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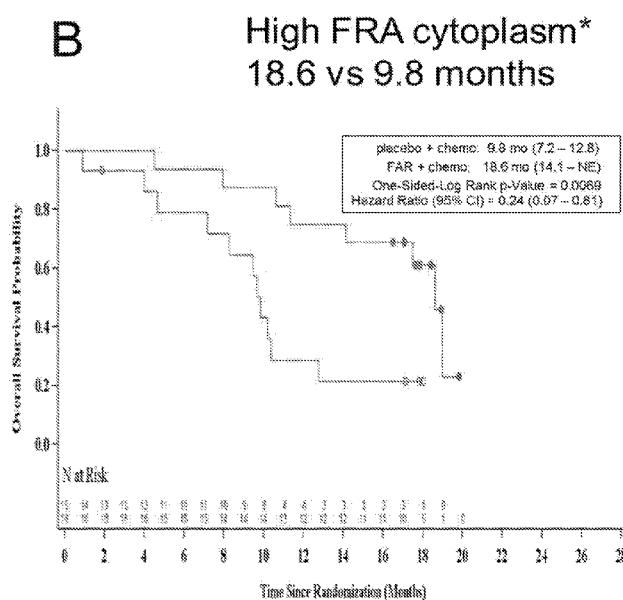
#### Declarations under Rule 4.17:

- as to applicant's entitlement to apply for and be granted a patent (Rule 4.17(ii))
- as to the applicant's entitlement to claim the priority of the earlier application (Rule 4.17(iii))

*[Continued on next page]*

(54) **Title:** METHODS FOR TREATING LUNG CANCER

**Figure 5**



(57) **Abstract:** Provided herein are methods for predicting a likelihood of responsiveness to treatment with a folate receptor alpha (FRA)-targeting agent in a patient having a FRA-expressing lung cancer. Also provided are methods for treating a FRA-expressing lung cancer in a patient with a FRA-targeting agent. The methods involve quantifying or determining the patient's FRA expression level in a biological sample and comparing the patient's FRA expression level to a reference standard used to quantify FRA expression level, wherein the patient is likely to respond to treatment with the FRA-targeting agent if the patient's FRA expression level equals or exceeds the reference FRA expression level. Further provided are related kits for prediction of response and treatment of the identified population of FRA-expressing lung cancer patients.



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## METHODS FOR TREATING LUNG CANCER

### CROSS-REFERENCE TO RELATED APPLICATIONS

**[0001]** This application claims the benefit of U.S. Provisional Application No. 62/149184, filed April 17, 2015, the content of which is incorporated herein by reference in its entirety.

### SEQUENCE LISTING

**[0002]** The instant application contains a Sequence Listing which has been submitted electronically in ASCII format and is hereby incorporated by reference in its entirety. Said ASCII copy, created on April 13, 2016, is named 104018.000950\_SL.txt and is 30,473 bytes in size.

### TECHNICAL FIELD

**[0003]** The subject matter described herein relates to methods for predicting a likelihood of responsiveness to treatment with a folate receptor alpha (FRA)-targeting agent in a patient having a FRA-expressing lung cancer and to methods of treating FRA-expressing lung cancer in a patient with a FRA-targeting agent.

### BACKGROUND

**[0004]** According to the American Cancer Society, an estimated 221,200 new cases of lung and bronchus cancer will be diagnosed in the United States in 2015, accounting for about 14% of all new cancer cases. In addition, an estimated 158,040 deaths from lung and bronchus cancer will occur in the United States in 2015, accounting for about 27% of all cancer deaths. Approximately 84% of all lung cancers are non-small cell lung cancer (NSCLC), and the five year survival rate is only 18%. American Cancer Society. *Cancer Facts & Figures 2015*. Atlanta: American Cancer Society. 2015.

**[0005]** Advanced NSCLC remains a difficult to treat cancer where durable long-term survival is exceedingly rare. Chemotherapy, especially platinum-based doublets, is the established treatment for patients with absent or unknown activating mutations for targeted therapy. Histology is an increasingly significant factor when

selecting chemotherapy for patients with advanced NSCLC. (Li et al., *J Clin Oncol* 2013;31:1039-1049.) A large Phase 3 study in advanced NSCLC found cisplatin plus pemetrexed provided statistically superior overall survival (OS) when compared with cisplatin plus gemcitabine in subjects with either adenocarcinoma or large-cell carcinoma. (Scagliotti, et al., *J Clin Oncol* 2008;26:3543-3551.) However, in the same study, those subjects with squamous cell histology had a better outcome when treated with gemcitabine plus cisplatin.

**[0006]** A number of driver mutations have been identified recently, allowing for targeted therapies with better outcomes in NSCLC. The Lung Cancer Mutation Consortium found such “actionable” mutations in 64% of the 1,007 patients with adenocarcinoma that they assessed, the most common being *KRAS*, *EGFR*, and *ALK*. The use of targeted therapies in this study resulted in median OS of 3.5 years in the subset of patients with oncogenic drivers who received targeted therapy versus OS of 2.4 years in those patients with identified mutations but no targeted therapy and 2.1 years in those who did not demonstrate oncogenic drivers ( $P = 0.001$ ). (Kris et al., *JAMA* 2014;311:1998-2006.)

**[0007]** Folate receptor alpha (FRA) is a cell surface GPI-anchored protein whose expression and biology is associated with a number of malignant cell types. Antibodies to FRA have been developed and tested in preclinical and clinical studies to evaluate their effect on suppressing cancer growth in patients whose tumors express this antigen. Recently trials in ovarian and lung cancer have been conducted to test the effect of anti-FRA antibodies on these diseases (Armstrong et al., *Gynecol. Oncol.* 2013 Jun;129(3):452-8; Thomas et al., *Lung Cancer*. 2013; 80(1):15-8.). Results from these trials demonstrate that the anti-FRA therapy has failed to provide a statistically significant clinical benefit in broad, non-enriched or biomarker-selected, heterogeneous intent-to-treat populations which consist in part of patients with cancers exhibiting varying levels of the FRA antigen as well as other factors that may affect the pharmacologic activity(s) of anti-FRA antibodies (Vergote et al., *Cancer Metastasis Rev.* 2015 Jan 7, DOI 10.1007/s10555-014-9539-8; Thomas et al., *Lung Cancer*. 2013; 80(1):15-8.). While most anti-FRA antibody clinical studies have included all comers, none has selected patients based on FRA expression levels

to determine if threshold levels of FRA expression in the tumor may correlate with therapeutic response to anti-FRA antibody therapy. Currently no data suggesting a correlation of the expression level of FRA to the therapeutic response to an anti-FRA antibody therapy exists. Indeed, findings of a correlative relationship between tumor antigen expression and response to antigen-targeted therapy have been difficult to ascertain for a number of tumor antigens. One of the best studied tumor antigens, HER2, is one of the many examples demonstrating lack of expression-clinical benefit correlation whereby patients treated with the anti-HER2 antibody Herceptin (trastuzumab) have been found to have clinical benefit regardless of high or low tumor antigen (Paik et.al; N Engl J Med 2008; 358:1409-1411).

**[0008]** A pressing need exists for methods for identifying NSCLC patients who would be responsive to treatment regimens that target FRA. The methods and kits described herein satisfy this need.

## SUMMARY

**[0009]** Provided herein are methods for predicting a likelihood of responsiveness to treatment with a folate receptor alpha (FRA)-targeting agent in a patient having folate receptor alpha (FRA)-expressing lung cancer. The methods for predicting such likelihood of responsiveness to treatment involve determining the patient's FRA expression level in a biological sample; and comparing the patient's FRA expression level to a reference FRA expression level. The patient is likely to respond to treatment with a FRA-targeting agent if the patient's FRA expression level equals or exceeds the reference FRA expression level.

**[0010]** Also provided herein are methods for treating folate receptor alpha FRA-expressing lung cancer in a patient with a FRA-targeting agent. Such methods involve determining the patient's FRA expression level in a biological sample; comparing the patient's FRA expression level to a reference FRA expression level; and administering the FRA-targeting agent to the patient if the patient's FRA expression level equals or exceeds the reference FRA expression level. In some embodiments, a chemotherapeutic agent (e.g., standard of care chemotherapy as

described herein) is administered to the patient with or without the FRA-targeting agent.

**[0011]** In some embodiments of the methods for predicting a likelihood of responsiveness to treatment with a folate receptor alpha (FRA)-targeting agent and the methods for treating folate receptor alpha FRA-expressing lung cancer in a patient with a FRA-targeting agent described herein, the FRA-targeting agent comprises an antibody that immunospecifically binds FRA, or an antigen-binding fragment thereof. In preferred embodiments, such antibody that immunospecifically binds FRA comprises farletuzumab. In some embodiments, the antibody that immunospecifically binds FRA is conjugated to a toxin, such as, for example, a microtubule inhibitor, a DNA damaging agent (e.g., a radionuclide), a DNA repair inhibitor, or a signal transduction inhibitor. An exemplary antibody-drug conjugate that can be employed as a FRA-targeting agent in accordance with the methods described herein is IMGN853. In some embodiments of the methods described herein, the FRA-targeting agent is vintafolide.

**[0012]** In some embodiments of the methods described herein, the FRA-expressing lung cancer is FRA-expressing non-small cell lung cancer (NSCLC). In some embodiments, the NSCLC is adenocarcinoma.

**[0013]** For each of the aforementioned methods, the reference FRA expression level corresponds to the FRA expression level above which a patient population afflicted with the FRA-expressing lung cancer that is administered the FRA-targeting agent demonstrates a statistically significant improvement in at least one clinical outcome relative to a patient population afflicted with the FRA-expressing lung cancer that is administered placebo. The improved clinical outcome may be, for example, progression free survival and/or overall survival. In some embodiments, the reference FRA expression level corresponds to 42% +1 or greater anti-FRA staining as described herein. In some embodiments, the reference FRA expression level is 21% +2 or greater anti-FRA staining as described herein. In some embodiments, the reference FRA expression level is 14% +3 or greater staining as described herein. In some embodiments, the reference FRA expression level is a

FRAMSCOR (or M-score) of 7. In some embodiments, the reference FRA expression level is a HBSCOR of 0.25.

**[0014]** In some embodiments, the patient populations afflicted with the FRA-expressing lung cancer are further administered a chemotherapeutic agent such as, for example, a taxane, a platinum-containing compound (e.g., cisplatin, carboplatin), and antifolate (e.g., pemetrexed), or any combination thereof.

**[0015]** FRA expression level may be measured in accordance with the methods described herein by protein quantification or RNA quantification. Either cytoplasmic or membranous FRA expression may be measured. In a preferred embodiment, the FRA expression level is measured by immunohistochemical analysis. In preferred embodiments, the FRA expression level is determined by immunoassay with at least one of the following antibodies:

- (a) an antibody that binds the same epitope as farletuzumab;
- (b) an antibody comprising SEQ ID NO:1 (GFTFSGYGLS) as CDRH1, SEQ ID NO:2 (MISSGGSYTYYADSVKG) as CDRH2, SEQ ID NO:3 (HGDDPAWFAY) as CDRH3, SEQ ID NO:4 (SVSSSISSNNLH) as CDRL1, SEQ ID NO:5 (GTSNLAS) as CDRL2 and SEQ ID NO:6 (QQWSSYPYMYT) as CDRL3;
- (c) an antibody comprising a mature light chain variable region comprising the amino acid sequence of SEQ ID NO:7 and a mature heavy chain variable region comprising the amino acid SEQ ID NO: 8;
- (d) farletuzumab;
- (e) the 548908 antibody;
- (f) an antibody that binds the same epitope as the 548908 antibody;
- (g) the 6D398 antibody;
- (h) an antibody that binds the same epitope as the 6D398 antibody;
- (i) the BN3.2 antibody (Leica Biosystems);
- (j) an antibody that binds the same epitope as the BN3.2 antibody;
- (k) an antibody that binds the same epitope as the 26B3 antibody;
- (l) an antibody comprising SEQ ID NO: 14 (GYFMN) as CDRH1, SEQ ID NO: 15 (RIFPYNGDTFYNQKFKG) as CDRH2, SEQ ID NO: 16 (GTHYFDY) as

CDRH3, SEQ ID NO: 17 (RTSENIFSYLA) as CDRL1, SEQ ID NO: 18 (NAKTLAE) as CDRL2 and SEQ ID NO: 19 (QHHYAFPWT) as CDRL3;

(m) the 26B3 antibody;

(n) an antibody that binds the same epitope as the 19D4 antibody;

(o) an antibody comprising SEQ ID NO: 20 (HPYMH) as CDRH1, SEQ ID NO: 21 (RIDPANGNTKYDPKFQG) as CDRH2, SEQ ID NO: 22 (EEVADYTM DY) as CDRH3, SEQ ID NO: 23 (RASESVDTYGNNFIH) as CDRL1, SEQ ID NO: 24 (LASNLES) as CDRL2 and SEQ ID NO: 25 (QQNNGDPWT) as CDRL3;

(p) the 19D4 antibody;

(q) an antibody that binds the same epitope as the 9F3 antibody;

(r) an antibody comprising SEQ ID NO: 26 (SGYYWN) as CDRH1, SEQ ID NO: 27 (YIKSDGSNNYNPSLKN) as CDRH2, SEQ ID NO: 28 (EWKAMDY) as CDRH3, SEQ ID NO: 29 (RASSTVSYSYLH) as CDRL1, SEQ ID NO: 30 (GTSNLAS) as CDRL2 and SEQ ID NO: 31 (QQYSGYPLT) as CDRL3;

(s) the 9F3 antibody;

(t) an antibody that binds the same epitope as the 24F12 antibody;

(u) an antibody comprising SEQ ID NO: 32 (SYAMS) as CDRH1, SEQ ID NO: 33 (EIGSGGSYTYYPDTVTG) as CDRH2, SEQ ID NO: 34 (ETTAGYFDY) as CDRH3, SEQ ID NO: 35 (SASQGINNFLN) as CDRL1, SEQ ID NO: 36 (YTSSLHS) as CDRL2 and SEQ ID NO: 37 (QHFSKLPWT) as CDRL3;

(v) the 24F12 antibody;

(w) an antibody that comprises a variable region light chain selected from the group consisting of:

- (i) LK26HuVK;
- (ii) LK26HuVKY;
- (iii) LK26HuVKPW; and
- (iv) LK26HuVKPW,Y;

(x) an antibody that comprises a variable region heavy chain selected from the group consisting of:

- (i) LK26HuVH;

- (ii) LK26HuVH FAIS,N;
- (iii) LK26HuVH SLF;
- (iv) LK26HuVH LI;
- (v) LK26KOLHuVH;

(y) an antibody that comprises the heavy chain variable region LK26KOLHuVH (SEQ ID NO: 46) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41);

(z) an antibody that comprises the heavy chain variable region LK26HuVH SLF (SEQ ID NO: 44) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41);

(aa) an antibody that comprises the heavy chain variable region LK26HuVH FAIS,N (SEQ ID NO: 43) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41); and

(bb) the murine monoclonal LK26 antibody.

**[0016]** In the methods described herein, FRA expression level may be assessed by digital imaging technology or manual pathology quantification. In some embodiments, the FRA expression level is assessed by FRAMSCOR or HBSCOR.

