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BODY FLUIDS

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(54) QUANTITATIVE ASSAYS FOR RAS P21 IN

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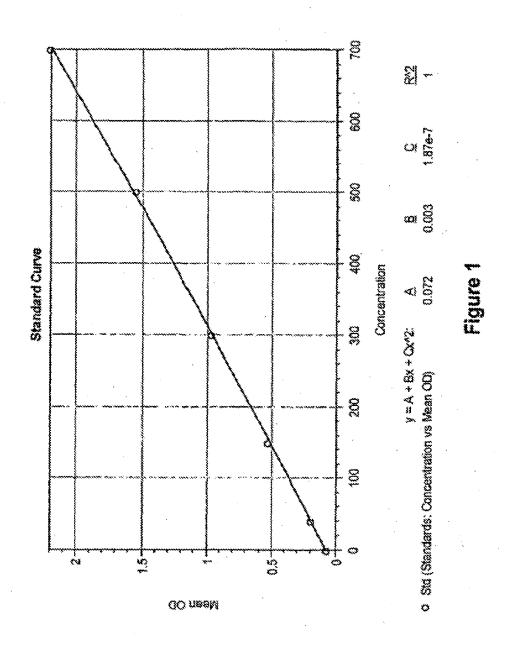
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(57) ABSTRACT

The present invention is directed to the detection and quantification of total ras p21 in body fluids, particularly serial changes of total ras p21 levels in a subject's body fluids. Further, the invention is directed to detecting and quantitating total ras p21 in conjunction with one or more other proteins, such as, oncoproteins, angiogenic factors, tumor markers, inhibitors, growth factor receptors, metastasis proteins, and tumor suppressors. The disclosed methods are diagnostic/prognostic for preneoplastic/neoplastic diseases, and useful to select therapies for patients with preneoplastic/neoplastic diseases. The disclosed methods are further useful to monitor the status of a patient's preneoplastic/neoplastic disease, and/or to monitor how a patient is responding to an anticancer therapy.



QUANTITATIVE ASSAYS FOR RAS P21 IN BODY FLUIDS

FIELD OF THE INVENTION

[0001] The present invention is in the general area of medical genetics and in the fields of oncology and immunology. More specifically, the invention is directed to the detection and quantification of total ras p21 in body fluids, particularly serial changes of total ras p21 levels in a subject's body fluids. Further, the invention is directed to detecting and quantitating total ras p21 in conjunction with one or more other proteins, such as, oncoproteins, angiogenic factors, tumor markers, inhibitors, growth factor receptors, metastasis proteins, and tumor suppressors.

BACKGROUND

[0002] The ras family of oncogenes (homologous to the rat sarcoma virus) belongs to the G protein family of cellular proteins engaged in signal transduction. The ras genes code for proteins that are located on the inner surface of the plasma membrane and have GTPase activity. Normally, when triggered by growth factor binding to plasma membrane receptors, ras proteins bind GTP, resulting in activation of the ras signal. The ras gene family has three primary members (H-ras, K-ras, and N-ras) that are among the most commonly activated oncogenes found in human cancer. Other ras gene family members include the Rho/Rac, Rab, Arf and Ran subfamilies.

[0003] The three main ras genes (H-ras, K-ras, and N-ras) encode 21 kD alpha/beta sheet proteins of 189 amino acids which are found in all eukaryotic cells on the inner surface of the plasma membrane. The isoforms (denoted H-Ras, N-Ras and K-Ras) encoded by the three ras genes are collectively known in the art as "p21" or "ras p21," and share a high degree of homology (>90%) as well as common downstream effectors and upstream guanine-nucleotide exchange factors (GEFS). As is true for the other members of the ras gene family, ras p21 is an intracellular GDP/GTP binding G protein important in cell signal transduction, converting signals from receptor tyrosine kinases in the cell membrane to the nucleus to regulate normal cell proliferation and differentiation [Pruitt and Der (2002)]. Ras p21 proteins activated by substitutions at amino acid positions 12, 13, or 61 can act as oncoproteins with an increased transforming capacity. Mutated forms of ras p21 are locked in a constitutively active state by reducing GTP hydrolysis, which results in uncontrolled signals for cell growth. When either mutated or overexpressed, p21 proteins are oncogenic. Ras p21 proteins are overexpressed in 50-60% of breast cancers and mutated in approximately 30% of all cancers, notably pancreatic and

[0004] Mutational activation of ras oncogenes has been identified in a variety of cancers as well as precursor lesions. Mutationally activated ras genes are uncommon in stomach, esophagus, ovary, prostate and breast tumors but are reported in 20-40% of lung carcinomas, 30% of acute myelogenous leukemia, 40-50% of colorectal tumors and thyroid tumors and >80% of pancreatic tumors. Overall, approximately 30% of all human neoplasms have been reported to contain a ras gene mutation. [Adjei (2001); Bos (1989).] H-ras mutations are common in bladder, kidney, and thyroid carcinomas; K-ms mutations occur in non-small-cell lung, colorectal and pancreatic carcinomas; and N-ras mutations are found in

melanoma, hepatocellular carcinoma, and hematologic malignancies. [Adjei (2001).] Transformation of cells by oncogenic ras mutants can increase the expression of metalloproteinases, such as gelatinase and stromelysin, which can enhance tumor metastasis. In addition, the expression of vascular endothelial growth factor (VEGF) is increased in K-rasand H-ras transformed epithelial cells through the Raf pathway.

[0005] There has been some controversy in the art about the prognostic significance of mutated versus overexpressed ras p21. For example, Watson et al. [Breast Cancer Res. Treat., 17(3): 161-169 (January-February 1991)] reported that elevated p21 levels detected by semiquantitative Western blot were an important prognostic indicator in early breast cancer patients. In 2001, however, Karelia et al. [Int. J. Biol. Markers, 16(2): 97-104 (2001)] investigated p21 ras oncoprotein expression in colorectal cancer using immunohistochemical studies, and concluded that genetic alteration of p21 ras, not ras p21 positivity, was important in determining biological aggressiveness of colorectal cancer.

[0006] Antibodies that are used frequently to detect ras in tumors are pan reactive antibodies, which bind to all three forms of ras p21 (H-, N- and K-ras), both mutated and wild-type. For example, Sun et al., [Eur. J. Cancer, 27(12): 1646-1649 (1991)] used the pan reactive Mab Ras 11 in immuno-histochemical studies to examine the significance of ras p21 in colorectal adenocarcinoma, and found ras p21 positivity to correlate with proliferative activity and to have independent prognostic value.

[0007] Although quantitative assays of ras p21 in human tissues are available, earlier described assays of ras p21 in human body fluids are at best semi-quantitative. As ras is normally expressed, it would be useful to have an indicator of an activated (upregulated and/or mutationally stimulated) ras oncoprotein pathway, such as a quantitative assay of elevated or serial changes in circulating ras p21 levels, wherein elevated levels or changes in ras p21 levels could be used to detect ras-driven oncogenesis or changes in the status of a ras-driven tumor.

[0008] Carney, W. P., U.S. Pat. No. 5,443,956 (1995) describes semi-quantitative immunoassays of total ras p21 in normal human plasma but not in corresponding sera. That Carney patent also describes the use of the pan reactive Mab. Ras 17 to detect total ras p21 in ELISA assays of normal human plasma.

[0009] By 2000, at least two investigators had reported the presence of ras protein in the sera of lung cancer patients. In 1991, Brandt-Rauf, P. W., [J. Occup. Med., 33(9): 951-955 (September 1991)] detected ras in the serum of five out of 11 lung cancer patients and two out of 21 patients with nonmalignant lung diseases using immunoblots, and suggested the use of ras as a marker for the early detection of respiratory tract cancer. Anderson et al. also analyzed ras levels in the plasma of human lung cancer patients using Western blots. [Anderson et al., Mutat. Res, 349: 121-126 (1996); Anderson et al., Mutat. Res, 403(1-2): 229-235 (1998); Anderson et al., Int. J. Hyg. Environ. Health. 204(1): 55-60 (October 2001).] In the Anderson et al. 1996 study, ras levels were increased in 45% patients vs. 6% of normal controls. In some of the Anderson et al. studies, the primary antibody Ras 10, which detects total ras protein (reactive with normal and mutant ras) was used. Ras levels were not quantified (in picograms) in either the Brandt-Rauf (1991) or the Anderson et al. studies.

[0010] Schneider et al. [Clin. Chem. Lab. Med., 38(4): 301-305 (2000)] reported that ras p21 could not be detected in lung cancer patient sera, or in sera from any of the study groups, and therefore ras could not be considered useful as a marker for the early detection of lung cancer. Schneider et al. (2000) assayed sera of 65 male lung cancer patients, 29 non-malignant lung disease, and 44 controls, using Ab-1 (pan reactive anti-ras p21 rat Mabs from clone Y13-259) which reacts with H, K and N-ras, in three different Western blot methods. However, he could not detect ras in those serum samples, although 1 ng ras protein was measurable in standards.

[0011] Rundle et al. [Cancer Lett, 185(1): 71-78 (Nov. 8, 2002)] examined ras p21 levels in plasma from breast cancer patients, using Western blots, computer aided image analysis (integrated pixel units), and a pan reactive anti-ras antibody. Rundle at al. (2002) found detectable levels of wild-type ras in 53% of the breast cancer patients; the average levels in breast cancer patients were qualitatively higher than the average wild-type ras levels found in positive control subjects (27% of benign breast disease patients and 26% of healthy controls). Perera et al. [Arch. Toxicol., 64(5): 401-406 (1990)] studied serial changes in serum levels of 9 different oncogene protein products, including ras, both prior to and throughout the course of chemotherapy. Again, the ras determinations of both the Rundle at al. (2002) and Perera et al. (1990) assays were only semi-quantitative.

[0012] It may be critical to assay for both mutated and overexpressed, ras p21 even in those cancers that are not associated with mutated ras, as overexpressed ras may also be indicative of an activated ras pathway. For example, although ras is overexpressed in 50-60% and mutated in less than 5% of breast cancers, there is considerable experimental evidence that mutationally activated ras can promote breast cancer growth and development, possibly due to activation of HER-2/neu or other oncogenes. [Eckert et al., *Cancer Research.* 64: 4585-4592 (2004).]

[0013] The inventors have discovered a quantitative immunoassay that can be used to measure serial changes in total ras p21 levels in body fluids, which changes are useful as an indicator of an activated (upregulated and/or mutationally stimulated) ras pathway. The novel immunoassay comprises the use of the pan reactive monoclonal antibodies (Mabs) Ras 17 and Ras 10 or Ras 10.2, each of which bind total ras p21 protein (both normal and mutation-activated ras p21). The Mabs. Ras 17 and Ras 10 have been described previously [Carney, W. P., U.S. Pat. Nos. 5,262,523, 5,443,956, 5,081, 230 and 6,200,764]; however, the use of the combination of Mabs Ras 17 and Ras 10 in an immunoassay has not been previously described.

[0014] The present invention discloses quantitative assays to detect changes in total ras p21 levels in body fluid samples to detect activated ras pathways, whether upregulated or mutationally stimulated. Such assays alone or in combination with levels of cancer markers upstream of ras (such as TIMP-1), or with levels of other proteins, may be used to select patient therapies targeted to the ras pathway. Because of the importance of ras in tumorigenesis, several approaches have been taken for developing ras-targeted anticancer therapies, such as inhibition of ras protein expression (e.g., antisense oligonucleotides), prevention of membrane localization essential for ras activation (e.g. using Ras farnesylation inhibitors) [Kohl et al. (1995); Lerner et al. (1995)], or inhibition of downstream effectors of ras (e.g., Raf serine/threonine kinases) thereby preventing the constitutive activation of

the MAPK signal cascade. [Pruitt and Der (2002).] One promising candidate for a ras pathway-targeted therapy is the bis-aryl urea sorafenib (BAY-43-9006) [Wilhelm et al., *Cancer Research*, 64: 7099-7109 (Oct. 1, 2004)], which is known to inhibit both raf kinase and VEGFR.

[0015] Until the present invention, there had not been a sensitive, quantitative and routine serum test to determine whether a cancer patient has a ras p21-driven tumor. The assays of this invention provide such sensitive and quantitative assays that can be used routinely, not only for serum, but for plasma, and other body fluids.

SUMMARY OF THE INVENTION

[0016] The invention concerns the use of quantitative immunoassays to measure serial changes in the levels of total ras p21 protein in human body fluids. Said serial changes are particularly useful as an indicator of the activation status of the ras signal transduction pathway, wherein "activation" is defined to include both upregulation and mutational activation. According to the theories underlying the methods of the invention, increasing total ras p21 protein (detected by serial changes in total ras p21 levels) is considered to be an indicator of an activated ras signal transduction pathway, signifying the presence of preneoplastic/neoplastic disease in a human subject. The immunoassays of the invention can be used clinically—diagnostically and/or prognostically, to detect precancer and/or cancer (preneoplastic/neoplastic disease) and as a therapeutic aid for patient therapy selection, monitoring the status of a preneoplastic/neoplastic disease in a patient, or monitoring how a patient with a preneoplastic/neoplastic disease is responding to a therapy. A preferred use for the immunoassays of the invention is to monitor the status of patients, and to make decisions about the optimal method for patient therapy.

