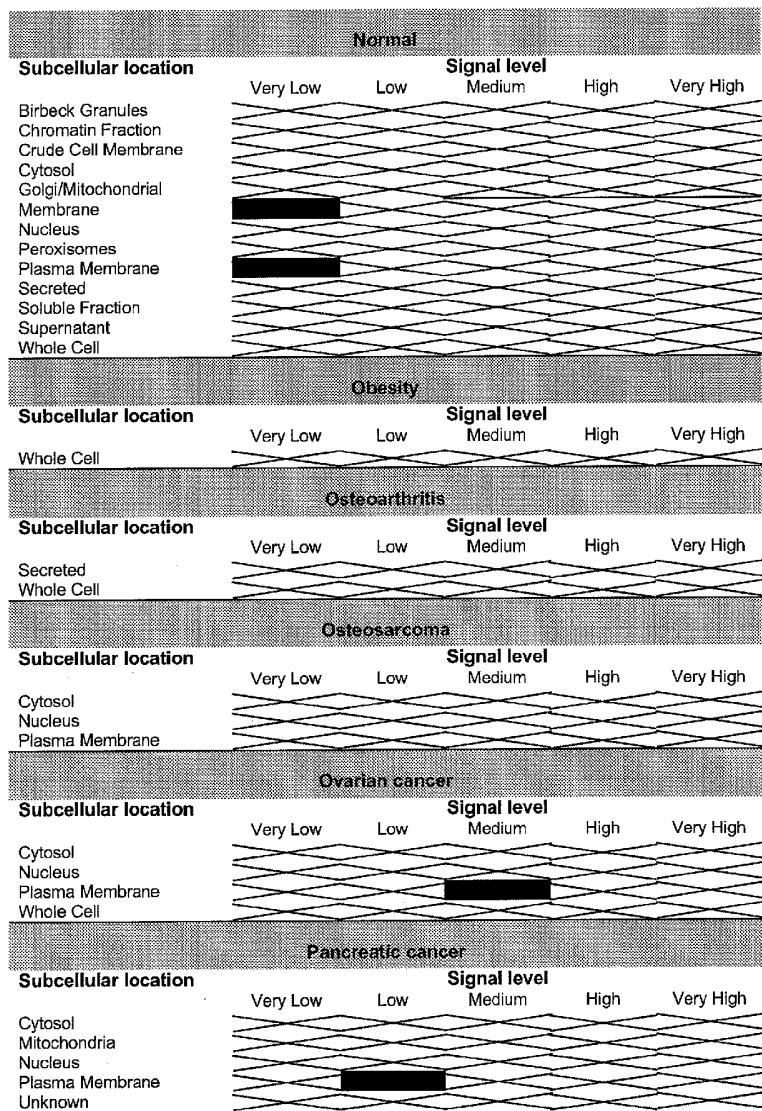


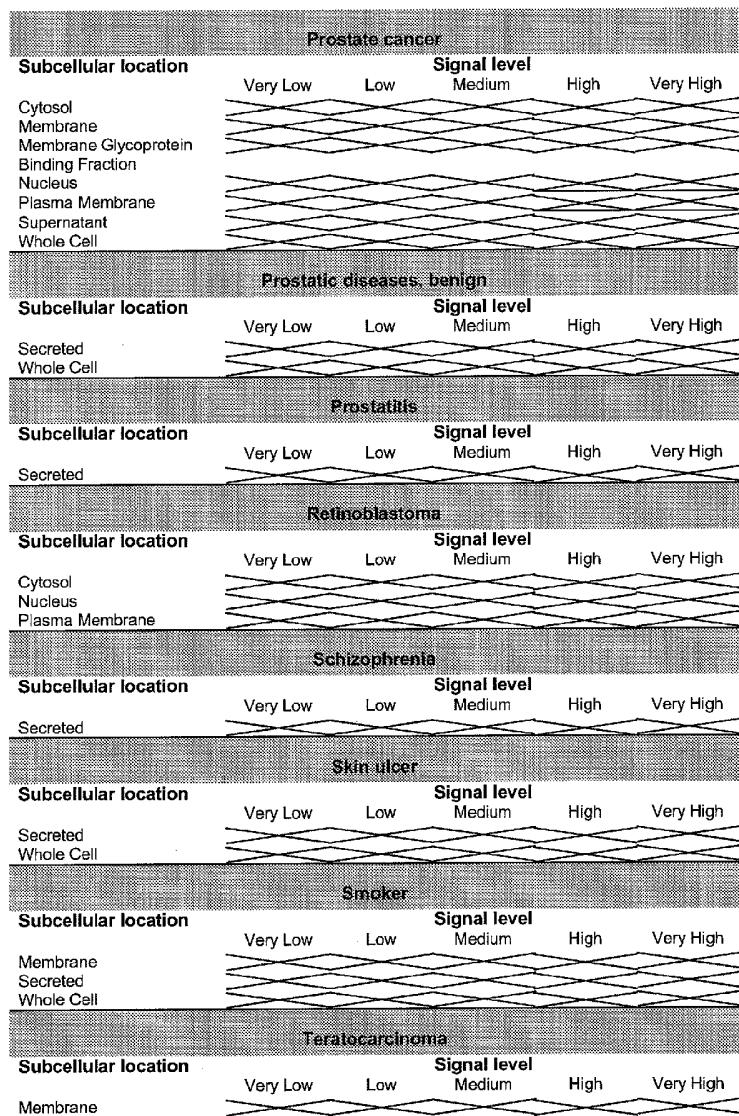












Whole Cell