**[0017]** The biological sample assayed in accordance with the methods described herein may be urine, blood, serum, plasma, saliva, circulating cells, circulating tumor cells, or tumor tissue (e.g., pleural tissue). In some embodiments, the biological sample comprises pleural cells derived from effusion.

**[0018]** The described methods may further involve administering a chemotherapeutic agent to the patient regardless of whether the patient's FRA expression level equals or exceeds the reference FRA expression level. In a preferred embodiment, the chemotherapeutic agent comprises a platinum-containing compound, such as cisplatin or carboplatin; a taxane (for example, paclitaxel); an antifolate (e.g., pemetrexed); or any combination thereof. For example, in the methods described herein, the patient may be administered carboplatin and paclitaxel; carboplatin and pemetrexed; or cisplatin and pemetrexed.

**[0019]** Additional aspects of the summarized subject matter are provided in greater detail in the detailed description and provided examples and associated figures.

#### BRIEF DESCRIPTION OF THE DRAWINGS

**[0020]** The summary, as well as the following detailed description, is further understood when read in conjunction with the appended drawings. For the purpose of illustrating the disclosed methods, there are shown in the drawings exemplary embodiments of the methods; however, the methods are not limited to the specific embodiments disclosed. In the drawings:

**[0021]** Figure 1 shows pleural tissue suitability for FRA quantification via FRAMSCOR and HBSCOR. Normal (left top panel) or malignant non-small cell lung adenocarcinoma (NSCLC) tissue samples (all other panels) were procured from patients. Tissues were formalin-fixed and sectioned onto glass slides and stained for FRA expression using the 26B3 anti-FRA antibody. Stained slides were visualized by microscopy and documented via photography. This figure represents pleural tissue specimen and staining that is suitable for quantified FRA cytoplasmic or membrane staining (top row). Specimens with poor tissue preservation, poor tissue morphology or over staining as well as those consisting of malignant cells in pleural effusions, not pleural tissue, are omitted from the FRA expression clinical outcome correlation study. Examples of the latter are shown on the bottom row. Cells derived from effusions may be used to measure the minimal FRA levels needed for anti-FRA therapeutic response.

**[0022]** Figure 2 shows a representative sample of a reference data set used for scoring +1 (low expression), +2 (moderate levels) and +3 (high levels) of FRA expression in malignant pleural tissue.

**[0023]** Figure 3 shows a representative expression cut-point analysis measuring cytoplasmic levels of FRA to clinical outcome (overall survival) using HBSCOR method. A significant improvement in response is observed in patients whose FRA expression level exceeds a HBSCOR of ~0.29 or greater. At this level,

significant clinical responses (Hazard Ratios of < 0.5) are observed as compared to patients whose tumor FRA expression level is less than this value. Positive clinical outcomes (Hazard Ratio < 0.7) are observed in patients treated with farletuzumab with a HBSCOR of 0.25 or greater.

**[0024]** Figures 4A and 4B illustrate clinical responses of patients treated with standard-of-care (SOC) chemotherapy +/- farletuzumab and expressing high levels of membrane localized (FRAMSCOR) FRA. Panel A shows a representative measurement of overall survival (OS) in patients with high levels of FRA using FRAMSCOR method whereby patients treated with farletuzumab + SOC (blue line; triangle) and a FRAMSCOR of greater than 7 showed a clinically meaningful improvement in OS, over placebo + SOC treated patients (red line; circle) 18.3 mo vs 10.0 mo (Hazard Ratio 0.54; p = 0.0266), respectively. Similar clinical benefit was observed when measuring PFS. Panel B shows that patients treated with farletuzumab + SOC (blue line; triangle) with a FRAMSCOR of less than 7 did not have a clinically meaningful improvement in OS over placebo + SOC treated patients (red line; circle) (p = 0.386).

**[0025]** Figures 5A and 5B illustrate clinical responses of patients treated with standard-of-care chemotherapy +/- farletuzumab and expressing high levels of cytoplasmic FRA (Panel B) vs patients expressing low levels of cytoplasmic FRA (Panel A) as determined using the HBSCOR method. Panel A shows a representative measurement of overall survival (OS) in patients with low levels of FRA whereby patients having a suboptimal HBSCOR (HBSCOR < 0.38) had no clinical improvement when treated with farletuzumab (closed circle) as compared to placebo control (open circle) (Hazard Ratio 1.03; p = 0.5389). Panel B shows statistically significant improvement in overall survival in patients having an optimal HBSCOR (HBSCOR  $\geq$  0.38; Hazard Ratio 0.24; p = 0.0069) which translates into an 8.8 month improvement in overall survival. Similar effects were observed when measuring PFS.

#### DETAILED DESCRIPTION OF ILLUSTRATIVE EMBODIMENTS

**[0026]** The disclosed methods may be understood more readily by reference to the following detailed description taken in connection with the accompanying figures, which form a part of this disclosure. It is to be understood that the disclosed methods are not limited to the specific methods described and/or shown herein, and that the terminology used herein is for the purpose of describing particular embodiments by way of example only and is not intended to be limiting of the claimed methods.

**[0027]** Similarly, unless specifically otherwise stated, any description as to a possible mechanism or mode of action or reason for improvement is meant to be illustrative only, and the disclosed methods are not to be constrained by the correctness or incorrectness of any such suggested mechanism or mode of action or reason for improvement.

**[0028]** It is to be appreciated that certain features of the disclosed methods which are, for clarity, described herein in the context of separate embodiments, may also be provided in combination in a single embodiment. Conversely, various features of the disclosed methods that are, for brevity, described in the context of a single embodiment, may also be provided separately or in any subcombination.

**[0029]** As used herein, the singular forms “a,” “an,” and “the” include the plural.

**[0030]** The term “about” as used herein when referring to a measurable value such as an amount, a temporal duration, and the like, is meant to encompass variations of up to  $\pm 10\%$  from the specified value, as such variations are appropriate to perform the disclosed methods. Unless otherwise indicated, all numbers expressing quantities of ingredients, properties such as molecular weight, reaction conditions, and so forth used in the specification and claims are to be understood as being modified in all instances by the term “about.” Accordingly, unless indicated to the contrary, the numerical parameters set forth in the following specification and attached claims are approximations that may vary depending upon the desired properties sought to be obtained by the present invention. At the very least, and not as an attempt to limit the application of the doctrine of equivalents to the scope of the

claims, each numerical parameter should at least be construed in light of the number of reported significant digits and by applying ordinary rounding techniques.

**[0031]** The term "antibody" refers to (a) immunoglobulin polypeptides, i.e., polypeptides of the immunoglobulin family that contain an antigen binding site that specifically binds to a specific antigen (e.g., folate receptor alpha), including all immunoglobulin isotypes (IgG, IgA, IgE, IgM, IgD, and IgY), classes (e.g. IgG1, IgG2, IgG3, IgG4, IgA1, IgA2), subclasses, and various monomeric and polymeric forms of each isotype, unless otherwise specified, and (b) conservatively substituted variants of such immunoglobulin polypeptides that immunospecifically bind to the antigen (e.g., folate receptor alpha). Antibodies are generally described in, for example, Harlow & Lane, *Antibodies: A Laboratory Manual* (Cold Spring Harbor Laboratory Press, 1988). Unless otherwise apparent from the context, reference to an antibody also includes antibody derivatives as described in more detail below.

**[0032]** "Antibody fragments" comprise a portion of a full length antibody, generally the antigen-binding or variable region thereof, such as Fab, Fab', F(ab')<sub>2</sub>, and Fv fragments; diabodies; biobodies; linear antibodies; single-chain antibody molecules; and multispecific antibodies formed from antibody fragments. Various techniques have been developed for the production of antibody fragments, including proteolytic digestion of antibodies and recombinant production in host cells; however, other techniques for the production of antibody fragments will be apparent to the skilled practitioner. In some embodiments, the antibody fragment of choice is a single chain Fv fragment (scFv). "Single-chain Fv" or "scFv" antibody fragments comprise the V<sub>H</sub> and V<sub>L</sub> 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<sub>H</sub> and V<sub>L</sub> domains which enables the scFv to form the desired structure for antigen binding. For a review of scFv and other antibody fragments, see James D. Marks, *Antibody Engineering*, Chapter 2, Oxford University Press (1995) (Carl K. Borrebaeck, Ed.).

**[0033]** An "antibody derivative" means an antibody, as defined above, that is modified by covalent attachment of a heterologous molecule such as, e.g., by attachment of a heterologous polypeptide (e.g., a cytotoxin) or therapeutic agent

(e.g., a chemotherapeutic agent), or by glycosylation, deglycosylation, acetylation or phosphorylation not normally associated with the antibody, and the like.

**[0034]** The term "monoclonal antibody" refers to an antibody that is derived from a single cell clone, including any eukaryotic or prokaryotic cell clone, or a phage clone, and not the method by which it is produced. Thus, the term "monoclonal antibody" is not limited to antibodies produced through hybridoma technology.

**[0035]** An "antigen" is an entity to which an antibody specifically binds. For example, folate receptor alpha is the antigen to which an anti-folate receptor-alpha antibody specifically binds.

**[0036]** A FRA targeting agent is a therapeutic agent that targets or exerts its effect through FRA. FRA targeting agents include but are not limited to FRA-binding proteins, such as antibodies that immunospecifically bind FRA and antigen-binding fragments thereof such as farletuzumab; drug conjugates of such antibodies and antigen-binding fragments such as IMGN853 (Immunogen); and small molecules such as vintafolide (EC 145; Endocyte). Vintafolide is a folate-desacetylvinblastine monohydrate conjugate, which allows liberation of the drug into the cytoplasm of cancerous cells via the FRA and endocytosis. In preferred embodiments, the FRA targeting agent is farletuzumab.

**[0037]** The terms "cancer" and "tumor" are well known in the art and refer to the presence, e.g., in a subject, of cells possessing characteristics typical of cancer-causing cells, such as uncontrolled proliferation, immortality, metastatic potential, rapid growth and proliferation rate, and certain characteristic morphological features. Cancer cells are often in the form of a tumor, but such cells may exist alone within a subject, or may be non-tumorigenic cancer cells, such as leukemia cells. As used herein, the term "cancer" includes pre-malignant as well as malignant cancers.

**[0038]** As used herein, the term "folate receptor alpha" (also referred to as FRA, FR-alpha, FOLR-1 or FOLR1) refers to the alpha isoform of the high affinity receptor for folate. Membrane bound FRA is attached to the cell surface by a glycosyl phosphatidyl inositol (GPI) anchor, recycles between membrane and endocytic compartments and is capable of transporting folate into the cell. FRA is

expressed in a variety of epithelial tissues including those of the female reproductive tract, placenta, breast, kidney proximal tubules, choroid plexus, lung and salivary glands. Soluble forms of FRA include but are not limited to those derived by the action of proteases or phospholipase on membrane anchored folate receptors.

**[0039]** The consensus nucleotide and amino acid sequences for human FRA are set forth herein as SEQ ID NOs: 9 and 10, respectively.

SEQ ID NO: 9

tcaaggtaa acgacaagga cagacatggc tcagcggatg acaacacagc tgctgctcct	60
tctagtgtgg gtggctgttag taggggaggc tcagacaagg attgcatggg ccaggactga	120
gcttctcaat gtctgcataa acgccaagca ccacaaggaa aagccaggcc ccgaggacaa	180
gttgcatacg cagtgtcgac cctggaggaa gaatgcctgc tgttctacca acaccagcca	240
ggaagcccat aaggatgttt cctacctata tagattcaac tggaaacctt gtggagagat	300
ggcacctgccc tgcaaaacggc atttcatcca ggacacctgc ctctacgagt gctcccccaa	360
cttggggccc tggatccagc aggtggatca gagctggcgcc aaagagcggg tactgaacgt	420
ccccctgtgc aaagaggact gtgagcaatg gtggaaagat tgctgcaccc cctacacctg	480
caagagcaac tggcacaagg gcttggactg gacttcaggg tttaacaagt ggcgactggg	540
agctgcctgc caacctttcc atttctactt ccccacaccc actgttctgt gcaatgaaat	600
ctggactcacttccatcaagg tcagcaacta cagccgaggg agtggccgct gcatccagat	660
gtggttcgac ccagcccagg gcaaccccaa tgaggaggtg gcgagggtct atgctgcagc	720
catgagtgccg gctggggccat gggcagcctg gcctttctg cttagcctgg ccctaattgt	780
gctgtggctg ctcagctgac ctcctttac cttctgatac ctggaaatcc ctgcctgtt	840
cagccccaca gctcccaact atttggttcc tgctccatgg tcgggcctct gacagccact	900
ttgaataaac cagacaccgc acatgtgtct tgagaattat ttggaaaaaaaaaaaaaaaaaa	960
aa	962

SEQ ID NO: 10

Met Ala Gln Arg Met Thr Thr Gln Leu Leu Leu Leu Val Trp Val
Ala Val Val Gly Glu Ala Gln Thr Arg Ile Ala Trp Ala Arg Thr Glu
Leu Leu Asn Val Cys Met Asn Ala Lys His His Lys Glu Lys Pro Gly
Pro Glu Asp Lys Leu His Glu Gln Cys Arg Pro Trp Arg Lys Asn Ala

Cys Cys Ser Thr Asn Thr Ser Gln Glu Ala His Lys Asp Val Ser Tyr  
Leu Tyr Arg Phe Asn Trp Asn His Cys Gly Glu Met Ala Pro Ala Cys  
Lys Arg His Phe Ile Gln Asp Thr Cys Leu Tyr Glu Cys Ser Pro Asn  
Leu Gly Pro Trp Ile Gln Gln Val Asp Gln Ser Trp Arg Lys Glu Arg  
Val Leu Asn Val Pro Leu Cys Lys Glu Asp Cys Glu Gln Trp Trp Glu  
Asp Cys Arg Thr Ser Tyr Thr Cys Lys Ser Asn Trp His Lys Gly Trp  
Asn Trp Thr Ser Gly Phe Asn Lys Cys Ala Val Gly Ala Ala Cys Gln  
Pro Phe His Phe Tyr Phe Pro Thr Pro Thr Val Leu Cys Asn Glu Ile  
Trp Thr His Ser Tyr Lys Val Ser Asn Tyr Ser Arg Gly Ser Gly Arg  
Cys Ile Gln Met Trp Phe Asp Pro Ala Gln Gly Asn Pro Asn Glu Glu  
Val Ala Arg Phe Tyr Ala Ala Ala Met Ser Gly Ala Gly Pro Trp Ala  
Ala Trp Pro Phe Leu Leu Ser Leu Ala Leu Met Leu Leu Trp Leu Leu  
Ser

Variants, for example, naturally occurring allelic variants or sequences containing at least one amino acid substitution, are encompassed by the terms as used herein.

**[0040]** As used herein, the term “not bound to a cell” refers to a protein that is not attached to the cellular membrane of a cell, such as a cancerous cell. In a particular embodiment, the FRA not bound to a cell is unbound to any cell and is freely floating or solubilized in biological fluids, e.g., urine, serum, plasma, or pleural effusion. For example, a protein that is not bound to a cell may be shed, secreted or exported from normal or cancerous cells, for example, from the surface of cancerous cells, into biological fluids.

**[0041]** The “level” or “expression level” of a specified RNA, as used herein, refers to the level of the RNA as determined using any method known in the art for the measurement of RNA levels. Such methods include, but are not limited to, spectrophotometry (e.g., ultraviolet absorbance), fluorometry, hybridization assays, and microcapillary electrophoresis.