[0017] Preferred samples in which to assay serial changes in total ras p21 by immmunoassay are samples of human body fluids, which can include, among other body fluids: blood, serum, plasma, urine, saliva, semen, breast exudate, cerebrospinal fluid, tears, sputum, mucous, lymph, cytosols, ascites, pleural effusions, amniotic fluid, bladder washes and bronchioalveolar lavages. Preferred body fluids to assay are blood, serum, and plasma, more preferably, serum or plasma. Particularly preferred body fluid samples include pretreatment samples and samples taken from a patient who has not responded to treatment.

[0018] In one aspect, the invention concerns diagnostic/prognostic methods to detect a preneoplastic/neoplastic disease associated with an activated ras pathway. One method of detecting a preneoplastic/neoplastic disease associated with an activated ras pathway in a human subject comprises the steps of:

- [0019] (a) immunologically detecting and quantifying the average level of total ras p21 protein in samples of a body fluid taken from individuals of a control population;
- [0020] (b) immunologically detecting and quantifying serial changes in total ras p21 protein levels in equivalent samples of body fluid taken from the subject over time; and
- [0021] (c) comparing the levels of total ras p21 protein in the subject's samples to the average level of total ras p21 protein in the control samples;

[0022] wherein a level of total ras p21 protein in the subject's samples that is above the average level of total ras p21

protein in the control samples is indicative of an activated ras pathway and the presence of preneoplastic/neoplastic disease in the subject. The individuals of a control population of step (a) may be of either gender, or may be of the same gender as the subject.

[0023] Said method of detecting a preneoplastic/neoplastic disease associated with an activated ras pathway in a human subject may be further prognostic for said preneoplastic/neoplastic disease, wherein the levels of total ras p21 protein in the subject's samples relative to the average level of total ras p21 in the control samples are indicative of a better or poorer prognosis for said subject. Preferably, said prognosis is a clinical outcome selected from the group consisting of response rate (RR), complete response clinical benefit (CR), partial response clinical benefit (PR), stable disease clinical benefit (SD), time to progression (TTP), and time to death (TTD).

[0024] Said methods can be in standard formats, as for example, by an immunoassay in the form of a sandwich immunoassay, preferably a sandwich enzyme-linked immunosorbent assay (ELISA) or an equivalent assay. A preferred format is a sandwich ELISA comprising the use of monoclonal antibodies, preferably anti-p21 pan reactive monoclonal antibodies, preferably a sandwich ELISA wherein the capture monoclonal antibody is designated Ras 17 which is secreted from hybridoma HB 10054 deposited at the American Type Culture Collection (ATCC), and wherein the detector monoclonal antibody is designated Ras 10 which is secreted from hybridoma HB 9426 deposited at the ATCC, or Ras 10.2, which is secreted from a subclone of the Ras 10 hybridoma HB 9426. More preferably, the detector monoclonal antibody Ras 10 or Ras 10.2 is biotinylated.

[0025] Said methods can be diagnostic and/or prognostic for cancerous or precancerous diseases, wherein said disease is associated with an activated ras pathway. Exemplary of such precancerous or cancerous diseases are those selected from the group consisting of colorectal cancer, colon cancer, lung cancer, non-small-cell lung cancer, small-cell lung cancer, acute myelogenous leukemia, thyroid cancer, pancreatic cancer, bladder cancer, kidney cancer, melanoma, breast cancer, prostate cancer, ovarian cancer, cervical cancer, head-and-neck cancer, hepatocellular carcinoma and hematologic malignancies, and precancers leading to such cancers. Colon, colorectal, lung cancer and breast cancer, and precancers leading thereto, are considered particular targets for the diagnostic/prognostic assay methods that detect serial changes in total ras p21 in human body fluids.

[0026] A second aspect of the invention concerns a quantitative immunoassay to measure serial changes in the levels of total ras p21 protein in human body fluids, as a method of therapy selection for a human patient with a preneoplastic/neoplastic disease. One such method of therapy selection comprises the steps of

[0027] (a) immunologically detecting and quantifying the average level of total ras p21 protein in samples of a body fluid taken from individuals of a control population;

[0028] (b) immunologically detecting and quantifying serial changes in total ras p21 protein levels in equivalent samples of body fluid taken from the patient over time;

[0029] (c) comparing the levels of total ras p21 protein in the patient's samples to the average level of total ras p21 protein in the control samples; and [0030] (d) deciding whether to use conventional cancer therapy and/or ras-directed cancer therapy to treat the patient based upon the differences between the levels of total ras p21 protein in the patient's samples and the average level of total ras p21 protein in the control samples, and in view of the serial changes among the levels of total ras p21 protein in the patient's samples. For example, if a level of total ras p21 protein in a patient's sample is found to be above the average level of total ras p21 protein in the control samples, and if the serial changes in the total ras p21 protein levels in the patient's samples reflect levels that are trending above the average level of total ras p21 protein in the control samples, the conclusion could be drawn that the patient has a ras oncogene driven preneoplastic/neoplastic disease, and the decision may be made to use ras-directed therapy to treat the patient, either alone or in conjunction with one or more other anticancer therapies.

[0031] Preferably, said ras-directed therapy is selected from the group consisting of farnesyltransferase inhibitors (FTI), tyrosine kinase inhibitors, bis-aryl ureas and antisense inhibitors of ras. More preferably, said ras-directed therapy is the bis aryl-urea sorafenib (BAY 43-9006).

[0032] Another aspect of the invention concerns the use of an immunoassay as a method of monitoring the status of a preneoplastic/neoplastic disease in a patient, and/or monitoring how a patient with a preneoplastic/neoplastic disease is responding to a therapy, comprising immunologically detecting and quantifying serial changes in total ras p21 protein levels in samples of a body fluid taken from said patient over time; wherein increasing levels of total ras p21 protein over time indicate disease progression or a negative response to said therapy, and wherein decreasing levels of total ras p21 protein indicate disease remission or a positive response to said therapy. Said therapy may be conventional anticancer therapy, unconventional anticancer therapy, and/or a ras-directed therapy. A preferred body fluid sample, among other types of samples, may be a pretreatment sample, or a sample from a cancer patient who has not responded to treatment.

[0033] Still another aspect of this invention concerns the use of quantitative immunoassays to detect changes in total ras p21 levels in combination with the levels of one or more other protein(s), in the methods of this invention as in the diagnostic/prognostic methods, in the methods of therapy selection for patients with a preneoplastic/neoplastic disease, in the methods of monitoring the status of a preneoplastic/ neoplastic disease in a patient, and of monitoring how a patient with a preneoplastic/neoplastic disease is responding to a ras-directed or other therapy. Preferably, said other proteins that would be useful to quantitate along with ras p21 in the assays of this invention are inhibitors, oncoproteins, growth factor receptors, angiogenic factors, metastasis proteins, tumor markers and tumor suppressors. Preferably, said inhibitor is tissue-inhibitor of metalloproteinase-1 (TIMP-1), said oncoprotein is HER-2/neu, said growth factor receptors are selected from the group consisting of epidermal growth factor receptor (EGFR) and platelet derived growth factor receptor (PDGFR), said angiogenic factor is vascular endothelial growth factor (VEGF), said metastasis protein is urokinase-type plasminogen activator (uPA), said tumor marker is carcinoembryonic antigen (CEA), and said tumor suppressor is p53.

[0034] Most preferably, said other proteins are proteins that activate the ras pathway, such as TIMP-1 and HER-2/neu/c-

erbB-2 ("HER-2/neu"). For example, increasing levels of total ras p21 and/or HER-2/neu may be indicative of a greater probability of early recurrence of a tumor or tumor metastasis. It would be advantageous to test cancer patients for serial changes in both total ras p21 and such other proteins, especially proteins that activate the ras pathway, to enlarge the clinical perspective, therapeutic resources and diagnostic/prognostic parameters to pick the optimal therapeutic combinations for the most promising treatment outcomes.

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ABBREVIATIONS

[0109] The following abbreviations are used herein:

[0110] ATCC—American Type Culture Collection

[0111] BSA—bovine serum albumin

[0112] ° C.—degrees centigrade

[0113] CEA—carcinoembryonic antigen

[0114] CV—coefficient of variation

[0115] EDTA—ethylenediaminetetraacetate

[0116] EGFR—epidermal growth factor receptor

[0117] ELISA—enzyme-linked immunosorbent assay

[0118] FTI—farnesyltransferase inhibitor

[0119] GDP—guanosine diphosphate

[0120] GEFS—guanine-nucleotide exchange factors

[0121] GTP—guanosine triphosphate

[0122] G—protein guanyl nucleotide dependent protein

[0123] HRP—horseradish peroxidase

[0124] kD—kilodalton

[0125] M—molar

[0126] Mab—monoclonal antibody

[0127] MAP—mitogen-activated protein

[0128] ml—milliliter

[0129] mM—millimolar

[0130] N—normal or number (depending on context)

[0131] ND—not detected

[0132] NED—no evidence of disease

[0133] Ng—nanogram

[0134] nm—nanometer

[0135] OD—optical density

[0136] pg—proteoglycan

[0137] PDGFR—platelet derived growth factor receptor

[0138] P. Resp—patient's response

[0139] Prog—progressive disease

[0140] Recur—recurrence

[0141] SD—standard deviation

[0142] Stab—stable

[0143] Stg—stage

[0144] TIMP-1—tissue inhibitor of metalloproteinase-1

[0145] TMB—tetramethylbenzidine

[0146] µg—microgram

[0147] µl—microliter

[0148] Unk—unknown

[0149] uPA—urokinase-type plasminogen activator

[0150] VEGF—vascular endothelial growth factor

[0151] VEGFR—vascular endothelial growth factor receptor

BRIEF DESCRIPTION OF THE FIGURES

[0152] FIG. 1 is an illustrative standard curve constructed by reacting a wild-type recombinant ras p21 standard at 6 different concentrations of 0, 40, 150, 300, 500 and 700 pg/ml in a preferred embodiment of the assays of this invention, a ras p21 ELISA, as described in detail in the Materials and Methods, and in Example 1, below.

DETAILED DESCRIPTION

[0153] The present invention is directed to quantitative immunoassays that measure serial changes in the levels of total ras p21 protein in human body fluids, which assays are useful diagnostically/prognostically to detect or monitor a preneoplastic/neoplastic disease in a human, or to select a therapy for a patient with said preneoplastic/neoplastic disease, wherein said disease is associated with an activated ras pathway. As used herein, an "activated ras pathway" is defined as a ras pathway activated by either overexpression or mutation of ras p21 protein, and therefore encompasses upregulated and/or mutationally stimulated ras pathways. Further, as used herein, "total ras p21 protein" comprises both mutation-activated and normal (wild-type) p21 protein (including H-ras, K-ras, and N-ras proteins).

[0154] Exemplary of preneoplastic/neoplastic diseases associated with an activated ras pathway are the following, as well as precancers leading to the following: colorectal cancer, colon cancer, lung cancer, non-small-cell lung cancer, small-cell lung cancer, acute myelogenous leukemia, thyroid cancer, pancreatic cancer, bladder cancer, kidney cancer, melanoma, breast cancer, prostate cancer, ovarian cancer, cervical cancer, head-and-neck cancer, hepatocellular carcinoma and hematologic malignancies. In particular, the levels of total ras p21 protein, alone or in combination with levels of other proteins, particularly other oncoproteins, can be used to predict clinical outcome and/or as an aid in therapy selection.

[0155] The assays of this invention can be diagnostic and/or prognostic, i.e., diagnostic/prognostic. The term "diagnostic/ prognostic" is herein defined to encompass the following processes either individually or cumulatively depending upon the clinical context: determining the presence of disease, determining the nature of a disease, distinguishing one disease from another, forecasting as to the probable outcome of a disease state, determining the prospect as to recovery from a disease as indicated by the nature and symptoms of a case, monitoring the disease status of a patient, monitoring a patient for recurrence of disease, and/or determining the preferred therapeutic regimen for a patient. The diagnostic/prognostic methods of this invention are useful, for example, for screening populations for the presence of neoplastic or preneoplastic disease, determining the risk of developing neoplastic disease, diagnosing the presence of neoplastic and/or preneoplastic disease, monitoring the disease status of patients with neoplastic disease, and/or determining the prognosis for the course of neoplastic disease.

[0156] The present invention is useful for screening for the presence of a wide variety of preneoplastic/neoplastic diseases as indicated above. Such an assay can be used to detect tumors, monitor their growth, and help in the diagnosis and prognosis of disease. The assays can also be used to detect the

presence of cancer metastasis, as well as confirm the absence of tumor tissue following cancer chemotherapy and/or radiation therapy. It can further be used to monitor cancer chemotherapy and tumor reappearance.

[0157] The methods include quantifying total ras p21 protein, if any, present in serial body fluid samples taken from a human subject diagnosed with, or suspected of having, a preneoplastic/neoplastic disease. The quantified total ras p21 protein levels are compared with the average levels in body fluid samples taken from individuals of a control population, wherein an above average level of total ras p21 protein is indicative of an activated ras pathway. As used herein, an "above average" level of total ras p21 protein indicates a level higher than two standard deviations (SD) above the mean level found in control samples. The individuals of the control population can be of either gender, or can be restricted to those who are of the same gender as the subject.

[0158] As used herein, "cancerous" and "neoplastic" have equivalent meanings, and "precancerous" and "preneoplastic" have equivalent meanings. The use of detection of gene expression products of oncogenes as diagnostic/prognostic indicators for preneoplastic/neoplastic diseases is considered conventional by those of skill in the art. However, the application of such approaches to measure quantitatively serial changes in ras levels in body fluids, particularly of circulating ras levels, to detect or monitor a preneoplastic/neoplastic disease, wherein said disease is associated with an activated ras pathway, is new.

Serial Samples of Body Fluids

[0159] In a preferred embodiment of the invention, total ras p21 protein is quantitated in human body fluid samples drawn serially over time. Such body fluid samples can be blood, serum, plasma, urine, saliva, semen, breast exudate, cerebrospinal fluid, tears, sputum, mucous, lymph, cytosols, ascites, pleural effusions, amniotic fluid, bladder washes and bronchioalveolar lavages, among other body fluid samples. Preferred body fluids, for example, include serum, or plasma samples treated with heparin, citrate or EDTA, among other body fluid samples, and can be fresh or frozen. Such body fluid specimens can be taken pretreatment, during treatment, or post-treatment, or can be taken from a patient who is not responding to therapy. As used herein, "serial changes over time" or "serial samples" denotes sequential testing of samples taken over time periods which would be considered relevant for the subject, depending on the context and the circumstances. For example, for cancer patient screening, serial samples might be drawn upon a patient's initial visit, after diagnosis, pre-surgery and/or post-surgery; whereas for population screening for a preneoplastic disease, serial samples might be drawn on a yearly basis.

Immunoassays

[0160] An exemplary and preferred immunoassay according to the methods of the invention is a sandwich ELISA described below in the Materials and Methods section, and was used to obtain the data for Examples 1 through 8. It can be appreciated that alternate methods, in addition to those disclosed herein, can be used to quantify total ras p21 protein in human body fluids. Other preferred sandwich assays could be used with other visualizing means, such as luminescent labels. Other labels are detailed below under the subsection labeled Labels.

[0161] Many formats can be adapted for use with the methods of the present invention. The detection and quantitation of total ras p21 protein in human body fluids can be performed, for example, by enzyme-linked immunosorbent assays, radioimmunoassays, dual antibody sandwich assays, agglutination assays, fluorescent immunoassays, immunoelectron and scanning microscopy using immunogold, among other assays commonly known in the art. The quantitation of total ras p21 protein in such assays can be adapted by conventional methods known in the art. In preferred embodiments, serial changes in circulating total ras p21 protein levels, or such levels in other body fluids, are detected and quantified by a sandwich assay in which the capture antibody has been immobilized, using conventional techniques, on the surface of the support.

[0162] Suitable supports used in assays include among other supports, synthetic polymer supports, such as polypropylene, polystyrene, substituted polystyrene, polyacrylamides (such as polyamides and polyvinylchloride), glass beads, agarose, and nitrocellulose, among other supports.

Exemplary Immunoassay

[0163] An exemplary and preferred ELISA sandwich immunoassay is described in the Materials and Methods section and in Examples 1-8. That exemplary ELISA uses Mab Ras 17 as the capture antibody and biotinylated Mab 10.2 (produced by a subclone of hybridoma Ras 10) as the detector antibody. The capture Mab Ras 17 is immobilized on microtiter plate wells; diluted human serum/plasma samples or ras p21 standards (recombinant wild-type H-ras protein) are incubated for three hours at 37° C. in the wells to allow binding of total ras p21 antigen by Mab 17. After washing of wells, the immobilized ras p21 antigen is exposed to the biotinylated detector antibody Mab Ras 10.2 for 30 minutes at room temperature (20-27° C.), after which the wells are again washed. A streptavidin-horseradish peroxidase conjugate is then added (30 minutes at room temperature). After a final wash, TMB Blue Substrate is added to the wells (and incubated for 30 minutes at room temperature) to detect bound peroxidase activity. The reaction is stopped by the addition of 2.5N sulfuric acid, and the absorbance is measured at 450 nm. Correlating the absorbance values of samples with the ras standards allows the determination of a quantitative value of total ras p21 in pg/ml of serum or plasma.

Combination Assays: Levels of Ras and Other Protein(s)

[0164] It can be appreciated that other proteins, such as inhibitors, oncoproteins, growth factor receptors, angiogenic factors, metastasis proteins, tumor markers, and tumor suppressors, particularly those associated with the ras pathway, may be suitable for detection and quantitation along with ras p21, depending on the subject preneoplastic/neoplastic disease. Exemplary and preferred other proteins suitable for testing along with ras p21 include tissue inhibitor of metalloproteinase-1 (TIMP-1), HER-2/neu, epidermal growth factor receptor (EGFR), platelet derived growth factor receptor (PDGFR), vascular endothelial growth factor (VEGF), urokinase-type plasminogen activator (uPA), carcinoembryonic antigen (CEA), and p53. Assays to detect preferred other proteins suitable for the methods of the invention are known to one of skill in the art. For example, immunoassays for the quantitation of HER-2/neu and TIMP-1 in human body fluids are commercially available, such as the Oncogene Science TIMP-1 ELISA [Oncogene Science, Cambridge, Mass. (USA), www.oncogene.com], which can determine ng/ml values of TIMP-1 levels in human serum or plasma. As described in Example 8, serial colorectal cancer serum samples and normal serum samples were assayed for both total ras p21 and TIMP-1 levels.

Prognosis

[0165] Monitoring the levels of ras p21 and/or other proteins can be indicative of clinical outcomes for preneoplastic/neoplastic diseases. A preferred method of evaluating a clinical outcome is one based on response rate (RR), clinical benefit [including complete response (CR), partial response (PR), and stable disease (SD)], time to progression (TTP), and time to death (TTD). Other methods of evaluating prognosis are known in the art and can be used.

Antibodies

[0166] The term "antibody" herein is used in the broadest sense and specifically covers monoclonal antibodies (including full length monoclonal antibodies), polyclonal antibodies, multispecific antibodies (e.g., bispecific antibodies), and antibody fragments so long as they exhibit the desired biological activity. Antibodies useful according to the methods of the invention may be prepared by conventional methodology and/or by genetic engineering. Preferred antibodies according to the invention are those that are "pan reactive," that is, antibodies that bind to total ras p21 (both wild-type and mutation-activated ras p21), as well as the three families of the Harvey, Kirsten and N-ras p21. Most preferred are the antibodies used in the exemplary immunoassay of the invention, the Ras 17 and Ras 10 antibodies, or antibodies produced by subclones of the Ras 17 and Ras 10 hybridomas (such as the Ras 10.2 Mab from a subclone of the parent Ras 10 hybridoma), so long as they exhibit the desired biological activity.

[0167] "Antibody fragments" comprise a portion of a full length antibody, generally the antigen binding or variable domain thereof. Examples of antibody fragments include Fab, Fab', F(ab')₂, and Fv fragments, diabodies, linear antibodies, and single-chain antibody molecules, as well as multispecific antibodies, including bispecific antibodies, formed from antibody fragments.

[0168] The term "monoclonal antibody" as used herein refers to an antibody obtained from a population of substantially homogeneous antibodies, i.e., the individual antibodies comprising the population are identical except for possible naturally occurring mutations that may be present in minor amounts. Monoclonal antibodies are highly specific, being directed against a single antigenic site. Furthermore, in contrast to conventional (polyclonal) antibody preparations which typically include different antibodies directed against different determinants (epitopes), each monoclonal antibody is directed against a single determinant on the antigen. The modifier "monoclonal" indicates the character of the antibody as being obtained from a substantially homogeneous population of antibodies, and is not to be construed as requiring production of the antibody by any particular method. For example, the monoclonal antibodies to be used in accordance with the present invention may be made by the hybridoma method first described by Kohler et al., Nature 256:495 (1975), or may be made by recombinant DNA methods (see, e.g., U.S. Pat. No. 4,816,567). The "monoclonal antibodies"

may also be isolated from phage antibody libraries using the techniques described in Clackson et al., *Nature*, 352: 624-628 (1991) and Marks et al., *J. Mol. Biol.*, 222: 581-597 (1991), for example.

[0169] The monoclonal antibodies herein specifically include "chimeric" antibodies (immunoglobulins) in which a portion of the heavy and/or light chain is identical with or homologous to corresponding sequences in antibodies derived from a particular species or belonging to a particular antibody class or subclass, while the remainder of the chain(s) is identical with or homologous to corresponding sequences in antibodies derived from another species or belonging to another antibody class or subclass, as well as fragments of such antibodies, so long as they exhibit the desired biological activity [U.S. Pat. No. 4,816,567; and Morrison et al., *Proc. Natl. Acad. Sci.*, USA, 81: 6851-6855 (1984)].

[0170] "Humanized" forms of non-human (e.g., murine) antibodies are chimeric antibodies which contain minimal sequence derived from non-human immunoglobulin. For the most part, humanized antibodies are human immunoglobulins (recipient antibody) in which hypervariable region residues of the recipient are replaced by hypervariable region residues from a non-human species (donor antibody) such as mouse, rat, rabbit or nonhuman primate having the desired specificity, affinity, and capacity. In some instances, framework region (FR) residues of the human immunoglobulin are replaced by corresponding non-human residues. Furthermore, humanized antibodies may comprise residues which are not found in the recipient antibody or in the donor antibody. Such modifications are made to further refine antibody performance. In general, the humanized antibody will comprise substantially all of at least one, and typically two, variable domains, in which all or substantially all of the hypervariable regions correspond to those of a non-human immunoglobulin and all or substantially all of the FRs are those of a human immunoglobulin sequence. The humanized antibody optionally also will comprise at least a portion of an immunoglobulin constant region (Fc), typically that of a human immunoglobulin. For further details, see Jones et al., Nature, 321:522-525 (1986); Reichmann et al., Nature 332: 323-329 (1988); and Presta, Curr. Op. Struct. Biol., 2: 593-596 (1992).

[0171] "Single-chain Fv" or "sFv" antibody fragments comprise the V_H and V_L domains of antibody, wherein these domains are present in a single polypeptide chain. Generally, the Fv polypeptide further comprises a polypeptide linker between the V_H and V_L domains which enables the sFv to form the desired structure for antigen binding. For a review of sFv see Pluckthun in *The Pharmacology of Monoclonal Antibodies*, Vol. 113, Rosenburg and Moore eds. Springer-Verlag, New York, pp. 269-315 (1994).

[0172] The term "diabodies" refers to small antibody fragments with two antigen-binding sites, which fragments comprise a heavy chain variable domain (V_H) connected to a light chain variable domain (V_L) in the same polypeptide chain $(V_{H^*}V_L)$. By using a linker that is too short to allow pairing between the two domains on the same chain, the domains are forced to pair with the complementary domains of another chain and create two antigen-binding sites. Diabodies are described more fully in, for example, EP 404,097; WO 93/11161; and Hollinger et al., *Proc. Natl. Acad. Sci.*, USA, 90: 6444-6448 (1993).

[0173] The expression "linear antibodies" refers to the antibodies described in Zapata et al., *Protein Eng.*, 8(10): 1057-

1062 (1995). Briefly, such antibodies comprise a pair of tandem Fd segments (V_H - C_H 1- V_H - C_H 1) which form a pair of antigen binding regions. Linear antibodies can be bispecific or monospecific.

[0174] Representative monoclonal antibodies useful according to this invention include Mabs Ras 17 and Ras 10 described in earlier Carney et al. patents [U.S. Pat. No. 5,081, 230; U.S. Pat. No. 5,443,956; U.S. Pat. No. 6,200,764]. Monoclonal antibodies useful according to this invention serve to identify total ras p21 proteins in various laboratory prognostic tests, for example, in clinical samples.

[0175] General texts describing additional molecular biological techniques useful herein, including the preparation of antibodies include Berger and Kimmel, Guide to Molecular Cloning Techniques, Methods in Enzymology, Vol. 152, Academic Press, Inc., Sambrook et al., Molecular Cloning: A Laboratory Manual, (Second Edition, Cold Spring Harbor Laboratory Press; Cold Spring Harbor, N.Y.; 1989) Vol. 1-3; Current Protocols in Molecular Biology, F. M. Ausabel et al. [Eds.], Current Protocols, a joint venture between Green Publishing Associates, Inc. and John Wiley & Sons, Inc. (supplemented through 2000), Harlow et al., Monoclonal Antibodies: A Laboratory Manual, Cold Spring Harbor Laboratory Press (1988), Paul [Ed.]; Fundamental Immunology, Lippincott Williams & Wilkins (1998), and Harlow et al., Using Antibodies: A Laboratory Manual, Cold Spring Harbor Laboratory Press (1998).