**[0042]** The “level” or “expression level” of a specified protein, as used herein, refers to the level of the protein as determined using any method known in

the art for the measurement of protein levels. Such methods include, for example, electrophoresis, capillary electrophoresis, high performance liquid chromatography (HPLC), thin layer chromatography (TLC), hyperdiffusion chromatography, fluid or gel precipitation reactions, absorption spectroscopy, colorimetric assays, spectrophotometric assays, flow cytometry, immunodiffusion (single or double), solution phase assay, immunofluorimetry, immunoprecipitation, equilibrium dialysis, immunodiffusion, solution phase assay, immunoelectrophoresis, Western blotting, radioimmunoassay (RIA), enzyme-linked immunosorbent assays (ELISAs), immunofluorescent assays, and electrochemiluminescence immunoassay (exemplified below), and the like. In a preferred embodiment, the level is determined using antibody-based techniques, as described in more detail herein.

**[0043]** Antibodies used in immunoassays to determine the level of expression of a specified protein, such as for example FRA, may be labeled with a detectable label. The term "labeled", with regard to the binding agent or antibody, is intended to encompass direct labeling of the binding agent or antibody by coupling (i.e., physically linking) a detectable substance to the binding agent or antibody, as well as indirect labeling of the binding agent or antibody by reactivity with another reagent that is directly labeled. An example of indirect labeling includes detection of a primary antibody using a fluorescently labeled secondary antibody. In one embodiment, the antibody is labeled, e.g., radio-labeled, chromophore-labeled, fluorophore-labeled, or enzyme-labeled. In another embodiment, the antibody is an antibody derivative (e.g., an antibody conjugated with a substrate or with the protein or ligand of a protein-ligand pair (e.g., biotin-streptavidin), or an antibody fragment (e.g., a single-chain antibody, an isolated antibody hypervariable domain).

**[0044]** Expression levels of a specific marker (e.g., FRA) may be determined by any methods of RNA or protein quantification known in the art. Such methods include, but are not limited to, spectrophotometry (e.g., ultraviolet absorbance), fluorometry, hybridization assays, electrophoresis, capillary electrophoresis, high performance liquid chromatography (HPLC), thin layer chromatography (TLC), hyperdiffusion chromatography, fluid or gel precipitation reactions, absorption spectroscopy, colorimetric assays, spectrophotometric assays,

flow cytometry, immunodiffusion (single or double), solution phase assay, immunofluorimetry, immunoprecipitation, equilibrium dialysis, immunodiffusion, solution phase assay, immunoelectrophoresis, Western blotting, radioimmunoassay (RIA), enzyme-linked immunosorbent assays (ELISAs), immunofluorescent assays, and electrochemiluminescence immunoassay (exemplified below), and the like. In a preferred embodiment, the level is determined using antibody-based techniques, as described in more detail herein. In a preferred embodiment, immunohistochemical analysis (e.g., of tumor tissue) is employed. In one embodiment, proteomic methods, e.g., mass spectrometry, are used. Mass spectrometry is an analytical technique that consists of ionizing chemical compounds to generate charged molecules (or fragments thereof) and measuring their mass-to-charge ratios. In a typical mass spectrometry procedure, a sample is obtained from a subject, loaded onto the mass spectrometry, and its components (e.g., FRA) are ionized by different methods (e.g., by impacting them with an electron beam), resulting in the formation of charged particles (ions). The mass-to-charge ratio of the particles is then calculated from the motion of the ions as they transit through electromagnetic fields.

**[0045]** For example, matrix-associated laser desorption/ionization time-of-flight mass spectrometry (MALDI-TOF MS) or surface-enhanced laser desorption/ionization time-of-flight mass spectrometry (SELDI-TOF MS) which involves the application of a sample, such as urine or serum, to a protein-binding chip (Wright, G.L., Jr., et al. (2002) Expert Rev Mol Diagn 2:549; Li, J., et al. (2002) Clin Chem 48:1296; Laronga, C., et al. (2003) Dis Markers 19:229; Petricoin, E.F., et al. (2002) 359:572; Adam, B.L., et al. (2002) Cancer Res 62:3609; Tolson, J., et al. (2004) Lab Invest 84:845; Xiao, Z., et al. (2001) Cancer Res 61:6029) can be used to determine the level of FRA.

**[0046]** Furthermore, *in vivo* techniques for determination of the level of a marker (e.g., FRA) include introducing into a subject a labeled antibody directed against the marker, which binds to the marker to enable its detection. The presence, level, or location of the detectable marker in a subject may be determined using standard imaging techniques (e.g., PET).

**[0047]** As used herein, a subject who is “afflicted with” or “having lung cancer” is one who is clinically diagnosed with lung cancer at any stage by a qualified clinician, or one who exhibits one or more signs or symptoms of such a cancer and is subsequently clinically diagnosed with such a cancer by a qualified clinician. A non-human subject that serves as an animal model of folate receptor-alpha-expressing lung cancer may also fall within the scope of a subject “afflicted with folate receptor-alpha-expressing lung cancer.”

**[0048]** As used herein, a “folate receptor-alpha-expressing lung cancer” includes any type of lung cancer characterized in that the cancer cells express or present on their surface folate receptor alpha. A lung cancer may have been, but is not required to have been, clinically diagnosed as expressing FRA to be encompassed by the term “folate receptor-alpha-expressing lung cancer” as used herein. The phrase “folate receptor-alpha-expressing lung cancer” specifically includes FRA-expressing non-small cell lung cancer (NSCLC) and FRA-expressing non-small cell lung adenocarcinoma.

**[0049]** The term “sample” or “biological sample” as used herein refers to a collection of similar fluids, cells, or tissues isolated from a subject, as well as fluids, cells, or tissues present within a subject. The sample assessed for the expression level of FRA may be derived from urine, blood, serum, plasma, pleural effusion, sputum, bronchial washings, circulating cells, circulating tumor cells, cells that are not tissue associated (i.e., free cells), tissues (e.g., pleural tissue, surgically resected tumor tissue, biopsies, including fine needle aspiration), histological preparations, and the like.

**[0050]** In some embodiments, only a portion of the sample is subjected to an assay for determining the level of a marker, or various portions of the sample are subjected to various assays for determining the level of a marker. Also, in many embodiments, the sample may be pre-treated by physical or chemical means prior to the assay. For example, samples, may be subjected to centrifugation, dilution and/or treatment with a solubilizing substance (e.g., guanidine treatment) prior to assaying the samples for a marker. Such techniques serve to enhance the accuracy, reliability and reproducibility of the assays.

**[0051]** The term “reference expression level” when used to describe the level of expression of a marker (e.g., FRA) refers to an accepted or pre-determined level of a marker which is used to compare with the level of the marker in a sample derived from a subject. In one embodiment, the reference expression level of FRA is predetermined using a population of patients afflicted with the FRA-expressing lung cancer that are treated with a FRA targeting agent (e.g., an antibody that immunospecifically binds FRA) relative to a population of patients afflicted with the FRA-expressing lung cancer that receive placebo in lieu of the FRA targeting agent. The patient populations may have received standard-of-care therapy for such FRA-expressing lung cancer in addition to the antibody or placebo.

**[0052]** As used herein, the term “contacting the sample” with a specific binding agent, e.g., an antibody, includes exposing the sample, or any portion thereof with the agent, such that at least a portion of the sample comes into contact with the agent. The sample or portion thereof may be altered in some way, such as by subjecting it to physical or chemical treatments (e.g., dilution or guanidine treatment), prior to the act of contacting it with the agent.

**[0053]** The term "inhibit" or "inhibition of" means to reduce by a measurable amount, or to prevent entirely.

**[0054]** The term "deplete," in the context of the effect of an anti-FRA therapeutic agent on folate receptor alpha-expressing cells, refers to a reduction in the number of, or elimination of, the folate receptor alpha-expressing cells.

**[0055]** The term "functional," in the context of an antibody to be used in accordance with the methods described herein, indicates that the antibody is (1) capable of binding to antigen and/or (2) depletes or inhibits the proliferation of antigen-expressing cells.

**[0056]** The term "treatment" or "treat" or "positive therapeutic response" refers to slowing, stopping, or reversing the progression of a folate receptor alpha-expressing lung cancer in a patient, as evidenced by a decrease or elimination of a clinical or diagnostic symptom of the disease, by administration of a FRA targeting agent to the subject after the onset of a clinical or diagnostic symptom of the folate receptor alpha-expressing lung cancer at any clinical stage. Treatment can include,

for example, a decrease in the severity of a symptom, the number of symptoms, or frequency of relapse, depletion of FRA-expressing lung cancer cells, inhibition of the growth of FRA-expressing lung cancer cells, or a statistically significant and/or clinically relevant improvement in a specific clinical outcome (e.g., progression-free survival, overall survival).

**[0057]** The phrase "responsive to treatment with a FRA targeting agent" is intended to mean that the candidate subject (i.e., an individual with FRA-expressing lung cancer), following administration of the FRA targeting agent, would have a positive therapeutic response with respect to the lung cancer.

**[0058]** The term "pharmaceutically acceptable" refers to those properties and/or substances which are acceptable to the patient from a pharmacological/toxicological point of view and to the manufacturing pharmaceutical chemist from a physical/chemical point of view regarding composition, formulation, stability, patient acceptance and bioavailability and includes properties and/or substances approved by a regulatory agency of the Federal or a state government or listed in the U.S. Pharmacopeia or other generally recognized pharmacopeia for use in animals, and more particularly in humans. The term "pharmaceutically compatible ingredient" refers to a pharmaceutically acceptable diluent, adjuvant, excipient, or vehicle with which an anti-folate receptor alpha antibody is administered. "Pharmaceutically acceptable carrier" refers to a medium that does not interfere with the effectiveness of the biological activity of the active ingredient(s) and is not toxic to the host to which it is administered.

**[0059]** The terms "effective amount" and "therapeutically effective amount" are used interchangeably herein and, in the context of the administration of a pharmaceutical agent, refer to the amount of the agent (e.g., a FRA targeting agent) that is sufficient to inhibit the occurrence or ameliorate one or more clinical or diagnostic symptoms of a folate receptor alpha-expressing lung cancer in a patient. A therapeutically effective amount of an agent may vary according to factors such as the disease state, age, sex, and weight of the individual, and the ability of the agent to elicit a desired response in the individual. Such results may include, but are not limited to, the treatment of a folate-receptor alpha-expressing lung cancer, as

determined by any means suitable in the art. An effective amount of an agent is administered according to the methods described herein in an "effective regimen." The term "effective regimen" refers to a combination of amount of the agent and dosage frequency adequate to accomplish treatment of a folate receptor alpha-expressing lung cancer.

**[0060]** The terms "patient" and "subject" are used interchangeably to refer to humans and other non-human animals, including veterinary subjects, that receive diagnostic, prophylactic or therapeutic treatment. The term "non-human animal" includes all vertebrates, e.g., mammals and non-mammals, such as non-human primates, mice, rabbits, sheep, dog, cat, horse, cow, chickens, amphibians, and reptiles. In a preferred embodiment, the subject is a human.

**[0061]** Therapeutic agents are typically substantially pure from undesired contaminants. This means that an agent is typically at least about 50% w/w (weight/weight) pure as well as substantially free from interfering proteins and contaminants. Sometimes the agents are at least about 80% w/w and, more preferably at least 90 or about 95% w/w pure. However, using conventional protein purification techniques, homogeneous peptides of at least 99% purity w/w can be obtained.

#### **Methods for predicting a likelihood of responsiveness to treatment with a FRA targeting agent**

**[0062]** Provided herein are methods for predicting a likelihood of responsiveness to treatment with a FRA targeting agent (e.g., an antibody that immunospecifically binds FRA) in a patient having a folate receptor alpha (FRA)-expressing lung cancer. In some embodiments of the methods for predicting a likelihood of responsiveness to treatment with a FRA targeting agent described herein, the FRA-expressing lung cancer is non-small cell lung cancer (NSCLC). In some embodiments, the NSCLC is adenocarcinoma.

**[0063]** The described methods for predicting a likelihood of responsiveness to treatment with a FRA targeting agent in a patient having a folate receptor alpha (FRA)-expressing lung cancer involve determining the patient's FRA expression level in a biological sample of the patient.

**[0064]** Determination of the level of expression of FRA in a biological sample of a patient may be performed upon diagnosis, upon surgical resection, upon initiation of first-line therapy, upon completion of first-line therapy, upon symptomatic progression, serologic progression, and/or radiologic progression of the cancer, upon initiation of second or greater-line therapy, and/or upon completion of such therapy.

**[0065]** In the disclosed methods for predicting a likelihood of responsiveness to treatment with a FRA targeting agent in a patient having a folate receptor alpha (FRA)-expressing lung cancer, the FRA expression level may be determined by any means known in the art including known methods of RNA or protein quantification. Such methods include, but are not limited to, use of antibody to detect protein expression, nucleic acid hybridization, quantitative RT-PCR, immunoprecipitation, equilibrium dialysis, immunodiffusion, immunohistochemistry, fluorescence-activated cell sorting (FACS), fluorimetry, hybridization assays, electrophoresis, capillary electrophoresis, high performance liquid chromatography (HPLC), thin layer chromatography (TLC), hyperdiffusion chromatography, fluid or gel precipitation reactions, absorption spectroscopy, colorimetric assays, spectrophotometric assays, flow cytometry, immunodiffusion (single or double), solution phase assay, immunoprecipitation, equilibrium dialysis, immunodiffusion, solution phase assay, immunoelectrophoresis, Western blotting, radioimmunoassay (RIA), enzyme-linked immunosorbent assays (ELISAs), immunofluorescent assays, and electrochemiluminescence immunoassay, and the like. In a preferred embodiment, the level is determined using antibody-based techniques, as described in more detail herein. In a preferred embodiment, immunohistochemical analysis (e.g., of pleural tissue) is employed. The step of determining the expression level of FRA may be performed *ex vivo* or *in vivo*.

**[0066]** For *ex vivo* assessments, the biological sample used in determining the level of FRA expression may be urine, whole blood, serum, plasma, pleural effusion, sputum, bronchial washings, circulating cells, circulating tumor cells, cells that are not tissue associated (i.e., free cells), tissues (e.g., pleural tissue, surgically resected tumor tissue, biopsies, including fine needle aspiration), histological

preparations, and the like. The tissue sample on which the assay is performed can be fixed or frozen to permit histological sectioning. Preferably, the excised tissue samples are fixed in aldehyde fixatives such as formaldehyde, paraformaldehyde, glutaraldehyde; or heavy metal fixatives such as mercuric chloride. More preferably, the excised tissue samples are fixed in formalin and embedded in paraffin wax prior to incubation with the antibody. Optionally, FFPE specimens can be treated with citrate, EDTA, enzymatic digestion or heat to increase accessibility of epitopes. Alternatively, a protein fraction can be isolated from cells from known or suspected lung cancer and analyzed by ELISA, Western blotting, immunoprecipitation or the like. In another variation, cells can be analyzed for expression of folate receptor alpha by FACS analysis. In a further variation, mRNA can be extracted from cells from known or suspected lung cancer. The mRNA or a nucleic acid derived therefrom, such as a cDNA can then be analyzed by hybridization to a nucleic probe binding to DNA encoding folate receptor alpha.