Labels

[0176] The antibodies useful according to this invention to identify total ras p21 proteins can be labeled in any conventional manner. A preferred label, according to this invention is horseradish peroxidase, and a preferred method of labeling the antibodies is by using biotin-strepavidin complexes.

[0177] As appropriate, antibodies used in the immunoassays of this invention that are used as tracers may be labeled in any manner, directly or indirectly, that results in a signal that is visible or can be rendered visible. Detectable marker substances include radionuclides, such as $^3\mathrm{H},\,^{125}\mathrm{I},$ and $^{131}\mathrm{I};$ fluorescers, such as, fluorescein isothiocyanate and other fluorochromes, phycobiliproteins, phycoerythin, rare earth chelates, Texas red, dansyl and rhodamine; colorimetric reagents (chromogens); electron-opaque materials, such as colloidal gold; bioluminescers; chemiluminescers; dyes; enzymes, such as, horseradish peroxidase, alkaline phosphatases, glucose oxidase, glucose-6-phosphate dehydrogenase, acetylcholinesterase, alpha-, beta-galactosidase, among others; coenzymes; enzyme substrates; enzyme cofactors; enzyme inhibitors; enzyme subunits; metal ions; free radicals; or any other immunologically active or inert substance which provides a means of detecting or measuring the presence or amount of immunocomplex formed. Exemplary of enzyme substrate combinations are horseradish peroxidase and tetramethyl benzidine (TMB), and alkaline phosphatases and paranitrophenyl phosphate (pNPP).

[0178] Another preferred detection, or detection and quantitation systems according to this invention produce luminescent signals, bioluminescent (BL) or chemiluminescent (CL). In chemiluminescent (CL) or bioluminescent (BL) assays, the intensity or the total light emission is measured and related to the concentration of the unknown analyte. Light can be measured quantitatively using a luminometer (photomultiplier tube as the detector) or charge-coupled device, or qualitatively by means of photographic or X-ray film. The

main advantages of using such assays are their simplicity and analytical sensitivity, enabling the detection and/or quantitation of very small amounts of analyte.

[0179] Exemplary luminescent labels are acridinium esters, acridinium sulfonyl carboxamides, luminol, umbelliferone, isoluminol derivatives, photoproteins, such as aequorin, and luciferases from fireflies, marine bacteria, Vargulla and Renilla. Luminol can be used optionally with an enhancer molecule, preferably selected from the group consisting of 4-iodophenol or 4-hydroxy-cinnamic acid. Acridinium esters are one of the preferred types of CL labels according to this invention. Typically, a CL signal is generated by treatment with an oxidant under basic conditions.

[0180] Also preferred luminescent detection systems are those wherein the signal (detectable marker) is produced by an enzymatic reaction upon a substrate. CL and BL detection schemes have been developed for assaying alkaline phosphatases (AP), glucose oxidase, glucose 6-phosphate dehydrogenase, horseradish peroxidase (HRP), and xanthine-oxidase labels, among others. AP and HRP are two preferred enzyme labels which can be quantitated by a range of CL and BL reactions. For example, AP can be used with a substrate, such as an adamantyl 1,2-dioxetane aryl phosphate substrate (e.g. AMPPD or CSPD; [Kricka, L. J., "Chemiluminescence and Bioluminescence, Analysis by," at p. 167, Molecular Biology and Biotechnology: A Comprehensive Desk Reference (ed. R. A. Meyers) (VCH Publishers; N.Y., N.Y.; 1995)); preferably a disodium salt of 4-methoxy-4-(3-phosphatephenyl)spiro[1,2-dioxetane-3,2'-adamantane], with or without an enhancer molecule, preferably, 1-(trioctylphosphonium methyl)-4-(tributylphosphonium methyl) benzene diochloride. HRP is preferably used with substrates, such as, 2',3',6'trifluorophenyl 3-methoxy-10-methylacridan-9-carboxylate. [0181] CL and BL reactions can be adapted for analysis not

only of enzymes, but also of other substrates, cofactors, inhibitors, metal ions and the like. For example, luminol, firefly luciferase, and marine bacterial luciferase reactions are indicator reactions for the production or consumption of peroxide, ATP, and NADPH, respectively. They can be coupled to other reactions involving oxidases, kinases, and dehydrogenases, and can be used to measure any component of the coupled reaction (enzyme, substrate, cofactor).

[0182] The detectable marker may be directly or indirectly linked to an antibody used in an assay of this invention. Exemplary of an indirect linkage of the detectable label is the use of a binding pair between an antibody and a marker or the use of a signal amplification system.

[0183] Exemplary of binding pairs that can be used to link antibodies of assays of this invention to detectable markers are biotin/avidin, streptavidin, or anti-biotin; avidin/anti-avidin; thyroxine/thyroxine-binding globulin; antigen/antibody; antibody/anti-antibody; carbohydrate/lectins; hapten/antihapten antibody; dyes and hydrophobic molecules/hydrophobic protein binding sites; enzyme inhibitor, coenzyme or cofactor/enzyme; polynucleic acid/homologous polynucleic acid sequence; fluorescein/anti-fluorescein; dinitrophenol/anti-dinitrophenol; vitamin B12/intrinsic factor; cortisone, cortisol/cortisol binding protein; and ligands for specific receptor proteins. Preferred binding pairs according to this invention are biofin/avidin or streptavidin, more preferably biotin/streptavidin.

[0184] Various means for linking labels directly or indirectly to antibodies are known in the art. For example, labels

may be bound either covalently or non-covalently. Exemplary antibody conjugation methods are described in: Avarmeas et al., *Scan. J. Immunol.*, 8 (Suppl. 7): 7 (1978); Bayer et al., *Meth. Enzymol.*, 62: 308 (1979); Chandler et al., *J. Immunol. Meth.*, 53: 187 (1982); Ekeke and Abuknesha, *J. Steroid Biochem.*, 11: 1579 (1979); Engvall and Perlmann, *J. Immunol.*, 109: 129 (1972); Geoghegan et al., *Immunol. Comm.*, 7: 1 (1978); and Wilson and Nakane, *Immunofluorescence and Related Techniques*, p. 215 [Elsevier/North Holland Biomedical Press; Amsterdam (1978)].

[0185] Depending upon the nature of the label, various techniques can be employed for detecting and quantitating the label. For fluorescers, a large number of fluorometers are available. For chemiluminescers, luminometers or films are available. With enzymes, a fluorescent, chemiluminescent, or colored product can be determined or measured fluorometrically, luminometrically, spectrophotometrically or visually.

[0186] Various types of chemiluminescent compounds having an acridinium, benzacridinium, or acridan type of heterocyclic ring systems are preferred labels. Acridinium and benzacridinium esters are currently the more preferred chemiluminescent compounds, with preferred acridinium esters including those compounds having heterocyclic rings or ring systems that contain the heteroatom in a positive oxidation state including such ring systems as acridinium, benz[a]acridinium, benz[b]acridinium, benz[c]acridinium, a benzimidazole cation, quinolinium, isoquinolinium, quinolizinium, a cyclic substituted quinolinium, phenanthridinium, and quinoxalinium, as are well-known in the art.

[0187] The tracer may be prepared by attaching to the selected antibody either directly or indirectly a reactive functional group present on the acridinium or benzacridinium ester, as is well known to those skilled in the art, e.g. Weeks et al., *Clinical Chemistry*, 29(8), 1474-1479, (1983). Particularly preferred compounds are acridinium and benzacridinium esters with an aryl ring leaving group and the reactive functional group present in either the para or the meta position of the aryl ring. [See, U.S. Pat. No. 4,745,181 and WO 94/21823.]

Ras-Directed Therapies

[0188] As used herein, "ras-directed therapies" include any therapies that are targeted to the ras pathway, including inhibition of ras protein expression (e.g., antisense oligonucleotides), prevention of membrane localization essential for ras activation (ras farnesylation inhibitors), or inhibition of downstream effectors of ras (e.g., Raf serine/threonine kinases). Preferred ras-directed therapies include farnesyltransferase inhibitors (FTIs), tyrosine kinase inhibitors, or bis-aryl ureas. Exemplary FTIs are BMS-214662 [an imidazole-containing nonpeptidomimetic FTI; Cortes et al., J. Clin. Oncol, 23 (12): 2805-2812 (Apr. 20, 2005); Bristol-Myers Squibb, Princeton, N.J. (USA)], FTI-2153 and FTI-277 [EMD Biosciences, www.emdbiosciences.com (Cat. No. 3444555); Lerner et al. (1995); Sun et al., Cancer Res., 59: 4919-4926 (1999)], R115777 [methylquinolone; Janssen Research Foundation, Titusville, N.J. (USA)], and Sch66336 [Sarasar® or Lonafarniber; a benzocycloheptapyridyl FTI; Taveras et al., Curr. Topics Med. Chem., 3(10): 1103-1114 (June 2003); Schering Plough, Kenilworth, N.J. (USA)].

[0189] An exemplary tyrosine kinase inhibitor is AZD3049 [prenyl transferase inhibitor; AstraZeneca, www.astrazeneca. com]. An exemplary antisense inhibitor of ras is ISIS 2503 [ISIS Pharmaceuticals, Inc., Carlsbad, Calif. (USA)]. [See

also, for example, http://www.webtie.org/sots/html/LungAgents.htm, for other ras-directed therapies targeted to ras or at relevant effectors downstream from ras.]

[0190] A preferred ras-directed therapy according to the invention is the bis-aryl urea sorafenib (BAY 43-9006) [Onyx Pharmaceuticals, Richmond, Calif. (USA), and Bayer Corporation, West Haven, Conn. (USA); Lyons et al., *Endocrine-Related Cancer*, 8:219-225 (2001) and Wilhelm et al. (2004)], a small molecule which inhibits the enzyme Raf kinase. Sorafenib targets both the RAF/MEK/ERK [MEK: MAP/ERK kinase; ERK: extracellular signal related kinase] signaling pathway to inhibit cell proliferation, and the VEGFR-2/PDGFR-β signaling pathway to inhibit tumor angiogenesis. Sorafenib then would be a particularly preferred anticancer therapy for a patient found to have an activated ras pathway and an activated VEGFR-2/PDGFR-β signaling pathway.

EXAMPLES

[0191] The following examples are for purposes of illustration only and are not meant to limit the invention in any way.

Materials and Methods

[0192] Solid Phase Sandwich Microtiter ELISA for Human Serum and Plasma

[0193] Sample Preparation

[0194] Suitable samples for analysis by the ras p21 ELISA include human plasma treated with heparin, citrate, or EDTA, and human serum. Due to possible interfering factors, special care must be taken in the preparation and assay of human serum and plasma. Any flocculant material should be removed from samples by microcentrifugation prior to dilution. The initial concentration of the serum or plasma specimen to be examined should not exceed a concentration of 50% (a 1:2 dilution of specimen in sample diluent). For example, 0.150 ml of sample may be diluted into 0.150 ml of sample diluent, and 100 μ l added to the microplate wells.

[0195] Exemplary Assay Procedure

[0196] Purified Ras 17 Mabs [secreted from the Ras 17 hybridoma deposited at the ATCC under HB 10054] are diluted to 1 $\mu\text{g/ml}$ in coating buffer (0.03 M sodium carbonate anhydrous and 0.07 M sodium bicarbonate, pH 9.6) and are adsorbed to wells (at 100 µl per well) of 96-well microplates [Nunc MaxiSorpTM F8 Immuno Module, Part #1001; Nalge Nunc International Corp., Naperville, Ill. (USA)]. Adsorption is carried out overnight (at least 12 hours) at room temperature (20°-27° C.). At the end of the incubation period, blocking solution is added to the wells (0.002M sodium phosphate monobasic, 0.008M sodium phosphate dibasic heptahydrate, 0.15M sodium chloride, 0.29M beta D-lactose and 2% BSA reagent grade; pH 7.45) and aspirated using an Oyster Bay Microplate Coater (Model MPCS-3) [Oyster Bay Pump Works, Hicksville, N.Y. (USA)]. Coated microplates are dried using a LyoStar® Freeze-Dryer [FTS Systems, Stone Ridge, N.Y. (USA)] at 22° C. and stored at 2-8° C.

[0197] Human serum samples and ras p21 standards [Invitrogen Corporation, Catalog #P2138; Carlsbad, Calif. (USA)] are diluted serially with sample diluent [4%. BSA Protease Free (Serologicals Product #82-045; www.serologicals.com), 10 μ g/ml Purified Mouse IgG (Scantibody Laboratories Product #3BM222; Santel, Calif., USA) and 0.09% sodium azide]. Plasma samples treated with heparin, citrate or EDTA may be used instead of serum. Diluted samples are then added to microplate wells (100 μ l per well) and the

sealed plates incubated for 3 hours at 37° C. Each well is washed six times with 300 μL per well of platewash, pH 7.4 [0.137M sodium chloride, 2.7 mM potassium chloride, 1.45 mM potassium phosphate monobasic, 0.08 M sodium phosphate dibasic heptahydrate, and 0.05% Tween 20]. To each well is then added 100 μl of biotinylated Ras 10.2 Mabs [whose parent hybridoma Ras 10 is on deposit at the ATCC under HB 9426] diluted in detector diluent [0.4% casein, 1.7 mM sodium phosphate monobasic, 8.1 mM sodium phosphate dibasic heptahydrate, 0.15M sodium chloride, and 0.1% sodium azide, pH 7.15±0.1] (final Mab concentration 0.5-3.5 $\mu g/ml$; optimal concentration determined by titration and functional testing of detector Mab).