**[0067]** For example, the step of determining FRA expression level may involve determining the level of FRA expression in a biological sample of the lung cancer tissue obtained from the subject. FRA expression levels may be determined by an immunoassay in which a sample containing cells known or suspected to be from a cancer (e.g., lung cancer) is contacted with an anti-FRA antibody or antigen-binding fragment. After contact, the presence or absence of a binding event of the antibody or antigen-binding fragment to the cells in the specimen is determined. The binding is related to the presence or absence of the antigen expressed on cancerous cells in this specimen. Generally, the sample is contacted with a labeled specific binding partner of the anti-FRA antibody or antigen-binding fragment capable of producing a detectable signal. Alternatively, the anti-FRA antibody or fragment itself can be labeled. Examples of types of labels include enzyme labels, radioisotopic labels, nonradioactive labels, fluorescent labels, toxin labels and chemiluminescent labels. Many such labels are readily known to those skilled in the art. For example, suitable labels include, but should not be considered limited to, radiolabels, fluorescent labels (such as DyLight® 649), epitope tags, biotin, chromophore labels, ECL labels, or enzymes. More specifically, the described labels include ruthenium,

<sup>111</sup>In-DOTA, <sup>111</sup>In- diethylenetriaminepentaacetic acid (DTPA), horseradish peroxidase, alkaline phosphatase and beta-galactosidase, poly-histidine (HIS tag), acridine dyes, cyanine dyes, fluorone dyes, oxazin dyes, phenanthridine dyes, rhodamine dyes, Alexafluor® dyes, and the like. Detection of a signal from the label indicates the presence of the antibody or fragment specifically bound to folate receptor alpha in the sample.

**[0068]** As mentioned above, in some embodiments, FRA expression level may be determined by an immunoassay in which a sample containing cells known or suspected to be from a cancer (e.g., lung cancer) is contacted with an anti-FRA antibody or antigen-binding fragment. In some embodiments, the FRA is not bound to a cell in the sample. Methods for determining the expression level of FRA in a sample derived from the subject are disclosed, for example, in U.S. Publ. No. 20130017195, incorporated herein by reference. Methods for determining the expression level of FRA which is not bound to a cell in a sample derived from the subject are disclosed, for example, in U.S. Publ. No. 20120207771, incorporated herein by reference. The sample employed in the determination of the expression level of FRA may be derived from urine, blood, serum, plasma, pleural effusion, sputum, bronchial washings, circulating cells, circulating tumor cells, cells that are not tissue associated (i.e., free cells), tissues (e.g., pleural tissue, surgically resected tumor tissue, biopsies, including fine needle aspiration), histological preparations, and the like, for example. In preferred embodiments, the sample is tissue, urine, or serum.

**[0069]** In various aspects, the expression level of FRA is determined by contacting the sample with an antibody that binds FRA. For example, the anti-FRA antibody may be selected from the group consisting of:

- (a) an antibody that binds the same epitope as the MORAb-003 antibody;
- (b) an antibody comprising SEQ ID NO:1 (GFTFSGYGLS) as CDRH1, SEQ ID NO:2 (MISSGGSYTYYADSVKG) as CDRH2, SEQ ID NO:3 (HGDDPAWFAY) as CDRH3, SEQ ID NO:4 (SVSSSISSNNLH) as CDRL1, SEQ ID NO:5 (GTSNLAS) as CDRL2 and SEQ ID NO:6 (QQWSSYPYMYT) as CDRL3; );

(c) an antibody comprising a mature light chain variable region comprising the amino acid sequence of SEQ ID NO:7:

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1  DIQLTQSPSS LSASVGDRVT ITCSVSSSIS SNNLHWYQQK PGKAPKWY
51  GTSNLASGVP SRFSGSGSGT DYTFTISSLQ PEDIATYYCQ QWSSYPYMYT
101 FGQGTKVEIK RTVAAPSVFI FPPSDEQLKS GTASVVCLLN NFYPREAKVQ
151 WKVDNALQSG NSQESVTEQD SKDSTYSLSS TLTLSKADYE KHKVYACEVT
201 HQGLSSPVTK SFNRGEC

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and a mature heavy chain variable region comprising the amino acid SEQ ID NO: 8:

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1  EVQLVESGGG VVQPGRSLRL SCSASGFTFS GYGLSWVRQA PGKGLEWVAM
51  ISSGGSYTYY ADSVKGRFAI SRDNAKNTLF LQMDSLRPED TGVYFCARHG
101 DDPAWFAYWG QGTPVTVSSA STKGPSVFP APSSKSTSGG TAALGCLVKD
151 YFPEPVTVSW NSGALTSGVH TFPAVLQSSG LYSLSSVVTV PSSSLGTQTY
201 ICNVNHKPSN TKVDKKVEPK SCDKTHTCPP CPAPELLGGP SVFLFPPKPK
251 DTLMISRTPE VTCVVVDVSH EDPEVKFNWY VDGVEVHNAK TKPREEQYNS
301 TYRVVSVLTV LHQDWLNGKE YKCKVSNKAL PAPIEKTISK AKGQPREPQV
351 YTLPPSRDEL TKNQVSLTCL VKGFYPSDIA VEWESNGOPE NNYKTTPPVL
401 DSDGSFFLYS KLTVDKSRWQ QGNVFSCSVM HEALHNHYTQ KSLSLSPGK

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(CDRs underlined);

(d) the MORAb-003 antibody (USAN name: farletuzumab), as described in U.S. Publ. No. 20090274697 and U.S. Patent No. 8,124,083, the entire contents of each of which are incorporated herein by reference;

- (e) the 548908 antibody (Novus; catalog no. MAB5646);
- (f) an antibody that binds the same epitope as the 548908 antibody;
- (g) the 6D398 antibody (USBiological Life Sciences);
- (h) an antibody that binds the same epitope as the 6D398 antibody;
- (i) the BN3.2 antibody (Leica Biosystems);
- (j) an antibody that binds the same epitope as the BN3.2 antibody;
- (k) an antibody that binds the same epitope as the 26B3 antibody;
- (l) an antibody comprising SEQ ID NO: 14 (GYFMN) as CDRH1, SEQ ID NO: 15 (RIFPYNGDTFYNQKF~~K~~G) as CDRH2, SEQ ID NO: 16 (GTHYFDY) as CDRH3, SEQ ID NO: 17 (RTSENIFSYLA) as CDRL1, SEQ ID NO: 18 (NAKTLAE) as CDRL2 and SEQ ID NO: 19 (QHHYAFPWT) as CDRL3;
- (m) the 26B3 antibody;

- (n) an antibody that binds the same epitope as the 19D4 antibody;
- (o) an antibody comprising SEQ ID NO: 20 (HPYMH) as CDRH1, SEQ ID NO: 21 (RIDPANGNTKYDPKFQG) as CDRH2, SEQ ID NO: 22 (EEVADYTM DY) as CDRH3, SEQ ID NO: 23 (RASESVDTYGNNFIH) as CDRL1, SEQ ID NO: 24 (LASNLES) as CDRL2 and SEQ ID NO: 25 (QQNNGDPWT) as CDRL3;
- (p) the 19D4 antibody (see U.S. Patent 8,475,795);
- (q) an antibody that binds the same epitope as the 9F3 antibody;
- (r) an antibody comprising SEQ ID NO: 26 (SGYYWN) as CDRH1, SEQ ID NO: 27 (YIKSDGSNNYNPSLKN) as CDRH2, SEQ ID NO: 28 (EWKAMDY) as CDRH3, SEQ ID NO: 29 (RASSTVSYSYLH) as CDRL1, SEQ ID NO: 30 (GTSNLAS) as CDRL2 and SEQ ID NO: 31 (QQYSGYPLT) as CDRL3;
- (s) the 9F3 antibody (see U.S. Patent 8,475,795);
- (t) an antibody that binds the same epitope as the 24F12 antibody;
- (u) an antibody comprising SEQ ID NO: 32 (SYAMS) as CDRH1, SEQ ID NO: 33 (EIGSGGSYTYYPDTVTG) as CDRH2, SEQ ID NO: 34 (ETTAGYFDY) as CDRH3, SEQ ID NO: 35 (SASQGINNFLN) as CDRL1, SEQ ID NO: 36 (YTSSLHS) as CDRL2 and SEQ ID NO: 37 (QHFSKLPWT) as CDRL3;
- (v) the 24F12 antibody (see U.S. Patent 8,475,795);
- (w) an antibody that comprises a variable region light chain selected from the group consisting of:

- (i) LK26HuVK as set forth in SEQ ID NO: 38:  
Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
Asp Arg Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Asn  
Asn Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu  
Ile Tyr Gly Thr Ser Asn Leu Ala Ser Gly Val Pro Ser Arg Phe Ser  
Gly Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu Gln  
Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Tyr Pro  
Tyr Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys,

- (ii) LK26HuVKY as set forth in SEQ ID NO: 39:  
Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly

Asp Arg Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Asn Asn Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile Tyr Gly Thr Ser Asn Leu Ala Ser Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Tyr Thr Phe Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Tyr Pro Tyr Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys,

(iii) LK26HuVKPW as set forth in SEQ ID NO: 40:

Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Asn Asn Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Pro Trp Ile Tyr Gly Thr Ser Asn Leu Ala Ser Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Tyr Pro Tyr Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys,

and

(iv) LK26HuVKPW,Y as set forth in SEQ ID NO: 41:

Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Asn Asn Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Pro Trp Ile Tyr Gly Thr Ser Asn Leu Ala Ser Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Tyr Thr Phe Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Tyr Pro Tyr Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys;

(x) an antibody that comprises a variable region heavy chain selected from the group consisting of:

(i) LK26HuVH as set forth in SEQ ID NO: 42:

Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg Pro Ser Gln Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly Phe Thr Phe Ser Gly Tyr Gly Leu Ser Trp Val Arg Gln Pro Pro Gly Arg Gly Leu Glu Trp Val Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val Lys Gly Arg Val Thr Met Leu Arg Asp Thr Ser Lys Asn Gln Phe Ser Leu Arg Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly

Ser Leu Val Thr Val Ser Ser,

(ii) LK26HuVH FAIS,N as set forth in SEQ ID NO: 43:

Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg Pro Ser Gln Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly Phe Thr Phe Ser Gly Tyr Gly Leu Ser Trp Val Arg Gln Pro Pro Gly Arg Gly Leu Glu Trp Val Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ser Lys Asn Gln Phe Ser Leu Arg Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly Ser Leu Val Thr Val Ser Ser,

(iii) LK26HuVH SLF as set forth in SEQ ID NO: 44:

Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg Pro Ser Gln Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly Phe Thr Phe Ser Gly Tyr Gly Leu Ser Trp Val Arg Gln Pro Pro Gly Arg Gly Leu Glu Trp Val Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val Lys Gly Arg Val Thr Met Leu Arg Asp Thr Ser Lys Asn Ser Leu Phe Leu Arg Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser,

(iv) LK26HuVH I,I as set forth in SEQ ID NO: 45:

Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg Pro Ser Gln Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly Phe Thr Phe Ser Gly Tyr Gly Leu Ser Trp Val Arg Gln Pro Pro Gly Arg Gly Leu Glu Trp Val Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val Lys Gly Arg Val Thr Met Leu Arg Asp Thr Ser Lys Asn Gln Phe Ser Leu Arg Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Ile Tyr Ile Cys Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly Ser Leu Val Thr Val Ser Ser,

(v) LK26KOLHuVH as set forth in SEQ ID NO: 46:

Glu Val Gln Leu Val Glu Ser Gly Gly Val Val Gln Pro Gly Arg Ser Leu Arg Leu Ser Cys Ser Ala Ser Gly Phe Thr Phe Ser Gly Tyr Gly Leu Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val

Ala Met Ile Ser Ser Gly Gly Ser Tyr Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Phe Leu Gln Met Asp Ser Leu Arg Pro Glu Asp Thr Gly Val Tyr Phe Cys Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly Thr Pro Val Thr Val Ser Ser;

(y) an antibody that comprises the heavy chain variable region LK26KOLHuVH (SEQ ID NO: 46) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41);

(z) an antibody that comprises the heavy chain variable region LK26HuVH SLF (SEQ ID NO: 44) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41);

(aa) an antibody that comprises the heavy chain variable region LK26HuVH FAIS,N (SEQ ID NO: 43) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41); and

(bb) the murine monoclonal LK26 antibody, the heavy and light chains of which are presented herein as SEQ ID NOs: 11 and 12, respectively:

SEQ ID NO: 11

Gln Val Xaa Leu Gln Xaa Ser Gly Gly Asp Leu Val Lys Pro Gly Gly Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Gly Tyr Gly Leu Ser Trp Val Arg Gln Thr Pro Asp Lys Arg Leu Glu Trp Val Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Phe Leu Gln Met Ser Ser Leu Lys Ser Asp Asp Thr Ala Ile Tyr Ile Cys Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ala (wherein Xaa refers to any amino acid)

SEQ ID NO: 12

Asp Ile Glu Leu Thr Gln Ser Pro Ala Leu Met Ala Ala Ser Pro Gly Glu Lys Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Asn Asn Leu His Trp Tyr Gln Gln Lys Ser Glu Thr Ser Pro Lys Pro Trp Ile Tyr Gly Thr Ser Asn Leu Ala Ser Gly Val Pro Leu Arg Phe Arg Gly Phe Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser Ser Met Glu Ala Glu Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Tyr Pro

Tyr Met Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys

as described in European Patent Application No. 86104170.5 (Rettig), the entire contents of which are incorporated herein by reference. In certain embodiments, the anti-FRA antibody includes (i) the heavy chain variable region LK26KOLHuVH (SEQ ID NO: 46) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41); the heavy chain variable region LK26HuVH SLF (SEQ ID NO: 44) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41); or the heavy chain variable region LK26HuVH FAIS,N (SEQ ID NO: 43) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41). Chinese hamster ovary (CHO) cells producing MORAb-003 have been deposited with the ATCC (10801 University Boulevard, Manassas, VA 20110) on April 24, 2006 and assigned accession no. PTA-7552.

**[0070]** Other useful antibodies that immunospecifically bind to folate receptor alpha comprise mature light and heavy chain variable regions having at least 90% and preferably at least 95% or 99% sequence identity to SEQ ID NO: 7 and SEQ ID NO: 8, respectively. Other useful antibodies that immunospecifically bind to folate receptor alpha or derivatives thereof can competitively inhibit binding of farletuzumab to folate receptor alpha, as determined, for example, by immunoassay. Competitive inhibition means that an antibody when present in at least a two-fold and preferably five-fold excess inhibits binding of farletuzumab to folate receptor alpha by at least 50%, more typically at least 60%, yet more typically at least 70%, and most typically at least 75%, at least 80%, at least 85%, at least 90%, or at least 95%.

**[0071]** The antibody that immunospecifically binds to folate receptor alpha may also be a derivative of an anti-folate receptor alpha antibody disclosed above. Typical modifications include, e.g., glycosylation, deglycosylation, acetylation, pegylation, phosphorylation, amidation, derivatization by known protecting/blocking groups, proteolytic cleavage, linkage to a cellular ligand or other protein, and the like. Additionally, the derivative may contain one or more non-classical amino acids.

**[0072]** In certain embodiments, the anti-FRA antibody is selected from the group consisting of a murine antibody, a human antibody, a humanized antibody, a bispecific antibody, a chimeric antibody, a Fab, Fab'2, ScFv, SMIP, affibody, avimer, versabody, nanobody, biobody, and a domain antibody. Alternatively, or in combination, the anti-FRA antibody is labeled, for example, with a label selected from the group consisting of a radio-label, a biotin-label, a chromophore-label, a fluorophore-label, or an enzyme-label.

**[0073]** In a further aspect, the level of folate receptor alpha (FRA) expression in a sample derived from the subject is assessed by a two-antibody sandwich assay. In some embodiments of the sandwich assay, the sample is contacted with (a) 9F3 antibody immobilized to a solid support and labeled 24F12 antibody, (b) 26B3 antibody immobilized to a solid support and labeled 19D4 antibody, and (c) 9F3 antibody immobilized to a solid support and labeled 26B3 antibody. For example, the sample may be urine, whole blood, serum, plasma, pleural effusion, sputum, bronchial washings, circulating cells, circulating tumor cells, cells that are not tissue associated (i.e., free cells), tissues (e.g., pleural tissue, surgically resected tumor tissue, biopsies, including fine needle aspiration), histological preparations, and the like.

**[0074]** In certain embodiments, the sample is treated with guanidine prior to determining the level of FRA expression in the sample. Alternatively or in combination, the sample is diluted prior to determining the level of FRA expression in the sample. Alternatively, or in combination, the sample is centrifuged, vortexed, or both, prior to determining the level of FRA expression in the sample.

**[0075]** In another variation, FRA expression level in known or suspected lung cancer can be detected *in vivo* by administering a labeled anti-FRA antibody or antigen-binding fragment thereof to a patient and detecting the antibody or fragment by *in vivo* imaging. Any of the antibodies described above may likewise be employed in the *in vivo* imaging analysis.

**[0076]** The level of FRA in a lung tissue sample can (but need not) be determined with respect to one or more standards. The standards can be historically or contemporaneously determined. The standard can be, for example, a FRA-

expressing lung tissue sample known to be cancerous from a different subject, a FRA-expressing lung tissue sample known not to be cancerous from a different subject, a tissue from either the patient or other subject known not to express FRA, or a FRA-expressing lung cancer cell line.

**[0077]** The presence of detectable signal from binding of an anti-FRA antibody or fragment to FRA relative to a standard (if used) indicates the presence of FRA in the tissue sample, and the level of detectable binding provides an indication of the level of expression of FRA. In assays performed on tissue sections, the level of expression can be expressed as a percentage of the surface area of the sample showing detectable expression of FRA. Alternatively, or additionally, the level (intensity) of expression can be used as a measure of the total expression in the sample or of the cells expressing FRA in the sample. The intensity of expression can be determined, for example, via digital imaging or manual microscopic assessment of tissue sections using methods as previously described (Potts, Drug Discov Today, 2009; 14(19-20):935-41; O'Shannessy et al., Oncotarget, 2012; 3(4):414-25; U.S. Patent 8,475,795; manufacturer's instructions, Catalog no. IPI4006K G10 (Biocare Medical; Concord, CA). The intensity of FRA expression may be used to determine a FRAMSCOR or HBSCOR as described herein. In particular, the FRAMSCOR (M-score) is calculated as a weighted average, assuming a 0, 1+, 2+, 3+ scoring system as follows (see Figure 2):

$$x = \% \text{ tumor stained at } 1+$$

$$y = \% \text{ tumor stained at } 2+$$

$$z = \% \text{ tumor stained at } 3+$$

Then

$$M = \frac{x + 2y + 3z}{6}$$

For purposes of example, if a patient's tumor has 20% FRA staining at +1, 10% FRA staining at +2 and 20% FRA staining at +3, then the M score = 16.6. The HBSCOR (H-Score) reports the mean optical density value for biomarker staining (in this case, staining of FRA) computed from all the cells in a target tissue compartment. It utilizes proprietary tissue recognition features to determine the tissue

compartment via a linear score and a continuous extension of the H-score with no cell classification. The H-score is a standard scoring method that is commonly used by pathologists and those skilled in the art to score biomarker expression in tissues, which is basically the sum of the intensity scores at all intensity levels (1+, + 2 x 2+, + 3x 3+). The HBSCOR is derived from the sum of cell measurements (optical density) divided by the total number of cells. HBSCOR in turn reports the value for biomarker staining computed from all the cells in a target tissue compartment. This calculation is quantified using the following formula:

$$\text{HBSCOR} = \frac{\sum \text{Cell Measurement}}{\text{Number of Cells}}$$

[0078] Once the patient's FRA expression level is determined, it is compared to a reference FRA expression level. In a preferred embodiment, the patient's FRA expression level is presented in terms of a FRAMSCOR (*i.e.*, M-score) or HBSCOR (*i.e.*, H-score) for comparison to a reference FRA expression level. In a preferred embodiment, the reference FRA expression level is predetermined. For example, a reference data set may be established using samples from unrelated patients with low, moderate and high FRA expression levels. This data set represents a standard by which relative FRA expression levels are compared among patients and quantified using the manual and digital analysis FRAMSCOR and HBSCOR methods. In some embodiments, the reference FRA expression level is determined by comparison of a patient population afflicted with the FRA-expressing lung cancer that is administered the FRA targeting agent to a patient population afflicted with the FRA-expressing lung cancer that is administered placebo. The patient populations afflicted with the FRA-expressing lung cancer may also have received standard-of-care chemotherapy. The FRA expression level for each patient in the respective populations afflicted with the FRA-expressing lung cancer is determined in accordance with the methods described above. Clinical outcomes (e.g., progression-free survival or overall survival) for the patient

populations are monitored. Clinical outcomes for the patient populations relative to FRA expression levels are then compared as described in the examples provided below. The reference FRA expression level corresponds to the FRA expression level above which the patient population afflicted with the FRA-expressing lung cancer that is administered the FRA targeting agent (e.g., an antibody that immunospecifically binds FRA) demonstrates a statistically significant improvement in at least one clinical outcome relative to the patient population afflicted with the FRA-expressing lung cancer that is administered placebo. A patient FRA expression level that equals or exceeds the reference FRA expression level is indicative that the patient will benefit from treatment with the FRA targeting agent.

**[0079]** Methods of Treatment

**[0080]** Also provided herein are methods of treating a patient having a folate receptor alpha (FRA)-expressing lung cancer. In some embodiments of the methods for treating a patient having FRA-expressing lung cancer, the cancer is NSCLC. In some embodiments, the NSCLC is adenocarcinoma. The disclosed methods for treating a FRA-expressing lung cancer in a patient include methods in which an antibody that immunospecifically binds folate receptor alpha (FRA) is administered to a patient whose FRA expression level equals or exceeds a reference FRA expression level.

**[0081]** In accordance with the methods of treating a patient with folate receptor alpha (FRA)-expressing lung cancer described herein, the patient's FRA expression level in a biological sample of the patient is quantified and compared to a reference FRA expression level as described above. If the patient's FRA expression level equals or exceeds the reference FRA expression level, then the patient is administered a FRA targeting agent (e.g., an antibody that immunospecifically binds FRA).

**[0082]** In some embodiments of the methods of treatment described herein, the FRA targeting agent is vintafolide. In some embodiments, the FRA targeting agent is an antibody that immunospecifically binds FRA, such as but not limited to an antibody that immunospecifically binds to folate receptor alpha expressed on lung

cancer cells; antigen-binding fragments of such an antibody; derivatives; and variants thereof. In a preferred embodiment, the antibody that immunospecifically binds to folate receptor alpha is an antibody selected from the group consisting of:

- (a) an antibody comprising SEQ ID NO:1 as CDRH1, SEQ ID NO:2 as CDRH2, SEQ ID NO:3 as CDRH3, SEQ ID NO:4 as CDRL1, SEQ ID NO:5 as CDRL2 and SEQ ID NO:6 as CDRL3;
- (b) an antibody comprising a mature light chain variable region comprising the amino acid sequence of SEQ ID NO:7 and/or a mature heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 8;
- (c) farletuzumab;
- (d) an antibody that specifically binds to folate receptor alpha comprising a mature light and heavy chain variable regions having at least 90% and preferably at least 95% or 99% sequence identity to SEQ ID NO: 7 and SEQ ID NO: 8, respectively;
- (e) an antibody or derivative thereof can competitively inhibit binding of farletuzumab to folate receptor alpha, as determined, for example, by immunoassay.

A derivative of an antibody that immunospecifically binds FRA can also be used in the practice of the present methods. Typical modifications include, e.g., glycosylation, deglycosylation, acetylation, pegylation, phosphorylation, amidation, derivatization by known protecting/blocking groups, proteolytic cleavage, linkage to a cellular ligand or other protein, and the like. Additionally, the derivative may contain one or more non-classical amino acids. In some embodiments, that antibody that immunospecifically binds FRA is conjugated to a toxin, such as but not limited to a microtubule inhibitor, a DNA damaging agent (e.g., a radionuclide), a DNA repair inhibitor, or a signal transduction inhibitor. Linkers for antibody conjugation and methods for conjugating antibodies are known in the art. An exemplary antibody-drug conjugate that can be employed as a FRA-targeting agent in accordance with the methods described herein is IMGN853.

**[0083]** The present methods can be combined with other means of treatment such as surgery (e.g., debulking surgery), radiation, targeted therapy,

chemotherapy, immunotherapy, use of growth factor inhibitors, or anti-angiogenesis factors. The FRA targeting agent can be administered concurrently to a patient undergoing surgery, chemotherapy or radiation therapy treatments. Alternatively, a patient can undergo surgery, chemotherapy or radiation therapy prior or subsequent to administration of the FRA targeting agent by at least an hour and up to several months, for example at least an hour, five hours, 12 hours, a day, a week, a month, or three months, prior or subsequent to administration of the FRA targeting agent. For example, some embodiments of the methods of treatment provided herein further involve administration of a therapeutically effective amount of a platinum-containing compound, an antifolate, and/or a taxane to the subject in addition to the FRA targeting agent. Exemplary platinum-containing compounds are cisplatin or carboplatin. Examples of taxanes for use in the methods of treatment include but are not limited to paclitaxel, docetaxel, and semi-synthetic, synthetic, and/or modified versions and formulations thereof, including but not limited to nab-paclitaxel (Abraxane®), cabazitaxel (Jevtana®), DJ-927 (Tesetaxel®), paclitaxel poliglumex (Opaxio®), XRP9881 (Larotaxel®), EndoTAG + paclitaxel (EndoTAG®-1), Polymeric-micellar paclitaxel (Genexol-PM®), DHA-paclitaxel (Taxoprexin®), BMS-184476. An exemplary antifolate is pemetrexed. The platinum-containing compound may be administered to the patient once every week, once every two weeks, once every three weeks, or once every four weeks. The taxane may be administered to the patient once every week, once every two weeks, once every three weeks, or once every four weeks. The antifolate may be administered to the patient once every week, once every two weeks, once every three weeks, or once every four weeks. In embodiments in which both a platinum-containing compound and a taxane or antifolate are administered to the patient as part of the treatment regimen, the taxane or antifolate may be administered before, after, or simultaneously with the platinum-containing compound.

**[0084]** In some embodiments of the methods of treatment described herein, the patient may have received surgical resection of the lung cancer, prior platinum-based therapy, prior taxane-based therapy, and/or prior platinum and taxane-based therapy for treatment of the cancer prior to quantifying the patient's FRA expression

level. In some embodiments of the methods in which the patient received surgical resection of the cancer, prior platinum-based therapy, prior taxane-based therapy, and/or prior platinum and taxane-based therapy for treatment of the cancer prior to determining the patient's FRA expression level, the patient may have exhibited symptomatic progression, serologic progression, and/or radiologic progression of the cancer prior to the step of determining the patient's FRA expression level.

**[0085]** Administration of the therapeutic agents (including the FRA targeting agent, the taxane, the antifolate, and/or the platinum-containing compound) in accordance with the methods of treatment described herein may be by any means known in the art.

**[0086]** Various delivery systems can be used to administer the therapeutic agents (including the FRA targeting agent, the taxane, the antifolate, and/or the platinum-containing compound) including intradermal, intramuscular, intraperitoneal, intravenous, subcutaneous, intranasal, epidural, and oral routes. The agents can be administered, for example by infusion or bolus injection, by absorption through epithelial or mucocutaneous linings (e.g., oral mucosa, rectal and intestinal mucosa, and the like). Administration can be systemic or local.

**[0087]** The therapeutic agents can be administered by injection, by means of a catheter, by means of a suppository, or by means of an implant, the implant being of a porous, non-porous, or gelatinous material, including a membrane, such as a sialastic membrane, or a fiber. The therapeutic agents and pharmaceutical compositions thereof for use as described herein may be administered orally in any acceptable dosage form such as capsules, tablets, aqueous suspensions, solutions or the like.

**[0088]** Preferred methods of administration of the therapeutic agents include but are not limited to intravenous injection and intraperitoneal administration.

**[0089]** Alternatively, the therapeutic agents can be delivered in a controlled release system. For example, a pump can be used (see Langer, 1990, *Science* 249:1527-1533; Sefton, 1989, *CRC Crit. Ref. Biomed. Eng.* 14:201; Buchwald et al., 1980, *Surgery* 88:507; Saudek et al., 1989, *N. Engl. J. Med.* 321:574). Alternatively,

polymeric materials can be used (see Medical Applications of Controlled Release (Langer & Wise eds., CRC Press, Boca Raton, Fla., 1974); Controlled Drug Bioavailability, Drug Product Design and Performance (Smolen & Ball eds., Wiley, New York, 1984); Ranger & Peppas, 1983, *Macromol. Sci. Rev. Macromol. Chem.* 23:61. See also Levy et al., 1985, *Science* 228:190; During et al., 1989, *Ann. Neurol.* 25:351; Howard et al., 1989, *J. Neurosurg.* 71:105.) Other controlled release systems are discussed, for example, in Langer, *supra*.

**[0090]** The therapeutic agents can be administered as pharmaceutical compositions comprising a therapeutically or prophylactically effective amount of the therapeutic agent(s) and one or more pharmaceutically acceptable or compatible ingredients. For example, the pharmaceutical composition typically includes one or more pharmaceutical carriers (e.g., sterile liquids, such as water and oils, including those of petroleum, animal, vegetable or synthetic origin, such as peanut oil, soybean oil, mineral oil, sesame oil and the like). Water is a more typical carrier when the pharmaceutical composition is administered intravenously. Saline solutions (e.g., phosphate buffered saline) and aqueous dextrose and glycerol solutions can also be employed as liquid carriers, particularly for injectable solutions. Suitable pharmaceutical excipients include, for example, starch, glucose, lactose, sucrose, gelatin, malt, rice, flour, chalk, silica gel, sodium stearate, glycerol monostearate, talc, sodium chloride, dried skim milk, glycerol, propylene, glycol, water, ethanol, and the like. The composition, if desired, can also contain minor amounts of wetting or emulsifying agents, pH buffering agents (e.g., amino acids) and/or solubilizing or stabilizing agents (e.g., nonionic surfactants such as tween or sugars such as sucrose, trehalose or the like). The preferred formulation of farletuzumab contains farletuzumab, sodium phosphate, sodium chloride (NaCl), and polysorbate-80, pH 7.2. A preferred final formulation of farletuzumab contains 5 mg/mL farletuzumab, 10 mM sodium phosphate, 150 mM NaCl, and 0.01% polysorbate-80, pH 7.2.