[0198] The plates are incubated for 30 minutes at room temperature. Each well is washed six times as above, and 100 μl of Strepavidin-HRP [Zymed Laboratories, Inc., Catalog #43-8323; www.zymed.com] diluted to 1 µg/mL or less in conjugate diluent [1.7 mM sodium phosphate monobasic, 8.1 mM sodium phosphate dibasic heptahydrate, 0.145M sodium chloride, 0.1% 2-chloroacetamide, 1% BSA Protease Free, and 0.05% Tween 20; pH 7.3] is added to each well. The plates are incubated at room temperature for 30 minutes. Each well is again washed six times as above, and 100 µl of a color producing substrate solution [TMB Blue Substrate, DakoCytomation, Catalog # S1601; www.dakocytomation.com] is added to each well to detect bound peroxidase activity. The plates are incubated 30 minutes at room temperature, and the reaction is stopped by the addition of 100 µl 2.5N sulfuric acid. The absorbance is measured at 450 nm using a Molecular Devices spectrophotometric plate reader (Model #Vmax; www.moleculardevices.com) and analyzed using a Softmax Pro™ (Molecular Devices Corporation; www.moleculardevices.com) software program.

Human Serum and Plasma Samples

[0199] Frozen plasma and serum samples from normal humans and cancer patients were obtained from Bioclinical Partners, Inc. [Franklin, Mass. (USA); now Zeptometrix (www.zeptometrix.com)].

Example 1

Ras Standard Curve

[0200] Quantitative analyses were made by constructing a standard curve using Ha ras p21 [recombinant wild-type ras p21 from Invitrogen Corporation, Catalog # P2138]. FIG. 1 presents the standard curve which was constructed by reacting various concentrations of the Ha-ras p21 protein with immobilized anti-p21 pan reactive monoclonal antibody Ras 17 and detecting captured ras p21 protein using biotinylated anti-total p21 monoclonal antibody Ras 10.2. The immunoassay procedure was as described above in Materials and Methods.

Example 2

Normal Human Serum

[0201] Microplate wells were coated with anti-total ras p21 capture antibody (Mab Ras 17), as described above. A total of 82 normal human sera (42 female, 40 male; obtained from Bioclinical Partners, Inc.) were individually tested in the coated microplates by ELISA, and ras p21 present in the sera was detected using biotinylated Ras 10.2 monoclonal anti-

body. The absorbance index at 450 nm was used to determine the level of total ras p21 in serum samples. The results show that, in general, higher levels of total ras p21 protein were found in normal females than in normal males (average of 122.4 pg/ml in females vs. 33.1 pg/ml in males).

TABLE 1

Sin	Single Point Serum samples from Normal Humans					
	rmal Female rum Samples		al Male Samples			
Sample	Ras p21 (pg/mL)	Sample	Ras p21 (pg/mL)			
3666	204.4	3552	54.3	-		
3668	148.4	3553	54.5			
3669	119.7	3554	73.9			
3670	137.5	3555	40.6			
3671	235.6	3556	36.2			
3672	134.8	3557	46.3			
3673	223.0	3558	41.9			
3674	146.8	3559	37.0			
3675	91.6	3560	2.2			
3676	75.2	3561	7.0			
3677	100.7	3562	5.0			
3678	337.9	3563	5.6			
3679	99.7	3564	0.8			
3680	303.8	3565	1.5			
3681	136.5	3566	5.4			
3682	76.1	3567	2.0			
3683	64.6	3568	4.6			
3684	80.6	3569	6.4			
3685	168.1	3570	4.0			
3686	85.9	3571	10.3			
3687	120.2	3572	1.1			
3688	138.9	3573	98.8			
3689	127.0	3574	3.7			
3690	57.6	3575	4.8			
3691	68.0	3576	35.9			
3692	49.4	3577	37.6			
3693	183.1	3578	5.4			
3694	50.1	3579	10.8			
3695	62.5	3580	5.0			
3696	49.2	3581	5.1			
3697	48.6	3582	4.7			
3698	59.1	3583	3.5			
3699	68.8	3584	4.4			
3700	63.6	3585	8.9			
3701	395.1	3586	10.6			
3702	49.7	3587	16.5			
3702	47.6	3588	6.6			
3704	60.8	3589	19.3			
3704	56.4	3590	453.1			
3706	62.8	3591	150.5			
3707	228.0	3391	150.5			
2.07	220.0			_		

]	RAS Leve		Summary I Human S	erum (pg/ml))
Gender	N	Range	Mean	Mean + 2SD	95% fall below:
Male	40	ND- 453.1	33.1	182.5	101.4
Female	41	ND- 395.1	122.4	289.7	303.8

Example 3

Plasma vs. Serum

[0202] Sixteen matched normal human serum and plasma samples were assayed for total ras p21 protein levels by ELISA, as described above in Materials and Methods. Plasma

levels averaged approximately 24% higher than serum levels, as demonstrated in Table 2 below.

TABLE 2

Matc	hed Normal Hu	man Serum/Plas	sma (pg/ml)
Sample#	Plasma	Serum	% Difference
1602	166.659	146.964	11.8
1603	131.963	121.974	7.6
1604	111.289	87.124	21.7
1605	116.629	87.124	25.3
1606	171.328	102.398	40.2
1607	167.74	102.398	39.0
1608	178.863	106.667	40.4
1609	477.954	147.325	69.2
1610	147.328	140.53	4.6
1611	149.834	85.351	43.0
1612	154.124	127.678	17.2
1613	73.658	79.324	-7.7
1614	116.634	110.937	4.9
1615	125.182	117.341	6.3
4029	256.486	160.924	37.3
4031	132.319	98.486	25.6
Mean	167.37	113.91	24.14
$^{\mathrm{SD}}$	91.9	25.0	19.8

Example 4

Assay Characteristics

[0203] The sensitivity, specificity, precision and parallelism of the ELISA assay for total ras p21 in human serum or plasma was characterized (in all instances, using Mab Ras 17 as the capture antibody and Mab Ras 10.2 as the detector antibody).

1. Analytical Sensitivity

[0204] A minimal detectable concentration of analyte was determined by repeated measurement of a zero dose sample (Standard Level 1) and calculation of mean+2 standard deviations. The ras ELISA assay will detect 4.6 picograms/ml of total ras p21 protein.

2. Specificity

[0205] Cross Reactivity:

[0206] Rab 3A, Rac-1, and Ran are other members of the G-protein family. The Oncogene Science RAS p21 ELISA was challenged with high levels of each analyte. Only Rab-3A showed any cross-reactivity at a minimal 1.22%.

[0207] 3. Precision

[0208] Inter-Assay Precision:

[0209] The reproducibility of the ELISA test for determining total ras p21 protein in human samples was confirmed by repeated testing of the same sera on different occasions (Table 3). Normal human serum and plasma were spiked with recombinant wild-type ras p21 [Invitrogen] at three different concentrations and tested in 8 assays with 8 replicates per test point. Between-assay variability was less than or equal to 10.0%.

TABLE 3

	High	Medium	Low
n	64	64	64
Serum Mean (pg/ml)	472.9	320.0	115.7
Plasma Mean (pg/ml)	371.3	167.8	71.5
Serum Inter-Assay % CV	7.1	8.6	10.0
Plasma Inter-Assay %	8.7	7.5	8.0

[0210] Intra-Assay Precision:

[0211] Normal human serum and plasma were spiked with ras p21 protein at three different concentrations and tested in 8 assays with 8 replicates per test point. Mean intra-assay variability was less than 7% (Table 4).

TABLE 4

	High	Medium	Low
n	64	64	64
Serum Mean (pg/ml)	472.9	320.0	115.7
Plasma Mean (pg/ml)	371.3	167.8	71.5
Serum Intra-Assay % CV	5.8	6.6	6.9
Plasma Intra-Assay % CV	6.5	6.0	6.6

[0212] 4. Parallelism

[0213] Spiked serum and plasma samples were initially-diluted 1:2, subsequent 2 fold dilutions were made, and all dilutions were tested in the ras p21 ELISA. Recoveries (pg/ml) for each dilution were calculated by correcting with the appropriate dilution factor. Results show that spiked plasma and serum samples diluted in sample diluent result in accurate recovery of ras p21 protein (Table 5).

TABLE 5

		Parallelism		
Dilution	OD	pg/ml	OD	pg/ml
	Plasma	(EDTA)	Se	rum
1:2	1.516	485.909	1.707	549.869
1:4	0.869	538.608	0.937	554.674
1:8	0.496	577.219	0.470	564.503
1:16	0.264	534.753	0.237	524.047
	Plasma	(Citrate)	Plasma	(Heparin)
1:2	1.728	556.734	1.689	543.841
1:4	0.932	580.448	0.883	547.645
1:8	0.475	549.780	0.455	522.341
1:16	0.279	573.564	0.262	529.400

Example 5

Ras Levels in Serial Samples from Normal Serum

[0214] Serial serum samples from fifteen normal humans (thirteen males, two females) were assayed by ELISA for total ras p21 levels as described above under Materials and Methods. In general, the samples were drawn at least three weeks apart.

TABLE 6

		TABLE 6		
Seria	l Samples from	Normal individuals (I	Ras Levels)	
	Dog m21			
Sample	Ras p21 (pg/mL)	Date of Draw	Gender	
3651-1	159.0	May 20, 2001	Male	
3651-2	221.1	Jun. 12, 2001		
3651-3	141.3	Jul. 14, 2001		
3652-1	50.6	Mar. 20, 2001	Male	
3652-2	56.0	May 8, 2001		
3652-3	58.8	Jun. 19, 2001		
3653-1	246.7	Jan. 12, 2001	Male	
3653-2	560.0	Apr. 4, 2001		
3653-3	244.4	Sep. 17, 2001		
3654-1	113.9	Jan. 5, 2001	Male	
3654-2	127.9	Jan. 31, 2001		
3654-3	170.2	Feb. 21, 2001		
3654-4	126.7	Sep. 6, 2001		
3655-1	101.0	Mar. 22, 2001	Male	
3655-2	99.6	May 9, 2001		
3655-3	132.3	Jun. 7, 2001		
3656-1	150.3	May 30, 2001	Male	
3656-2	109.4	Aug. 29, 2001		
3656-3	109.7	Sep. 12, 2001		
3657-1	153.8	May 7, 2001	Male	
3657-2	122.3	Jun. 6, 2001		
3657-3	100.7	Jul. 10, 2001		
3658-1	75.8	Apr. 17, 2001	Male	
3658-2	74.0	May 18, 2001		
3658-3	70.8	Jun. 21, 2001		
3658-4	79.0	Sep. 13, 2001		
3659-1	82.7	Jul. 6, 2001	Male	
3659-2	109.2	Aug. 19, 2001		
3659-3	99.7	Sep. 24, 2001		
3660-1	96.3	Sep. 23, 2001	Male	
3660-2	146.3	Oct. 16, 2001		
3660-3	115.0	Nov. 21, 2001		
3661-1	172.0	Aug. 7, 2001	Male	
3661-2	179.1	Aug. 30, 2001		
3661-3	149.5	Oct. 10, 2001	3.6.1	
3662-1	130.6	Sep. 11, 2001	Male	
3662-2	116.7	Oct. 1, 2001		
3662-3	154.3	Dec. 3, 2001	3.6.1	
3663-1	75.5	Aug. 22, 2001	Male	
3663-2	167.4	Sep. 20, 2001		
3663-3	124.2	Oct. 24, 2001	T 1	
3664-1	67.6	May 9, 2000	Female	
3664-2	69.7	Jul. 11, 2000		
3664-3	40.2	Aug. 9, 2000		
3664-4	105.5	Jul. 11, 2001	T 1	
3665-1	86.4	Mar. 13, 2001	Female	
3665-2	121.4	Apr. 25, 2001		
3665-3	86.1	Jun. 20, 2001		

Example 6

Single Point and Serial Ras Levels in Colon Cancer Patients

[0215] Ras is mutated in approximately 20% of all cancers, particularly in colon and pancreatic cancer. Based on the hypothesis of the invention, that a ras pathway activated by either mutation or overexpression may result in elevated lev-

els of total ras p21 protein, 48 sera from colon cancer patients were assayed for total ras p21 levels (Table 7). At least four sera from colon cancer patients (samples #2004, #2065, #2784 and #2785) showed extremely high total ras p21 levels.