**[0091]** The pharmaceutical compositions provided herein can take the form of solutions, suspensions, emulsion, tablets, pills, capsules, powders, sustained-release formulations and the like. Also included are solid form preparations which are intended to be converted, shortly before use, to liquid preparations. The

composition can be formulated as a suppository, with traditional binders and carriers such as triglycerides. Oral formulation can include standard carriers such as pharmaceutical grades of mannitol, lactose, starch, magnesium stearate, sodium saccharine, cellulose, magnesium carbonate, etc. Examples of suitable pharmaceutical carriers are described in "Remington's Pharmaceutical Sciences" by E. W. Martin. Such compositions will contain a therapeutically effective amount of the nucleic acid or protein, typically in purified form, together with a suitable amount of carrier so as to provide the form for proper administration to the patient. The formulations correspond to the mode of administration.

**[0092]** Typically, compositions for intravenous administration are solutions in sterile isotonic aqueous buffer. When necessary, the pharmaceutical can also include a solubilizing agent and a local anesthetic such as lignocaine to ease pain at the site of the injection. Generally, the ingredients are supplied either separately or mixed together in unit dosage form, for example, as a dry lyophilized powder or a concentrate in a hermetically sealed container such as an ampoule or sachette indicating the quantity of active agent. When the pharmaceutical composition is to be administered by infusion, it can be dispensed with an infusion bottle containing sterile pharmaceutical grade water or saline. When the pharmaceutical composition is administered by injection, an ampoule of sterile water for injection or saline can be provided so that the ingredients can be mixed prior to administration.

**[0093]** The amount of the therapeutic agent that is effective in the treatment of lung cancer can be determined by standard clinical techniques. In addition, *in vitro* assays may optionally be employed to help identify optimal dosage ranges. The precise dose to be employed in the formulation also depends on the route of administration, and the stage of the cancer, and should be decided according to the judgment of the practitioner and each patient's circumstances. Effective doses may be extrapolated from dose-response curves derived from *in vitro* or animal model test systems. A dose can be formulated in animal models to achieve a circulating plasma concentration range that includes the IC<sub>50</sub> (i.e., the concentration of the test compound that achieves a half-maximal inhibition of symptoms) as determined in cell culture.

**[0094]** For example, toxicity and therapeutic efficacy of the agents can be determined in cell cultures or experimental animals by standard pharmaceutical procedures for determining the LD<sub>50</sub> (the dose lethal to 50% of the population) and the ED<sub>50</sub> (the dose therapeutically effective in 50% of the population). The dose ratio between toxic and therapeutic effects is the therapeutic index and it can be expressed as the ratio LD<sub>50</sub>/ED<sub>50</sub>. Agents that exhibit large therapeutic indices are preferred. When an agent exhibits toxic side effects, a delivery system that targets the agent to the site of affected tissue can be used to minimize potential damage to non-folate receptor alpha-expressing cells and, thereby, reduce side effects.

**[0095]** In some embodiments, the subject can be administered a therapeutic agent described herein in a daily dose range of about 0.01 µg to about 500 mg per kg of the weight of the subject. Typically, the dosage of the therapeutic agent (e.g., the FRA targeting agent, such as an antibody that immunospecifically binds FRA, preferably farletuzumab) administered to a patient with a folate receptor alpha-expressing lung cancer is about 0.1 mg/kg to about 100 mg/kg of the subject's body weight. More typically, the dosage administered to a subject is about 1.25 mg/kg to about 12.5 mg/kg of the subject's body weight, or even more typically about 2.5 mg/kg to about 10.0 mg/kg of the subject's body weight. In some embodiments, the dosage of the FRA targeting agent (e.g., an antibody that immunospecifically binds FRA, preferably farletuzumab) administered to a subject having folate receptor alpha-expressing lung cancer is about 5.0 mg/kg to about 7.5 mg/kg of the subject's body weight. In some embodiments of the methods of treatment described herein, a loading dose of the FRA targeting agent (e.g., an antibody that immunospecifically binds FRA) of about 7.5 mg/kg to about 12.5 mg/kg, preferably about 10 mg/kg, is administered to the subject. In some embodiments of the methods of treatment described herein, two loading doses of the FRA targeting agent (e.g., an antibody that immunospecifically binds FRA) of about 7.5 mg/kg to about 12.5 mg/kg weekly, preferably about 10 mg/kg, is administered to the subject in the first two weeks of treatment. In some embodiments, the dosage of the taxane administered to a subject having folate receptor alpha-expressing lung cancer is about 50 mg/m<sup>2</sup> to about 250 mg/m<sup>2</sup> of the subject's body weight, preferably about 75 mg/m<sup>2</sup> to about 200 mg/m<sup>2</sup>.

In some embodiments, the dosage of carboplatin administered to a subject having folate receptor alpha-expressing lung cancer is about AUC 3, preferably about AUC 4, more preferably about AUC 5-6, and in some preferred embodiments, about AUC 6. In some embodiments, the dosage of cisplatin administered to a subject having folate receptor alpha-expressing lung cancer is about 50 mg/m<sup>2</sup> to about 250 mg/m<sup>2</sup> of the subject's body weight, preferably about 75 mg/m<sup>2</sup> to about 200 mg/m<sup>2</sup>. In some embodiments, the dosage of antifolate administered to a subject having folate receptor alpha-expressing lung cancer is about 400 to about 600 mg/m<sup>2</sup>. In a preferred embodiment, at least four to six cycles of chemotherapy in combination with administration of the FRA targeting agent are administered to the patient.

**[0096]** For effective treatment, one skilled in the art may recommend a dosage schedule and dosage amount of the therapeutic agent(s) adequate for the subject being treated. It may be preferred that dosing occur one to four or more times daily, once per week, once per every two weeks, once per every three weeks, or once per every four weeks for as long as needed. Typically, the FRA targeting agent is administered to the subject weekly.

**[0097]** The dosing may occur less frequently if the compositions are formulated in sustained delivery vehicles. The dosage schedule may also vary depending on the active drug concentration, which may depend on the needs of the subject.

**[0098]** Kits

**[0099]** Further provided herein are kits for predicting a likelihood of responsiveness to treatment with a FRA targeting agent in a patient having FRA-expressing lung cancer. In some embodiments, the kits contain an anti-FRA antibody, a vessel for containing the antibody when not in use, and instructions for using the anti-FRA antibody for determining the level of FRA expression of a subject. One or more additional containers may enclose elements, such as reagents or buffers, to be used in the marker assay(s). Such kits can also, or alternatively, contain a detection reagent that contains a reporter group suitable for direct or indirect detection of antibody binding.

**[0100]** Also provided herein are kits for treating a FRA-expressing lung cancer in a patient comprising a FRA targeting agent (e.g., vintafolide, an antibody that immunospecifically binds FRA such a farletuzumab, or an antibody-drug conjugate such as IMGN853), a vessel for containing the FRA targeting agent when not in use, and instructions for use of the FRA targeting agent. Farletuzumab is the preferred antibody that immunospecifically binds FRA in the kits. In some embodiments, the kits for treating a subject having FRA-expressing lung cancer also contain an anti-FRA antibody for use in quantifying the level of FRA expression in a biological sample of the patient. This latter anti-FRA antibody may be the same or different than the antibody that immunospecifically binds FRA that is administered therapeutically. In some embodiments, the kits also contain a vessel for containing the anti-FRA antibody when not in use and instructions for using the anti-FRA antibody for determining the expression level of FRA of a subject.

**[0101]** The kits for treating a subject having FRA-expressing lung cancer also may contain additional therapeutic agents (e.g., a platinum-containing compound, a taxane, and/or an antifolate) as described herein. Examples of platinum-containing compounds for inclusion in the kits include, but are not limited to, cisplatin and carboplatin. Examples of taxanes for inclusion in the kits include, but are not limited to, paclitaxel, docetaxel, and semi-synthetic, synthetic, and/or modified versions and formulations thereof, including but not limited to nab-paclitaxel (Abraxane®), cabazitaxel (Jevtana®), DJ-927 (Tesetaxel®), paclitaxel poliglumex (Opaxio®), XRP9881 (Larotaxel®), EndoTAG + paclitaxel (EndoTAG®-1), Polymeric-micellar paclitaxel (Genexol-PM®), DHA-paclitaxel (Taxoprexin®), BMS-184476. An example of an antifolate for inclusion in the kits is pemetrexed. The therapeutic agents can be in any of a variety of forms suitable for distribution in a kit. Forms of the therapeutic agents suitable for distribution in the kits can include a liquid, powder, tablet, suspension and the like formulation for providing the therapeutic agent. The kits can also include a pharmaceutically acceptable diluent (e.g., sterile water) for injection, reconstitution or dilution of the therapeutic agent(s). One or more additional containers may enclose elements, such as reagents or buffers, to be used in the marker assay(s). Such kits can also, or

alternatively, contain a detection reagent that contains a reporter group suitable for direct or indirect detection of antibody binding.

**[0102]** Kits also typically contain a label or instructions for use in the methods described herein. The label or instruction refers to any written or recorded material that is attached to, or otherwise accompanies a kit at any time during its manufacture, transport, sale or use. It can be a notice in the form prescribed by a governmental agency regulating the manufacture, use or sale of pharmaceuticals or biological products, which notice reflects approval by the agency of manufacture, use or sale for human administration. The label or instruction can also encompass advertising leaflets and brochures, packaging materials, instructions, audio or videocassettes, computer discs, as well as writing imprinted directly on the pharmaceutical kits.

**[0103]** The following example is provided to further describe some of the embodiments disclosed herein. The example is intended to illustrate, not to limit, the disclosed embodiments.

**[0104]** EXAMPLE 1

**[0105]** Patients

**[0106]** Eligible subjects must have had newly diagnosed, unresectable, histologically or cytologically confirmed adenocarcinoma of the lung, with FRA expression (defined by 1+ or greater membranous staining) in at least 5% of tumor cells by immunohistochemistry, classified as Stage IV with at least one unidimensionally measurable lesion according to Response Evaluation Criteria in Solid Tumors (RECIST) version 1.1 using computed tomography (CT) or magnetic resonance imaging (MRI) (ClinicalTrials.gov identifier NCT01218516). As no data exists regarding the degree and frequency of FRA positivity that theoretically may be needed for a maximal anti-FRA antibody pharmacologic effect(s), patients whose tumors were at least 5% positive at a +1 intensity membrane FRA expression (using methods typically employed by those skilled in the art) were designated as having

FRA-positive tumors and eligible for study inclusion. Subjects had received no prior chemotherapy, radiation therapy, or surgery with curative intent for their lung cancer.

**[0107]** Treatment

**[0108]** Subjects were randomized in a 1:1 ratio to receive the selected platinum doublet (carboplatin area under the curve (AUC) pharmacokinetic exposure level of 6 and paclitaxel 200 mg/m<sup>2</sup>; carboplatin AUC 5-6 and pemetrexed 500 mg/m<sup>2</sup>; or cisplatin 75 mg/m<sup>2</sup> and pemetrexed 500 mg/m<sup>2</sup>) with either farletuzumab or placebo. Randomization was stratified by Eastern Cooperative Oncology Group (ECOG) performance status (0 or 1) and type of chemotherapy regimen chosen. All randomized subjects were treated with farletuzumab 7.5 mg/kg or placebo in combination with one of the 3 acceptable chemotherapy regimens, administered intravenously (IV) on Day 1 of a 21-day cycle except for the first cycle when farletuzumab or placebo monotherapy was administered on Day 8 as a loading dose. Beginning at Cycle 2, farletuzumab or placebo was administered on Day 1 of all additional cycles in combination with chemotherapy.

**[0109]** All protocol-acceptable platinum doublets were administered once every 3 weeks for at least 4, but no more than 6, cycles. Chemotherapy regimens were not changed once initiated. Chemotherapy doses could be reduced or delayed due to toxicity according to the country-specific, approved package inserts and based on the degree of toxicity experienced by the subject. In the event that chemotherapy was discontinued prior to the 4th cycle or was held for more than 6 weeks, the subject was discontinued from protocol treatment.

**[0110]** Those subjects who experienced clinical benefit from combination therapy could go on to receive farletuzumab or placebo on Day 1 of every 21-day cycle until documented radiographic progression or other protocol-approved measures of disease progression. Once disease progression had occurred, each subject was followed for survival status and use of additional systemic therapy for Stage IV adenocarcinoma of the lung. During this follow-up period, subjects were contacted monthly for the first 9 months, and every 2 months thereafter until death.

**[0111]** Assessments

**[0112]** Radiographic disease evaluations occurred every 2 cycles during combination therapy and every 3 cycles during monotherapy, read locally per independent review. CT scans (or MRIs) were also evaluated independently by central review using RECIST v.1.1. Subjects who discontinued study treatment prior to radiographic progression for any reason were followed radiographically every 9 weeks until documented radiographic progression or other protocol-approved measures of disease progression.

**[0113]** Safety evaluations were conducted throughout the study and included review of adverse events (AEs), physical examinations, laboratory evaluations, anti-drug antibodies (ADA), and electrocardiography (ECG).

**[0114]** PFS was defined as the time (in months) from the date of randomization to the date of the first observation of progression based on radiologic assessment by RECIST or definitive clinical disease progression as assessed by the investigators (e.g., new occurrence of positive fluid cytology), or the date of death, whatever the cause, in the absence of progressive disease.

**[0115]** To demonstrate that relative FRA expression levels in the tumor tissue or a threshold level of expression are important for improved farletuzumab-mediated clinical improvements in patients with non-small cell lung adenocarcinoma (NSCLC) in combination with standard-of-care chemotherapies (carboplatin + paclitaxel; carboplatin + pemetrexed; or cisplatin + pemetrexed), 5 $\mu$ m tissue section slides prepared from tumor lesions from chemo-naive, stage IV patients with NSCLC adenocarcinoma were analyzed for cytoplasmic or membrane FRA expression levels.

**[0116]** FRA protein levels were determined by immunohistochemical (IHC) analysis for this study using the 26B3 anti-FRA antibody (O'Shannessy et al., *Oncotarget*, 2012; 3(4):414-25; also contained within Catalog no. IPI4006K G10 (Biocare Medical; Concord, CA); see also U.S. Patent 8,475,795) following standard protocol instructions. To quantify FRA expression, levels were determined by one of two procedures and analyzed by trained pathologists in a blinded fashion. Either cytoplasmic or membrane FRA expression was determined via digital imaging or manual microscopic assessment of 26B3-stained tissue sections using methods as previously described (Potts, *Drug Discov Today*, 2009; 14(19-20):935-41;

O'Shannessy et al., *Oncotarget*, 2012; 3(4):414-25; U.S. Patent 8,475,795; manufacturer's instructions, Catalog no. IPI4006K G10 (Biocare Medical; Concord, CA).

**[0117]** Digital analysis of FRA in 26B3-stained sectioned tissue slides was conducted using Image Analysis via the Cell Map and/or Stain Map algorithms according to Flagship Biosciences (Westminster, CO). Manual analysis of FRA expression was conducted by microscopic evaluation by two independent labs with trained pathologists skilled in the art.