TABLE 7

2000 C2 G3 Prog 135.09 2001 C1 3 Stab 62.86 2002 D G4 Recur 57.86 2003 C1 3 Prog 75.80 2004 C2 G2 Recur 1147.18 2005 B2 G1 NED 24.00 2006 B2 G1 Prog 38.39 2007 C1 4 P. Resp 39.44 2008 C2 G3 Prog 121.38 2010 C2 3 Prog 105.74 2011 C2 3 Prog 118.44 2014 C1 4 Prog 118.44 201	Sample	Stage	Grade	Status	Ras p21 (pg/mL)
2002 D G4 Recur 57.86 2003 C1 3 Prog 75.80 2004 C2 G2 Recur 1147.18 2005 B2 G1 NED 24.00 2006 B2 G1 Prog 38.39 2007 C1 4 P. Resp 39.44 2008 C2 G3 Prog 121.38 2010 C2 3 Prog 65.14 2011 C2 3 Prog 105.74 2012 C2 3 Stab 40.04 2011 C2 3 Stab 40.04 2012 C2 3 Stab 40.04 2013 D 4 Prog 118.44 2014 C1 4 P. Resp 57.85 2015 C2 3 P. Resp 98.02 2016 D G3 Prog 61.04 2020<	2000		G3	Prog	135.09
2003 C1 3 Prog 75.80 2004 C2 G2 Recur 1147.18 2005 B2 G1 NED 24.00 2006 B2 G1 Prog 38.39 2007 C1 4 P. Resp 39.44 2008 C2 G3 Prog 121.38 2010 C2 3 Prog 15.74 2011 C2 3 Prog 105.74 2011 C2 3 Prog 105.74 2012 C2 3 Stab 40.04 2013 D 4 Prog 118.44 2014 C1 4 P. Resp 57.85 2015 C2 3 P. Resp 98.02 2016 D G3 Prog 53.02 2017 D 4 Prog 113.46 2019 C2 3 Prog 61.04 2020<	2001			Stab	62.86
2004 C2 G2 Recur 1147.18 2005 B2 G1 NED 24.00 2006 B2 G1 Prog 38.39 2007 C1 4 P. Resp 39.44 2008 C2 G3 Prog 121.38 2010 C2 3 Prog 121.38 2011 C2 3 Prog 121.38 2011 C2 3 Prog 105.74 2012 C2 3 Stab 40.04 2012 C2 3 Stab 40.04 2013 D 4 Prog 118.44 2014 C1 4 Pr. Resp 57.85 2015 C2 3 P. Resp 98.02 2016 D G3 Prog 53.02 2017 D 4 Prog 113.46 2020 C1 G1 Prog 149.69 2	2002	D		Recur	57.86
2005 B2 G1 NED 24.00 2006 B2 G1 Prog 38.39 2007 C1 4 P. Resp 39.44 2008 C2 G3 Prog 121.38 2010 C2 3 Prog 105.74 2011 C2 3 Prog 105.74 2012 C2 3 Stab 40.04 2013 D 4 Prog 118.44 2014 C1 4 P. Resp 57.85 2015 C2 3 P. Resp 98.02 2016 D G3 Prog 113.46 2019 C2 3 Prog 13.46 2019 C2 3 Prog 49.69 2021 D G3 Prog 49.69 2021 D G3 Prog 43.51 2022 C1 G1 Prog 84.97 2025 <td>2003</td> <td>C1</td> <td>3</td> <td>Prog</td> <td>75.80</td>	2003	C1	3	Prog	75.80
2006 B2 G1 Prog 38.39 2007 C1 4 P. Resp 39.44 2008 C2 G3 Prog 121.38 2010 C2 3 Prog 65.14 2011 C2 3 Prog 105.74 2012 C2 3 Stab 40.04 2013 D 4 Prog 118.44 2014 C1 4 P. Resp 57.85 2015 C2 3 P. Resp 98.02 2016 D G3 Prog 53.02 2016 D G3 Prog 53.02 2016 D G3 Prog 98.02 2017 D 4 Prog 913.46 2019 C2 3 Prog 61.04 2020 C1 G1 Prog 49.69 2021 D G3 Prog 49.69 2021	2004	C2	G2	Recur	1147.18
2007 C1 4 P. Resp 39.44 2008 C2 G3 Prog 121.38 2010 C2 3 Prog 65.14 2011 C2 3 Prog 105.74 2012 C2 3 Prog 118.44 2013 D 4 Prog 118.44 2014 C1 4 P. Resp 57.85 2015 C2 3 P. Resp 98.02 2016 D G3 Prog 53.02 2016 D G3 Prog 53.02 2016 D G3 Prog 53.02 2017 D 4 Prog 113.46 2019 C2 3 Prog 61.04 2020 C1 G1 Prog 49.69 2021 D G3 Prog 43.51 2022 C1 G1 Prog 96.02 2025 <td>2005</td> <td>B2</td> <td>G1</td> <td>NED</td> <td>24.00</td>	2005	B2	G1	NED	24.00
2008 C2 G3 Prog 121.38 2010 C2 3 Prog 65.14 2011 C2 3 Prog 105.74 2012 C2 3 Stab 40.04 2013 D 4 Prog 118.44 2014 C1 4 P. Resp 57.85 2015 C2 3 P. Resp 98.02 2016 D G3 Prog 53.02 2016 D G3 Prog 53.02 2017 D 4 Prog 113.46 2019 C2 3 Prog 61.04 2020 C1 G1 Prog 49.69 2021 D G3 Prog 49.69 2021 D G3 Prog 43.51 2022 C1 G1 Prog 84.97 2025 D G4 Prog 96.02 2026	2006	B2	G1	Prog	38.39
2010 C2 3 Prog 65.14 2011 C2 3 Prog 105.74 2012 C2 3 Stab 40.04 2013 D 4 Prog 118.44 2014 C1 4 P. Resp 57.85 2015 C2 3 P. Resp 98.02 2016 D G3 Prog 53.02 2017 D 4 Prog 53.02 2017 D 4 Prog 53.02 2017 D 4 Prog 61.04 2020 C1 G1 Prog 61.04 2020 C1 G1 Prog 49.69 2021 D G3 Prog 43.51 2022 C1 G1 Prog 84.97 2025 D G4 Prog 96.02 2026 C2 G2 P. Resp 54.98 2027	2007	C1	4	P. Resp	39.44
2011 C2 3 Prog 105.74 2012 C2 3 Stab 40.04 2013 D 4 Prog 118.44 2014 C1 4 P. Resp 57.85 2015 C2 3 P. Resp 98.02 2016 D G3 Prog 53.02 2017 D 4 Prog 53.02 2019 C2 3 Prog 61.04 2020 C1 G1 Prog 49.69 2021 D G3 Prog 43.51 2022 C1 G1 Prog 84.97 2025 D G4 Prog 96.02 2027	2008	C2	G3	Prog	121.38
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2013 D 4 Prog 118.44 2014 C1 4 P. Resp 57.85 2015 C2 3 P. Resp 98.02 2016 D G3 Prog 53.02 2017 D 4 Prog 113.46 2019 C2 3 Prog 61.04 2020 C1 G1 Prog 49.69 2021 D G3 Prog 49.69 2021 D G3 Prog 43.51 2022 C1 G1 Prog 84.97 2025 D G4 Prog 96.02 2026 C2 G2 P. Resp 54.98 2027 D G3 Recur 47.42 2028 B2 G2 Stab 73.05 2030 C1 G1 P. Resp 70.46 2034 D G3 P. Resp 76.25 20	2011	C2	3	Prog	105.74
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2015 C2 3 P. Resp 98.02 2016 D G3 Prog 53.02 2017 D 4 Prog 113.46 2019 C2 3 Prog 61.04 2020 C1 G1 Prog 49.69 2021 D G3 Prog 49.69 2022 C1 G1 Prog 49.69 2025 D G4 Prog 96.02 2025 D G4 Prog 96.02 2025 D G4 Prog 96.02 2027 D G3 Recur 47.42 2028 B2 G2 Stab 73.05 2034	2013	D	4	Prog	118.44
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2017 D 4 Prog 113.46 2019 C2 3 Prog 61.04 2020 C1 G1 Prog 49.69 2021 D G3 Prog 43.51 2022 C1 G1 Prog 84.97 2025 D G4 Prog 96.02 2026 C2 G2 P. Resp 54.98 2027 D G3 Recur 47.42 2028 B2 G2 Stab 73.05 2030 C1 G1 P. Resp 70.46 2034 D G3 P. Resp 76.25 2035 C1 G1 Prog 64.53 2036 III 2 NED 38.24 2038 III 2 NED 38.24 2054 IV 2 Prog 150.49 2054 IV 2 Prog 150.49 2055<	2015	C2	3	P. Resp	98.02
2017 D 4 Prog 113.46 2019 C2 3 Prog 61.04 2020 C1 G1 Prog 49.69 2021 D G3 Prog 43.51 2022 C1 G1 Prog 84.97 2025 D G4 Prog 96.02 2026 C2 G2 P. Resp 54.98 2027 D G3 Recur 47.42 2028 B2 G2 Stab 73.05 2030 C1 G1 P. Resp 70.46 2034 D G3 P. Resp 76.25 2035 C1 G1 Prog 64.53 2036 III 2 NED 38.24 2038 III 2 NED 38.24 2054 IV 2 Prog 150.49 2054 IV 2 Prog 150.49 2055<	2016	D	G3		53.02
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2022 C1 G1 Prog 84.97 2025 D G4 Prog 96.02 2026 C2 G2 P. Resp 54.98 2027 D G3 Recur 47.42 2028 B2 G2 Stab 73.05 2030 C1 G1 P. Resp 70.46 2034 D G3 P. Resp 76.25 2035 C1 G1 Prog 64.53 2036 III 2 NED 38.24 2038 III 2 NED 38.24 2052 III 2 NED 38.24 2054 IV 2 Prog 150.49 2055 IV 3 Stab 56.65 2056 III 2 NED 158.35 2057 III 2 Prog 137.91 2058 IV 3 Prog 90.18 2059	2021				
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2030 C1 G1 P. Resp 70.46 2034 D G3 P. Resp 76.25 2035 C1 G1 Prog 64.53 2036 III 2 NED 38.24 2038 III 2 NED 38.24 2052 III 2 NED 38.24 2052 III 2 NED 69.24 2054 IV 2 Prog 150.49 2055 IV 3 Stab 56.65 2056 III 2 NED 158.35 2057 III 2 Prog 137.91 2058 IV 3 Prog 90.18 2059 IV 3 Stab 48.03 2061 IV Unk Stab 198.64 2064 IV 4 Prog 319.91 2065 IV 4 Prog 318.55 2772 </td <td></td> <td>_</td> <td></td> <td></td> <td></td>		_			
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2781 III Unk Unk 209.4 2784 III Unk Unk 664.9					
2784 III Unk Unk 664.9					
	2784	III	Unk Unk	Unk	1032.2
2786 III Unk Unk 1032.2					

[0216] However, serial changes in total ras p21 protein may be a more useful indicator of preneoplastic/neoplastic disease than a single point assay. Increasing levels of ras p21 may be indicative of colorectal cancer recurrence or progression and identify patients who should be treated with either conventional therapies or ras pathway targeted therapies. Increasing levels of ras oncoprotein may be used in conjunction with other oncology tests, such as carcinoembryonic antigen (CEA).

[0217] As shown in Table 8, serial serum samples from seven colon cancer patients were assayed for total ras p21 levels using the ELISA immunoassay format described in Materials and Methods. In one particular patient with stage IV colon cancer, the inventors made the following observa-

tions: the ras level initially remained relatively stable (Table 8, samples #3766-2 to #3766-4), but as the cancer progressed the ras levels increased to 69 pg/ml and then to 124 pg/ml. That patient showed progressive cancer while on conventional therapy. The immunoassay of the invention therefore may be used to identify a patient not responding to conventional therapy who may respond to therapies specifically targeted to inhibit the ras oncogene pathway or components of the pathway. The immunoassays of the invention may also be used to monitor response or lack of response to the ras pathway targeted therapies.

[0218] Another example is that of a stage II colon cancer patient, whose ras p21 level initially remained relatively stable (Table 8, samples #3767-1 to #3767-4). As the cancer progressed, however (samples #3767-5 and #3767-6), the ras level increased to 142 pg/ml and remained elevated at 95 pg/ml, while the patient's CEA level decreased to a normal 1.8 ng/ml. As shown in this patient's case, CEA decreased in the face of progressive disease, whereas ras levels increased, a pattern which is consistent with progressive disease. Elevated levels of ras may therefore be used to identify patients eligible for ras inhibitor therapies, or a patient not responding to conventional therapy that may be responsive to therapies specifically targeted to inhibit the ras oncogene pathway.