**[0118]** To quantify membrane or cytoplasmic FRA expression levels derived by digital imaging or manual analysis, an algorithm was developed to quantify FRA staining as a percentage of tumor positivity and signal intensity. These values identified for use in the manual pathology scoring method are represented as FRA M Score (FRAMSCOR) for membrane staining. The values identified for use in the digital imaging analysis are represented as HBS Score (HBSCOR) for cytoplasmic staining. Previous methods used to accurately quantify biomarker expression levels determined that, in order to achieve highly accurate scoring, tumor sections must be well defined and IHC staining needs to be uniform. To assess the quality of tissue sections and staining obtained for patient inclusion into this study, a blinded pathologist skilled in the art independently evaluated the integrity of the FRA 26B3- stained NSCLC tumor tissue sections to compare FRA expression in pleural tissue using pleural tissue derived standards. After full assessment, the pathologist deemed 85 of the 130 tissue slides suitable for FRAMSCOR and HBSCOR pleural tissue analysis. While suitable for detecting the "presence" of FRA expression, detailed expression analysis of the remaining 45 slides was not possible due to: 1.) poor tissue preservation that precluded the ability to accurately analyze membrane expression; 2.) inability to accurately determine FRA expression in pleural tissue (a subset of slides only contained malignant cells within a pleural effusion background); and/or 3.) gross over staining that prohibited precise staining analysis in a linear range. Figure 1 shows the suitability of tissue integrity/morphology and staining versus examples of those not suitable for FRA

expression quantification by FRAMSCOR or HBSCOR that were part of the original clinical set of the intent-to-treat population (ITT).

[0119] To determine FRA staining intensity in the 85 suitable pleural tumor sections, a pleural tissue reference data set was first established using samples from unrelated patients with low, moderate and high FRA expression levels. This data set represents a standard by which relative FRA expression levels are compared among patients and quantified using the manual and digital analysis FRAMSCOR and HBSCOR methods. Figure 2 provides an example of the reference data set used for scoring +1 (low expression), +2 (moderate expression) and +3 (high expression) as known by those skilled in the art of FRA in NSCLC adenocarcinoma samples. In addition, the percentage of tumor positivity is also factored into the algorithm for each method (1 to 100% positivity of the tumor surface area) (not shown) using the formulas below.

[0120] The FRAMSCOR (M-score) is calculated as a weighted average, assuming a 0, 1+, 2+, 3+ scoring system as follows (see Figure 2):

$$x = \% \text{ tumor stained at } 1+$$

$$y = \% \text{ tumor stained at } 2+$$

$$z = \% \text{ tumor stained at } 3+$$

Then

$$M = \frac{x + 2y + 3z}{6}$$

For purposes of example, if a patient's tumor has 20% 26B3-FRA staining at +1, 10% 26B3-FRA staining at +2 and 20% 26B3-FRA staining at +3, then the M score = 16.6

[0121] The HBSCOR reports the mean optical density value for biomarker staining (in this case 26B3-staining of FRA) computed from all the cells in a target tissue compartment. It utilizes proprietary tissue recognition features to determine the tissue compartment via a linear score and a continuous extension of the H-score with no cell classification. The H-score is a standard scoring method that is commonly used by pathologists and those skilled in the art to score biomarker expression in

tissues, which is basically the sum of the intensity scores at all intensity levels (1+, + 2 x 2+, + 3x 3+). The HBSCOR is derived from the sum of cell measurements (optical density) divided by the total number of cells. HBSCOR in turn reports the value for biomarker staining computed from all the cells in a target tissue compartment. This calculation is quantified using the following formula:

$$\text{HBSCOR} = \frac{\sum_{\text{Cells}} \text{Cell Measurement}}{\text{Number of Cells}}$$

**[0122]** Evaluable slides containing patient-derived tissue stained for FRA using 26B3 antibody were analyzed and quantified using the FRAMSCOR and HBSCOR methods described above under blinded analysis. The data was first analyzed using an expression cut-point analysis. Folate Receptor Alpha (FRA) expression correlation to overall survival (OS) was conducted to determine if high or low FRA expression correlated with therapeutic response to farletuzumab treatment over placebo control treated patients. Figure 3 shows an example of this analysis whereby patient specimens were quantified for FRA expression via the HBSCOR method. As shown, a significant response in patient OS is observed, as clinically positive Hazard Ratios occur in farletuzumab-treated patients that express higher levels of FRA versus those with low expression. To determine the overall impact of FRA expression and farletuzumab treatment to patient clinical response as a function of progression free survival (PFS) or OS, Kaplan Meier analysis was conducted. As shown in Figure 4, patients selected for high FRA membrane expression using FRAMSCOR showed a statistically significant clinical improvement in OS (8.4 month improvement, HR 0.54; p = 0.0266) when treated with farletuzumab + SOC as compared to those treated with placebo + SOC only (p = 0.386). Improved PFS responses were also found in patients exhibiting high FRA membrane staining vs. low membrane staining (not shown). Similar effects were observed when patients with optimal FRA cytoplasmic expression using the HBSCOR method was used whereby patients treated with farletuzumab had a statistically significant clinical

benefit in PFS and OS (Figure 5, Panel B) as compared to those treated with farletuzumab expressing less than optimal FRA levels (Figure 5, Panel A). These findings teach the use of identifying FRA expression in biopsies of lung cancer patients to determine those exhibiting a minimal threshold level using the FRAMSCOR or HBSCOR method to improve therapeutic benefit than just recruiting patients with tumor exhibiting any FRA expression as was done for the ITT population which failed to demonstrate statistical clinical benefit in patients treated with farletuzumab + SOC over those treated with placebo + SOC (PFS HR 0.91, p = 0.7045 and OS HR 0.91, p = 0.6525).

**[0123]** Multivariate analysis was conducted using standard factors that may affect NSCLC lung cancer patient clinical responses to SOC treatment such as smoking, age or ECOG status. No effects were seen in the multivariate analysis that impacts the statistically positive effect of farletuzumab on clinical response in patients with high levels of FRA.

**[0124]** These findings teach the use of employing good quality tissue biopsies containing pleural malignant tissue and stained in a manner than enables quantifiable analysis of FRA expression either localized to the membrane or in the cytoplasm for use in identifying patients who may respond to anti-folate receptor alpha antibody therapy. Subsequent analysis has found that patient samples derived from pleural effusions and fine needle aspirates can also be quantified to identify those exhibiting suitable FRA expression for anti-FRA therapeutic effects. In these cases a non-pleural based reference standard may be developed for comparison. Moreover, these results demonstrate that patients with tumors expressing higher levels of FRA in either membrane and/or cytoplasm as determined via FRAMSCOR or HBSCOR methods have a significant improvement in clinical outcomes including PFS and OS. These teachings should position the use of anti-FRA antibody therapies in NSCLC patients most likely to respond to anti-FRA antibody therapy(s). Levels of FRA are important for defining patients who will respond to anti-FRA antibody therapy in combination with SOC. Future trials and clinical utility should incorporate the treatment of patients exhibiting necessary threshold levels of FRA above the minimal cutpoint for efficacy. Those levels defined here include those

with FRAMSCORs greater than or equal to 7 and HBSCORs greater than or equal to 0.25 (see Figure 3). On average this translates to NSCLC tumors with greater than 42% expression of FRA at a +1 intensity, 21% expression of FRA at a +2 intensity and/or tumors with greater than 14% expression of FRA at a +3 intensity.

What is claimed is:

1. A method for treating a folate receptor alpha (FRA)-expressing lung cancer in a patient with an antibody that immunospecifically binds folate receptor alpha (FRA), said method comprising:

    determining the patient's FRA expression level in a biological sample of the patient;

    comparing the patient's FRA expression level to a reference FRA expression level; and

    administering a therapeutically effective amount of said antibody to said patient if said patient's FRA expression level equals or exceeds said reference FRA expression level.

2. A method for treating a folate receptor alpha (FRA)-expressing lung cancer in a patient, the method comprising administering to the patient an antibody that immunospecifically binds folate receptor alpha (FRA), wherein the patient's FRA expression level equals or exceeds a reference FRA expression level.

3. A method for predicting a likelihood of responsiveness to treatment with an antibody that immunospecifically binds folate receptor alpha (FRA) in a patient having a folate receptor alpha (FRA)-expressing lung cancer, said method comprising:

    determining the patient's FRA expression level in a biological sample of the patient; and

    comparing said patient's FRA expression level to a reference FRA expression level,

    wherein said patient is likely to respond to treatment with said antibody that immunospecifically binds FRA if said patient's FRA expression level equals or exceeds said reference FRA expression level.

4. The method of any preceding claim wherein said reference FRA expression level corresponds to the FRA expression level above which a patient population afflicted

with said FRA-expressing lung cancer that is administered said antibody that immunospecifically binds FRA demonstrated a statistically significant improvement in at least one clinical outcome relative to a patient population afflicted with said FRA-expressing lung cancer that is administered placebo.

5. The method of claim 4 wherein said patient populations were further administered a chemotherapeutic agent.
6. The method of claim 5 wherein said chemotherapeutic agent comprises a taxane, cisplatin, carboplatin, and/or pemetrexed.
7. The method of any preceding claim wherein said FRA expression level is measured by protein quantification or RNA quantification.
8. The method of any preceding claim wherein said FRA expression level is measured by immunohistochemical analysis.
9. The method of any preceding claim wherein said FRA expression level is cytoplasmic FRA expression.
10. The method of any one of claims 1 to 8 wherein said FRA expression level is membranous FRA expression.
11. The method of any preceding claim wherein said antibody that immunospecifically binds FRA is conjugated to a toxin.
12. The method of claim 11 wherein said toxin comprises a microtubule inhibitor, a DNA damaging agent, a DNA repair inhibitor, or a signal transduction inhibitor.
13. The method of claim 11 wherein said DNA damaging agent comprises a radionuclide.

14. The method of any preceding claim wherein said antibody that immunospecifically binds FRA is:

an antibody comprising SEQ ID NO:1 as CDRH1, SEQ ID NO:2 as CDRH2, SEQ ID NO:3 as CDRH3, SEQ ID NO:4 as CDRL1, SEQ ID NO:5 as CDRL2 and SEQ ID NO:6 as CDRL3;

an antibody comprising a mature light chain variable region comprising the amino acid sequence of SEQ ID NO:7 and/or a mature heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 8;

farletuzumab;

an antibody that specifically binds to folate receptor alpha comprising a mature light and heavy chain variable regions having at least 90% and preferably at least 95% or 99% sequence identity to SEQ ID NO: 7 and SEQ ID NO: 8, respectively; or

an antibody or derivative thereof can competitively inhibit binding of farletuzumab to folate receptor alpha.

15. The method of any preceding claim further comprising administering a therapeutically effective amount of a chemotherapeutic agent to said patient.

16. The method of claim 15 wherein said chemotherapeutic agent comprises a platinum-containing compound.

17. The method of claim 16 wherein said platinum-containing compound comprises cisplatin or carboplatin.

18. The method of any preceding claim further comprising administering a therapeutically effective amount of a taxane to said patient.

19. The method of claim 18 wherein said taxane is paclitaxel.

20. The method of claim 15, 16, 17, 18, or 19, further comprising administering a therapeutically effective amount of pemetrexed to said patient.
21. The method of any one of claims 1 to 14, further comprising administering carboplatin and paclitaxel to said patient.
22. The method of claim 21 wherein carboplatin is administered to said patient to achieve an area under the curve (AUC) of about 6 or less.
23. The method of claim 21 or 22 wherein paclitaxel is administered to said patient at a dose of about 50 mg/m<sup>2</sup> to about 250 mg/m<sup>2</sup>.
24. The method of any one of claims 1 to 14, further comprising administering carboplatin and pemetrexed to said patient.
25. The method of claim 24 wherein said carboplatin is administered to said patient to achieve an area under the curve of about 5-6 or less.
26. The method of claim 24 or 25 wherein said pemetrexed is administered to said patient at a dose of about 400 to about 600 mg/m<sup>2</sup>.
27. The method of any one of claims 1 to 14, further comprising administering cisplatin and pemetrexed to said patient.
28. The method of claim 27 wherein cisplatin is administered to said patient at a dose of about 50 mg/m<sup>2</sup> to about 250 mg/m<sup>2</sup>.
29. The method of claim 27 or 28 wherein pemetrexed is administered at a dose of about 400 mg/m<sup>2</sup> to about 600 mg/m<sup>2</sup>.

30. The method of any preceding claim wherein said FRA expression level is determined by immunoassay with at least one of the following antibodies:

- (a) an antibody that binds the same epitope as farletuzumab;
- (b) an antibody comprising SEQ ID NO:1 (GFTFSGYGLS) as CDRH1, SEQ ID NO:2 (MISSGGSYTYYADSVKG) as CDRH2, SEQ ID NO:3 (HGDDPAWFAY) as CDRH3, SEQ ID NO:4 (SVSSSISSNNLH) as CDRL1, SEQ ID NO:5 (GTSNLAS) as CDRL2 and SEQ ID NO:6 (QQWSSYPYMYT) as CDRL3;
- (c) an antibody comprising a mature light chain variable region comprising the amino acid sequence of SEQ ID NO:7 and a mature heavy chain variable region comprising the amino acid SEQ ID NO: 8;
- (d) farletuzumab;
- (e) the 548908 antibody;
- (f) an antibody that binds the same epitope as the 548908 antibody;
- (g) the 6D398 antibody;
- (h) an antibody that binds the same epitope as the 6D398 antibody;
- (i) the BN3.2 antibody;
- (j) an antibody that binds the same epitope as the BN3.2 antibody;
- (k) an antibody that binds the same epitope as the 26B3 antibody;
- (l) an antibody comprising SEQ ID NO: 14 (GYFMN) as CDRH1, SEQ ID NO: 15 (RIFPYNGDTFYQNQKFKG) as CDRH2, SEQ ID NO: 16 (GTHYFDY) as CDRH3, SEQ ID NO: 17 (RTSENIFSYLA) as CDRL1 , SEQ ID NO: 18 (NAKTLAE) as CDRL2 and SEQ ID NO: 19 (QHHYAFPWT) as CDRL3;
- (m) the 26B3 antibody;
- (n) an antibody that binds the same epitope as the 19D4 antibody;
- (o) an antibody comprising SEQ ID NO: 20 (HPYMH) as CDRH1, SEQ ID NO: 21 (RIDPANGNTKYDPKFQG) as CDRH2, SEQ ID NO: 22 (EEVADYTM DY) as CDRH3, SEQ ID NO: 23 (RASESVDTYGNNFIH) as CDRL1, SEQ ID NO: 24 (LASNLES) as CDRL2 and SEQ ID NO: 25 (QQNNNGDPWT) as CDRL3;
- (p) the 19D4 antibody;

- (q) an antibody that binds the same epitope as the 9F3 antibody;
- (r) an antibody comprising SEQ ID NO: 26 (SGYYWN) as CDRH1, SEQ ID NO: 27 (YIKSDGSNNYNPSLKN) as CDRH2, SEQ ID NO: 28 (EWKAMDY) as CDRH3, SEQ ID NO: 29 (RASSTVSYSYLH) as CDRL1, SEQ ID NO: 30 (GTSNLAS) as CDRL2 and SEQ ID NO: 31 (QQYSGYPLT) as CDRL3;
- (s) the 9F3 antibody;
- (t) an antibody that binds the same epitope as the 24F12 antibody;
- (u) an antibody comprising SEQ ID NO: 32 (SYAMS) as CDRH1, SEQ ID NO: 33 (EIGSGGSYTYYPDTVTG) as CDRH2, SEQ ID NO: 34 (ETTAGYFDY) as CDRH3, SEQ ID NO: 35 (SASQGINNFLN) as CDRL1, SEQ ID NO: 36 (YTSSLHS) as CDRL2 and SEQ ID NO: 37 (QHFSKLPWT) as CDRL3;
- (v) the 24F12 antibody;
- (w) an antibody that comprises a variable region light chain selected from the group consisting of:
  - (i) LK26HuVK;
  - (ii) LK26HuVKY;
  - (iii) LK26HuVKPW; and
  - (iv) LK26HuVKPW,Y;
- (x) an antibody that comprises a variable region heavy chain selected from the group consisting of:
  - (i) LK26HuVH;
  - (ii) LK26HuVH FAIS,N;
  - (iii) LK26HuVH SLF;
  - (iv) LK26HuVH I,I;
  - (v) LK26KOLHuVH;
- (y) an antibody that comprises the heavy chain variable region LK26KOLHuVH (SEQ ID NO: 46) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41);
- (z) an antibody that comprises the heavy chain variable region LK26HuVH SLF (SEQ ID NO: 44) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41);

(aa) an antibody that comprises the heavy chain variable region LK26HuVH FAIS,N (SEQ ID NO: 43) and the light chain variable region LK26HuVKPW,Y (SEQ ID NO: 41); and

(bb) the murine monoclonal LK26 antibody.