TABLE 8

	111111111111	
Serial serum san	ples from Colon Cancer pa	atients (Ras p21 Levels)
Sample	Ras p21(pg/mL)	CEA (ng/mL)
3766-2	59.4	
3766-3	43.7	
3766-4	52.8	
3766-5	69.3	
3766-6	124.8	
3767-1	39.1	9.1
3767-2	46.5	6.5
3767-3	51.2	8.7
3767-4	38.5	6.4
3767-5	142.1	1.8
3767-6	95.6	1.8
3768-1	50.1	3.8
3768-2	76.8	4.5
3768-3	53.4	2.9
3768-4	62.7	2.5
3768-5	63.3	2.7
3768-6	103.3	2.8
3769-1	133.3	
3769-2	100.1	<0.7
3769-3	83.4	<0.7
3769-4	81.6	0.8
3769-5	75.4	
3769-6	87.1	<0.7
3770-1	81.6	6.8
3770-2	97.5	7.4
3770-3	92.7	7.5
3770-4	61.5	9.2
3770-5	93.1	10.0
3770-6	117.0	10.0
3771-1	43.5	26.7
3771-2	70.8	50.1
3771-3	50.9	81.0
3771-4	44.5	171.6
3771-5	41.3	368.7
3771-6	94.7	699.6
3772-1	74.0	1.8
3772-2	50.8	5.8
3772-3	154.8	7.1

TABLE 8-continued

Serial serum sar	nples from Colon Cancer pa	atients (Ras p21 Levels)
Sample	Ras p21(pg/mL)	CEA (ng/mL)
3772-4	50.2	
3772-5	83.4	
3772-6	158.0	20.0

Example 7

Ras p21 Serum Levels in Stage I/Stage II Breast Cancer

[0219] Ras p21 is overexpressed in 50-60% of breast cancers. Pretreatment elevated levels of the circulating ras oncoprotein in metastatic breast cancer (MBC) patients may indicate those patients eligible for either conventional or targeted anti-ras therapies.

[0220] In addition, increasing or decreasing levels of ras p21 alone, or in conjunction with levels of HER-2/neu, in breast cancer patients may be an early indicator of recurrent breast cancer. Pretreatment elevated circulating levels of ras or HER-2/neu oncoproteins in metastatic breast cancer (MBC) patients may indicate those patients eligible for antiras and/or anti-HER-2/neu therapies, either conventional or targeted therapies.

[0221] Table 9 shows total ras p21 levels found by immunoassay (as described under Materials and Methods) of single sera samples from Stage I or Stage II breast cancer patients. Individual breast cancer patients show high total ras p21 levels compared with normal levels in the 100-200 pg/ml range. Several patients show extremely high ras p21 levels in the 2600-2800 pg/ml range (see, for example, samples #1323, #1357 and #1432),

TABLE 9

Stage I Stage II Breast Cancer Serums Ras p21 Ras p21 (pg/mL) Ras p21 (pg/mL) 1318 312.9 1406 99.6 1320 471.8 1407 113.3 1321 2642.0 1408 171.0 1322 53.0 1409 82.6 1323 2646.0 1410 76.2 1324 2648.0 1411 140.2 1325 45.2 1412 53.5 1326 60.0 1413 83.6 1327 45.2 1414 207.6 1338 324.6 1415 76.7 1330 2660.0 1416 291.3 1331 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 284		Single Point Stage I/II Breast Cancer Patient Serum Samples (Ras p21 Levels)					
Sample (pg/mL) Sample (pg/mL) 1318 312.9 1406 99.6 1320 471.8 1407 113.3 1321 2642.0 1408 171.0 1322 53.0 1409 82.6 1323 2646.0 1410 76.2 1324 2648.0 1411 140.2 1325 45.2 1412 53.5 1326 60.0 1413 83.6 1327 45.2 1414 207.6 1328 324.6 1415 76.7 1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1337 2674.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423							
1320 471.8 1407 113.3 1321 2642.0 1408 171.0 1322 53.0 1409 82.6 1323 2646.0 1410 76.2 1324 2648.0 1411 140.2 1325 45.2 1412 53.5 1326 60.0 1413 83.6 1327 45.2 1414 207.6 1328 324.6 1415 76.7 1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4<	Sample		Sample				
1321 2642.0 1408 171.0 1322 53.0 1409 82.6 1323 2646.0 1410 76.2 1324 2648.0 1411 140.2 1325 45.2 1412 53.5 1326 60.0 1413 83.6 1327 45.2 1414 207.6 1328 324.6 1415 76.7 1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 <td>1318</td> <td>312.9</td> <td>1406</td> <td>99.6</td>	1318	312.9	1406	99.6			
1322 53.0 1409 82.6 1323 2646.0 1410 76.2 1324 2648.0 1411 140.2 1325 45.2 1412 53.5 1326 60.0 1413 83.6 1327 45.2 1414 207.6 1328 324.6 1415 76.7 1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0	1320	471.8	1407	113.3			
1323 2646.0 1410 76.2 1324 2648.0 1411 140.2 1325 45.2 1412 53.5 1326 60.0 1413 83.6 1327 45.2 1414 207.6 1328 324.6 1415 76.7 1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0 <td>1321</td> <td>2642.0</td> <td>1408</td> <td>171.0</td>	1321	2642.0	1408	171.0			
1324 2648.0 1411 140.2 1325 45.2 1412 53.5 1326 60.0 1413 83.6 1327 45.2 1414 207.6 1328 324.6 1415 76.7 1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1322	53.0	1409	82.6			
1325 45.2 1412 53.5 1326 60.0 1413 83.6 1327 45.2 1414 207.6 1328 324.6 1415 76.7 1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1323	2646.0	1410	76.2			
1326 60.0 1413 83.6 1327 45.2 1414 207.6 1328 324.6 1415 76.7 1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1324	2648.0	1411	140.2			
1327 45.2 1414 207.6 1328 324.6 1415 76.7 1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1325	45.2	1412	53.5			
1328 324.6 1415 76.7 1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1326	60.0	1413	83.6			
1330 2660.0 1416 291.3 1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1327	45.2	1414	207.6			
1332 66.9 1417 0.0 1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1328	324.6	1415	76.7			
1333 74.5 1418 50.5 1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1330	2660.0	1416	291.3			
1334 106.8 1419 114.7 1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1332	66.9	1417	0.0			
1335 89.6 1420 2840.0 1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1333	74.5	1418	50.5			
1336 2672.0 1421 2842.0 1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1334	106.8	1419	114.7			
1337 2674.0 1422 59.2 1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1335	89.6	1420	2840.0			
1338 90.4 1423 59.0 1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1336	2672.0	1421	2842.0			
1339 56.3 1424 106.3 1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1337	2674.0	1422	59.2			
1340 97.7 1425 110.4 1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1338	90.4	1423	59.0			
1341 66.9 1426 51.0 1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1339	56.3	1424	106.3			
1342 54.2 1427 51.0 1343 50.3 1428 2856.0	1340	97.7	1425	110.4			
1343 50.3 1428 2856.0	1341	66.9	1426	51.0			
	1342	54.2	1427	51.0			
1344 65.1 1429 138.5	1343	50.3	1428	2856.0			
1544 05.1 1425 150.5	1344	65.1	1429	138.5			
1345 295.9 1430 52.9	1345	295.9	1430	52.9			

TABLE 9-continued

	Single Point Stage I/II Breast Cancer Patient Serum Samples (Ras p21 Levels)					
	ge I acer Serums Ras p21		ge II ncer Serums Ras p21			
Sample	(pg/mL)	Sample	(pg/mL)			
1346	77.2	1431	2862.0			
1348	61.5	1432	2864.0			
1349	261.6	1433	44.9			
1350	40.3	1434	63.0			
1351	2702.0	1435	159.4			
1352	62.8	1436	111.7			
1353	56.8	1437	52.8			
1354	2708.0	1438	77.5			
1355	50.5	1439	74.5			
1356	44.3					
1357	2714.0					
1358	2716.0					
1359	2718.0					
1362	101.6					
1365	232.4					
1366	181.0					
1367	85.1					

Example 8

Co-Expression of Ras and TIMP-1 in Human Sera

[0222] Measuring ras p21 levels in combination with other tumor-associated proteins could aid in the selection of therapies for specific cancers. For example, by measuring levels of both ras and proteins upstream of ras, one may determine the most appropriate therapy for a neoplastic disease associated with an activated ras pathway.

[0223] One such upstream protein is TIMP-1 (tissue inhibitor of metalloproteinase-1). TIMP-1 is a multifunctional protein which is found in most tissues and body fluids; it inhibits the proteolytic activity of matrix metalloproteases and thereby prevents cancer invasion. However, TIMP-1 also possesses other functions such as inhibition of apoptosis, induction of malignant transformation and stimulation of cell-growth, and apparently activates ras. TIMP-1 is elevated in blood from colorectal cancer patients, and high TIMP-1 levels predict poor prognosis [38]. Monitoring levels of both ras and TIMP-1 would provide valuable information for therapy selection for a cancer patient, response or lack of response to therapy, as well as determining whether tumor progression is occurring.

[0224] Serial samples of ras p21 and TIMP-1 in colorectal cancer patients may be useful for predicting or monitoring response to targeted therapies (to ras and/or TIMP-1). There may also be a utility in measuring ras and TIMP-1 for predicting response to traditional anti-cancer therapies.

[0225] Pretreatment elevated ras p21 and TIMP-1 levels may be used for selecting patients for treatment with drugs meant to inhibit the ras and/or TIMP-1 proteins, or for predicting patient response to treatment to such drugs. Such treatment includes drugs that inhibit various components of the ras oncogene pathway, such as, raf kinase, and would apply to the raf kinase inhibitor drugs or to famesyltransferase inhibitors (FTIs). Such treatment also includes conventional therapies used against ras driven tumors, or drugs meant to inhibit TIMP-1 function.

Ras and TIMP-1 Levels in Serial Normal Samples

[0226] Serial serum samples from fifteen normal human controls (thirteen males, two females) were assayed by ELISA for total ras p21 and TIMP-1 levels as described above under Materials and Methods (shown in Table 10). In general, the samples were drawn at least three weeks apart. In the normal sera, total ras p21 levels varied between 50.6 pg/ml and 560 pg/ml; normal levels of TIMP-1 ranged between 167.8 ng/ml and 614.4 ng/ml.

TABLE 10

Serial normal serum samples evaluated for ras p21 and TIMP-1 levels Serial Normal Samples					
Sample	Ras p21 (pg/mL)	TIMP-1 (ng/mL)			
3651-1	159.0	242.0	_		
3651-2	221.1	342.9 368.5			
3651-3		214.4			
3652-1	141.3 50.6	330.7			
3652-2	56.0	395.0			
3652-3	58.8	279.8			
3653-1	246.7	201.4			
3653-2	560.0	192.9			
3653-3	244.4	269.0			
3654-1	113.9	354.7			
3654-2	127.9	324.1			
3654-3	170.2	399.2			
3654-4	126.7	471.9			
3655-1	101.0	465.0			
3655-2	99.6	445.9			
3655-3	132.3	430.8			
3656-1	150.3	368.6			
3656-2	109.4	398.3			
3656-3	109.7	336.2			
3657-1	153.8	351.5			
3657-2	122.3	375.3			
3657-3	100.7	238.4			
3658-1	75.8	282.3			
3658-2	74.0	390.5			
3658-3	70.8	348.0			
3658-4	79.0	346.9			
3659-1	82.7	218.7			
3659-2	109.2	167.8			
3659-3	99.7	247.4			
3660-1	96.3	563.5			
3660-2	146.3	532.1			
3660-3	115.0	614.4			
3661-1	172.0	481.5			
3661-2	179.1	349.5			
3661-3	149.5	309.7			
3662-1	130.6	258.5			
3662-2	116.7	446.4			
3662-3	154.3	261.8			
3663-1	75.5	219.6			
3663-2	167.4	195.7			
3663-3	124.2	343.2			
3664-1	67.6	270.3			
3664-2	69.7	279.3			
3664-3	40.2	239.4			
3664-4	105.5	326.3			
3665-1	86.4	318.8			
3665-2	121.4	357.0			
3665-3	86.1	411.1			
3665-4		369.5			

Ras p21, TIMP-1 and CEA Levels in Serial Colon Cancer Samples

[0227] As shown in Table 11, sera drawn serially from seven colon cancer patients were assayed by ELISA for total ras p21 and TIMP-1 levels as described above under Materials and Methods. Sera from six of those patients were also assayed for CEA.

[0228] In one particular stage IV colorectal cancer patient (Table 11, patient #3766), the ras level increased from 59 pg/ml to 125 pg/ml, whereas the TIMP-1 level increased from 263 to 703 ng/ml. The patient showed progressive disease while receiving conventional therapy. That data suggests that total ras p21 and TIMP-1 levels may be used in the management of such colorectal cancer patients.

TABLE 11

Serial Colorectal Cancer Serum Samples Evaluated for ras p21 and TIMP-1 levels							
Sample	Ras p21 (pg/mL)	CEA (ng/mL)	TIMP-1 (ng/mL)	Stage	Grade	Status	
3766-2	59.4		263.2	IV	3	Prog	
3766-3	43.7		258.6	IV	3	Prog	
3766-4	52.8		323.3	IV	3	Prog	
3766-5	69.3		439.1	IV	3	Prog	
3766-6	124.8		702.7	IV	3	Prog	
3767-1	39.1	9.1	383.4	II	1	Prog	
3767-2	46.5	6.5	415.3	II	1	Prog	
3767-3	51.2	8.7	423.6	II	1	Prog	
3767-4	38.5	6.4	438.2	II	1	Prog	
3767-5	142.1	1.8	430.1	II	1	Prog	
3767-6	95.6	1.8	534.5	II	1	Prog	
3768-1	50.1	3.8	390.2	III	2	Stab	
3768-2	76.8	4.5	311.8	III	2	Stab	
3768-3	53.4	2.9	409.5	III	2	Stab	
3768-4	62.7	2.5	262.7	III	2	Stab	
3768-5	63.3	2.7	302.1	III	2	Stab	
3768-6	103.3	2.8	290.0	III	2	Stab	
3769-1	133.3		212.7	III	1	Stab	
3769-2	100.1	< 0.7	197.6	III	1	Stab	
3769-3	83.4	< 0.7	219.6	III	1	Stab	
3769-4	81.6	0.8	181.5	$_{ m III}$	1	Stab	
3769-5	75.4		181.8	III	1	Stab	
3769-6	87.1	< 0.7	188.6	$_{ m III}$	1	Stab	
3770-1	81.6	6.8	488.4	$\Pi\Pi$	3	Prog	
3770-2	97.5	7.4	527.7	III	3	Prog	
3770-3	92.7	7.5	609.9	$\Pi\Pi$	3	Prog	
3770-4	61.5	9.2	724.9	III	3	Prog	
3770-5	93.1	10.0	1210.8	III	3	Prog	
3770-6	117.0	10.0	959.9	III	3	Prog	
3771-1	43.5	26.7	208.7	III	3	Prog	
3771-2	70.8	50.1	282.2	III	3	Prog	
3771-3	50.9	81.0	261.5	III	3	Prog	
3771-4	44.5	171.6	322.6	III	3	Prog	
3771-5	41.3	368.7	415.3	III	3	Prog	
3771-6	94.7	699.6	468.8	III	3	Prog	
3772-1	74.0	1.8	265.8	III	2	Prog	
3772-2	50.8	5.8	323.5	III	2	Prog	
3772-3	154.8	7.1	471.1	III	2	Prog	
3772-4	50.2		460.5	III	2	Prog	
3772-5	83.4		491.5	III	2	Prog	
3772-6	158.0	20.0	490.2	III	2	Prog	

Deposit of Hybridoma Cell Lines

[0229] ATCC Deposits.

[0230] The material listed below was deposited with the American Type Culture Collection (ATCC) now at 10801 University Boulevard, Manassas, Va., 20110-2209 (USA). The deposits were made under the provisions of the Budapest Treaty on the International Recognition of Deposited Microorganisms for the Purposes of Patent Procedure and Regulations thereunder (Budapest Treaty). Maintenance of a viable culture is assured for thirty years from the date of deposit. The hybridomas and plasmids will be made available by the ATCC under the terms of the Budapest Treaty, and subject to an agreement between the Applicants and the ATCC which

assures unrestricted availability of the deposited hybridomas and progeny thereof to the public upon issuance of a pertinent U.S. patent.

[0231] The assignee of the present application has agreed that if a culture of the material(s) on deposit should die or be lost or destroyed when cultivated under suitable conditions, to replace the material(s) promptly upon notification with another of the same.

i.	Hybridoma	Deposit Date	ATCC#
ii.	Ras 10	May 12, 1987	HB 9426
iii.	Ras 17	May 29, 1989	HB 10054

[0232] The description of the foregoing embodiments of the invention has been presented for purposes of illustration and description. They are not intended to be exhaustive or to limit the invention to the precise form disclosed, and obviously many modifications and variations are possible in light of the above teachings. The embodiments were chosen and described in order to explain the principles of the invention and its practical application to enable thereby others skilled in the art to utilize the invention in various embodiments and with various modifications as are suited to the particular use contemplated.

[0233] All references cited herein are hereby incorporated by reference.

- 1. A method to detect a preneoplastic/neoplastic disease associated with an activated ras pathway in a human subject comprising:
 - (a) immunologically detecting and quantifying the average level of total ras p21 protein in samples of a body fluid taken from individuals of a control population;
 - (b) immunologically detecting and quantifying serial changes in total ras p21 protein levels in equivalent samples of body fluid taken from the subject over time; and
 - (c) comparing the levels of total ras p21 protein in the subject's samples to the average level of total ras p21 protein in the control samples;
 - wherein a level of total ras p21 protein in the subject's samples that is above the average level of total ras p21 protein in the control samples is indicative of an activated ras pathway and the presence of preneoplastic/neoplastic disease in the subject.
- 2. The method of claim 1 which is further prognostic for said preneoplastic/neoplastic disease, wherein said levels of total ras p21 protein in the subject's samples relative to the average level of total ras p21 in the control samples, are indicative of a better or poorer prognosis for said subject.
- 3. The method of claim 2, wherein said prognosis is a clinical outcome selected from the group consisting of response rate (RR), complete response clinical benefit (CR), partial response clinical benefit (PR), stable disease clinical benefit (SD), time to progression (TTP), and time to death (TTD).
- **4**. The method of claim **2**, wherein the subject's samples are pretreatment samples.
- 5. The method of claim 1, wherein said body fluid is selected from the group consisting of blood, serum, plasma, urine, saliva, semen, breast exudate, cerebrospinal fluid,

tears, sputum, mucous, lymph, cytosols, ascites, pleural effusions, amniotic fluid, bladder washes and bronchioalveolar lavages.

- 6. The method of claim 1, wherein said body fluid is serum or plasma.
- 7. The method of claim 1, wherein said individuals of the control population of step (a) are of the same gender as the subject.
- **8**. The method of claim **1**, wherein said immunological detection and quantification of steps (a) and (b) is by an immunoassay in the form of a sandwich ELISA or equivalent assay.
- **9**. The method of claim wherein the sandwich ELISA or equivalent assay comprises the use of monoclonal antibodies.
- 10. The method of claim 9, wherein the capture monoclonal antibody is designated Ras 17 which is secreted from hybridoma HB 10054 deposited at the American Type Culture Collection (ATCC), and wherein the detector monoclonal antibody is designated Ras 10 which is secreted from hybridoma HB 9426 deposited at the ATCC.
- 11. The method of claim 10, wherein the detector monoclonal antibody Ras 10 is biotinylated.
- 12. The method of claim 1, wherein said preneoplastic/neoplastic disease associated with an activated ras pathway is selected from the group consisting of colorectal cancer, colon cancer, lung cancer, non-small-cell lung cancer, small-cell lung cancer, acute myelogenous leukemia, thyroid cancer, pancreatic cancer, bladder cancer, kidney cancer, melanoma, breast cancer, prostate cancer, ovarian cancer, cervical cancer, head-and-neck cancer, hepatocellular carcinoma and hematologic malignancies.
- 13. The method of claim 1, further comprising the use of an immunoassay to detect or detect and quantify levels of one or more other proteins in the subject's samples.
- 14. The method of claim 13, wherein said other protein is or said other proteins are selected from the group consisting of inhibitors, oncoproteins, growth factor receptors, angiogenic factors, metastasis proteins, tumor markers, and tumor suppressors.
- 15. The method of claim 14 wherein said inhibitor is tissue inhibitor of metalloproteinase-1 (TIMP-1), said oncoprotein is HER-2/neu, said growth factor receptors are selected from the group consisting of epidermal growth factor receptor (EGFR) and platelet derived growth factor receptor (PDGFR), said angiogenic factor is vascular endothelial growth factor (VEGF), said metastasis protein is urokinase-type plasminogen activator (uPA), said tumor marker is carcinoembryonic antigen (CEA), and said tumor suppressor is p53.
- **16**. A method of therapy selection for a human patient with a preneoplastic/neoplastic disease comprising:
 - (a) immunologically detecting and quantifying the average level of total ras p21 protein in samples of a body fluid taken from individuals of a control population;
 - (b) immunologically detecting and quantifying serial changes in total ras p21 protein levels in equivalent samples of body fluid taken from the patient over time;
 - (c) comparing the levels of total ras p21 protein in the patient's samples to the average level of total ras p21 protein in the control samples; and
 - (d) deciding whether to use conventional cancer therapy and/or ras-directed cancer therapy to treat the patient based upon the differences between the levels of total ras p21 protein in the patient's samples and the average level

- of total ras p21 protein in the control samples, and in view of the serial changes among the levels of total ras p21 protein in the patient's samples.
- 17. The method of claim 16, wherein said patient's samples are pretreatment samples.
- 18. The method of claim 16, wherein if a level of total ras p21 protein in a patient's sample is found to be above the average level of total ras p21 protein in the control samples, and if the serial changes in the total ras p21 protein levels in the patients samples reflect levels that are trending above the average level of total ras p21 protein in the control samples, concluding that the patient has a ras oncogene driven preneoplastic/neoplastic disease, and deciding to use a ras-directed therapy to treat the patient.
- 19. The method of claim 16 which is further prognostic for said preneoplastic/neoplastic disease, wherein said levels of total ras p21 protein in the subject's samples relative to the average level of total ras p21 in the control samples are indicative of a better or poorer prognosis for said subject, and wherein said decision in step (d) is based on said prognosis.
- 20. The method of claim 19, wherein said prognosis is a clinical outcome selected from the group consisting of response rate (RR), complete response clinical benefit (CR), partial response clinical benefit (PR), stable disease clinical benefit (SD), time to progression (TTP), and time to death (TTD).
- 21. The method of claim 16, wherein said ras-directed therapy is selected from the group consisting of farnesyltransferase inhibitors (FTI), tyrosine kinase inhibitors, bis-aryl ureas, and antisense inhibitors of ras.
- 22. The method of claim 21, wherein said ras-directed therapy is the bis aryl-urea sorafenib (BAY 43-9006).
- 23. The method of claim 16, further comprising the use of an immunoassay to detect or detect and quantify levels of one or more other proteins in the patient's samples.
- 24. The method of claim 23, wherein said other protein is or said other proteins are selected from the group consisting of inhibitors, oncoproteins, growth factor receptors, angiogenic factors, metastasis proteins, tumor markers, and tumor suppressors.
- 25. The method of claim 24 wherein said inhibitor is tissue inhibitor of metalloproteinase-1 (TIMP-1), said oncoprotein is HER-2/neu, said growth factor receptors are selected from the group consisting of epidermal growth factor receptor (EGFR) and platelet derived growth factor receptor (PDGFR), said angiogenic factor is vascular endothelial growth factor (VEGF), said metastasis protein is urokinase-type plasminogen activator (uPA), said tumor marker is carcinoembryonic antigen (CEA), and said tumor suppressor is p53.
- **26**. The method of claim **16**, wherein the body fluid samples are from a cancer patient who has not responded to treatment.
- 27. A method of monitoring the status of a preneoplastic/ neoplastic disease in a patient, and/or monitoring how a patient with a preneoplastic/neoplastic disease is responding to a therapy, comprising immunologically detecting and quantifying serial changes in total ras p21 protein levels in samples of a body fluid taken from said patient over time;
 - wherein increasing levels of total ras p21 protein over time indicate disease progression or a negative response to said therapy, and wherein decreasing levels of total ras p21 protein indicate disease remission or a positive response to said therapy.

- 28. The method of claim 27, wherein said therapy is a ras-directed therapy.
- 29. The method of claim 27 which is further prognostic for said preneoplastic/neoplastic disease, wherein said levels of total ras p21 protein in the subject's samples are indicative of a better or poorer prognosis for said subject.
- **30**. The method of claim **29**, wherein said prognosis is a clinical outcome selected from the group consisting of response rate (RR), complete response clinical benefit (CR), partial response clinical benefit (PR), stable disease clinical benefit (SD), time to progression (TTP), and time to death (TTD).
- 31. The method of claim 27, wherein said patient's samples are pretreatment samples.
- 32. The method of claim 27, further comprising the use of an immunoassay to detect or detect and quantify levels of one or more other proteins in the patient's samples.
- 33. The method of claim 32, wherein said other protein is or said other proteins are selected from the group consisting of

- inhibitors, oncoproteins, growth factor receptors, angiogenic factors, metastasis proteins, tumor markers, and tumor suppressors.
- 34. The method of claim 33 wherein said inhibitor is tissue inhibitor of metalloproteinase-1 (TIMP-1), said oncoprotein is HER-2/neu, said growth factor receptors are selected from the group consisting of epidermal growth factor receptor (EGFR) and platelet derived growth factor receptor (PDGFR), said angiogenic factor is vascular endothelial growth factor (VEGF), said metastasis protein is urokinase-type plasminogen activator (uPA), said tumor marker is carcinoembryonic antigen (CEA), and said tumor suppressor is p53.
- 35. The method of claim 27, wherein the body fluid samples are from a cancer patient who has not responded to treatment.
- **36.** The method of claim **34**, wherein increasing levels of total ras p21 and/or HER-2/neu are indicative of a greater probability of early recurrence or metastasis.

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