31. The method of any preceding claim wherein said patient's FRA expression level is assessed by digital imaging technology or manual pathology quantification.

32. The method of claim 7 wherein said patient's FRA expression level is assessed by FRAMSCOR or HBSCOR.

33. The method of claim 32 wherein said reference FRA expression level is 42% +1 or greater anti-FRA staining.

34. The method of claim 32 wherein said reference FRA expression level is 21% +2 or greater anti-FRA staining.

35. The method of claim 32 wherein said reference FRA expression level is 14% +3 or greater staining.

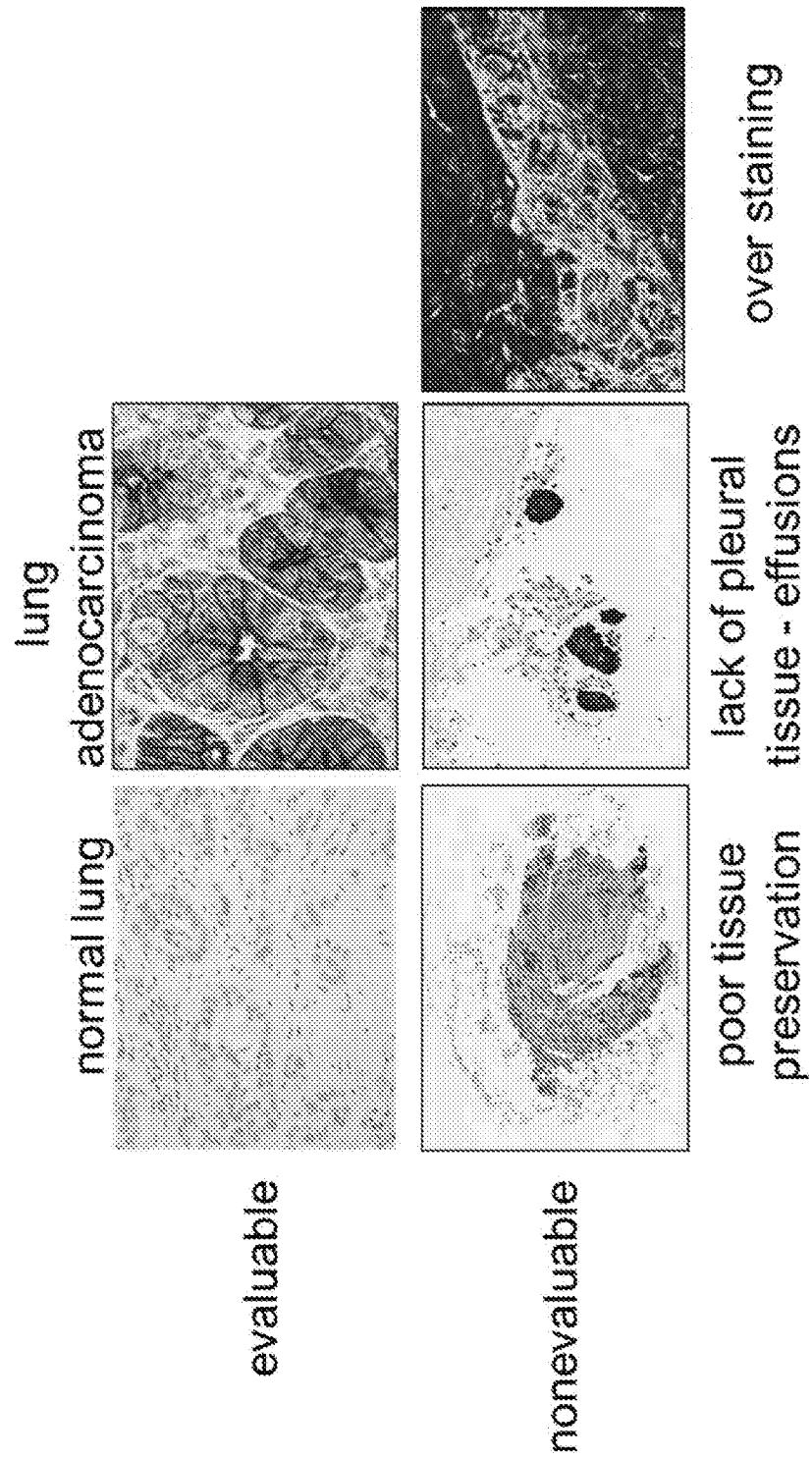
36. The method of claim 32 wherein said reference FRA expression level is a FRAMSCOR of 7.

37. The method of claim 32 wherein said reference FRA expression level is a HBSCOR of 0.25.

38. The method of any preceding claim wherein said biological sample is whole blood, serum, plasma, circulating cells, circulating tumor cells, free cells, tissue, pleural effusion, urine, saliva, sputum, or bronchial washing.

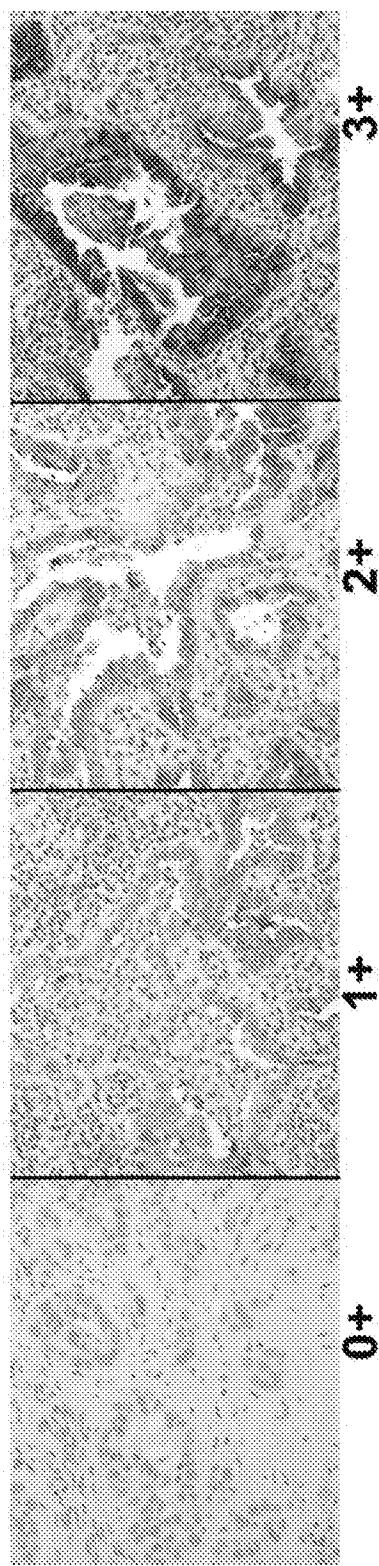
39. The method of any preceding claim wherein said biological sample comprises pleural tissue.
40. The method of any preceding claim wherein said biological sample comprises pleural cells derived from effusion.
41. The method of claim 4 wherein said at least one clinical outcome is progression free survival and/or overall survival.
42. The method of any preceding claim wherein said FRA-expressing lung cancer is FRA-expressing non-small cell lung cancer (NSCLC).
43. The method of claim 42 wherein said NSCLC is adenocarcinoma.

Figure 1



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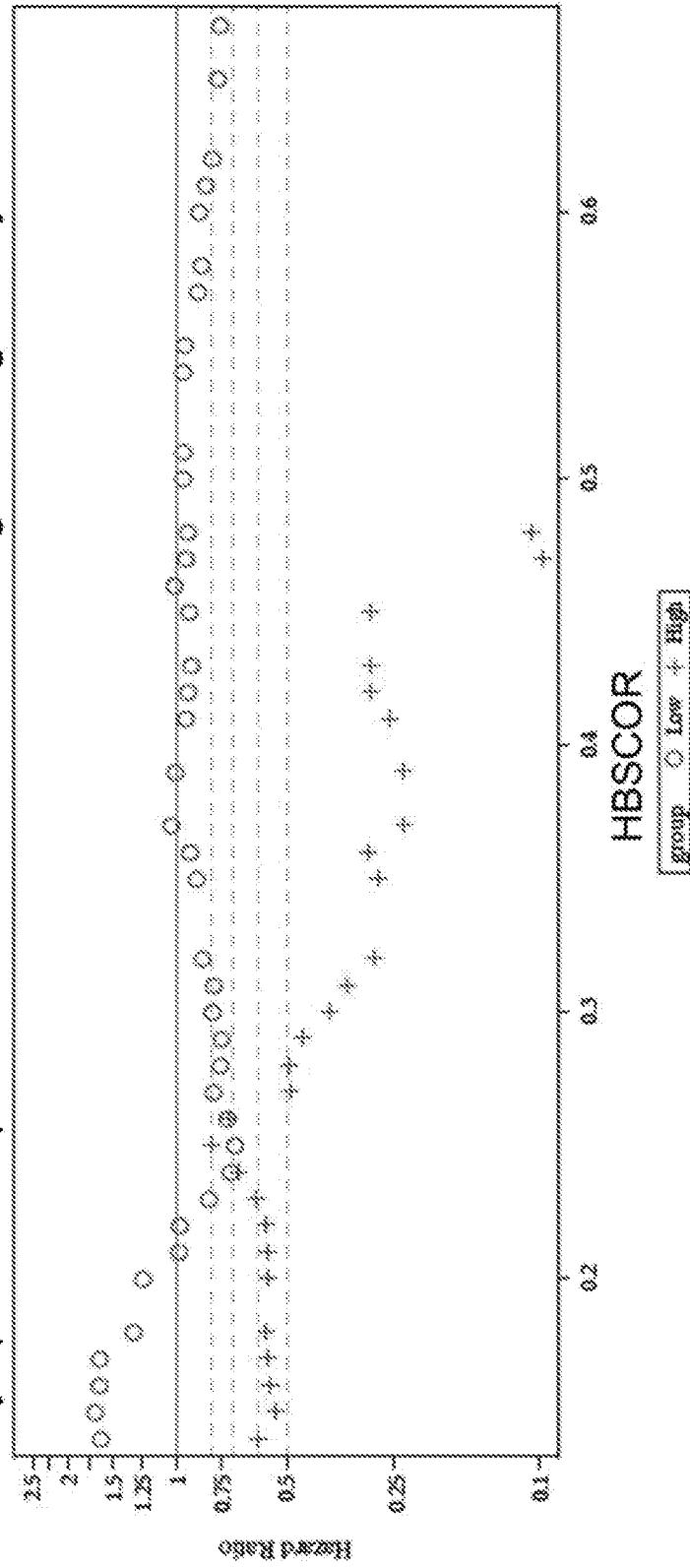
Figure 2  
Non-Small Cell Lung Adenocarcinoma staining  
gradation:



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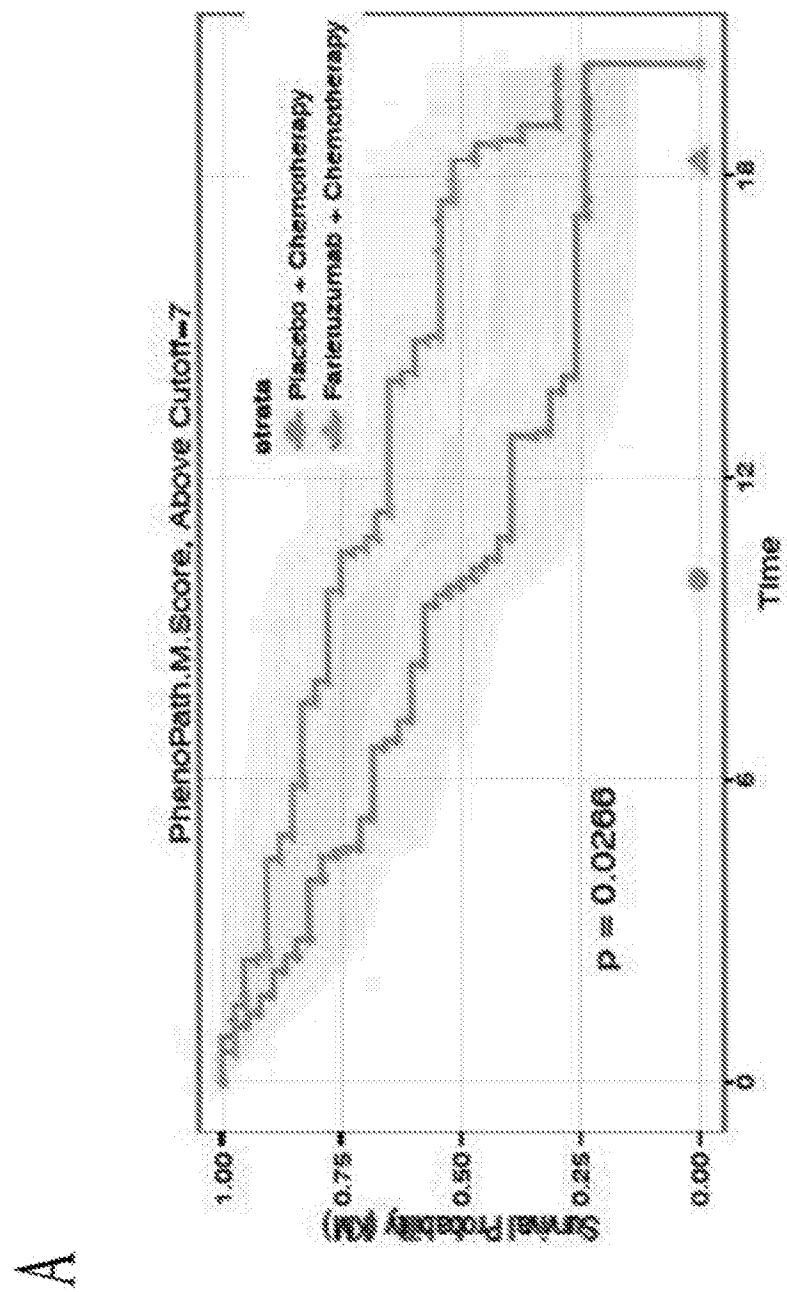
Figure 3

Hazard ratios of patient OS treated with farletuzumab at different FRA cytoplasmic expression levels detected via digital image analysis



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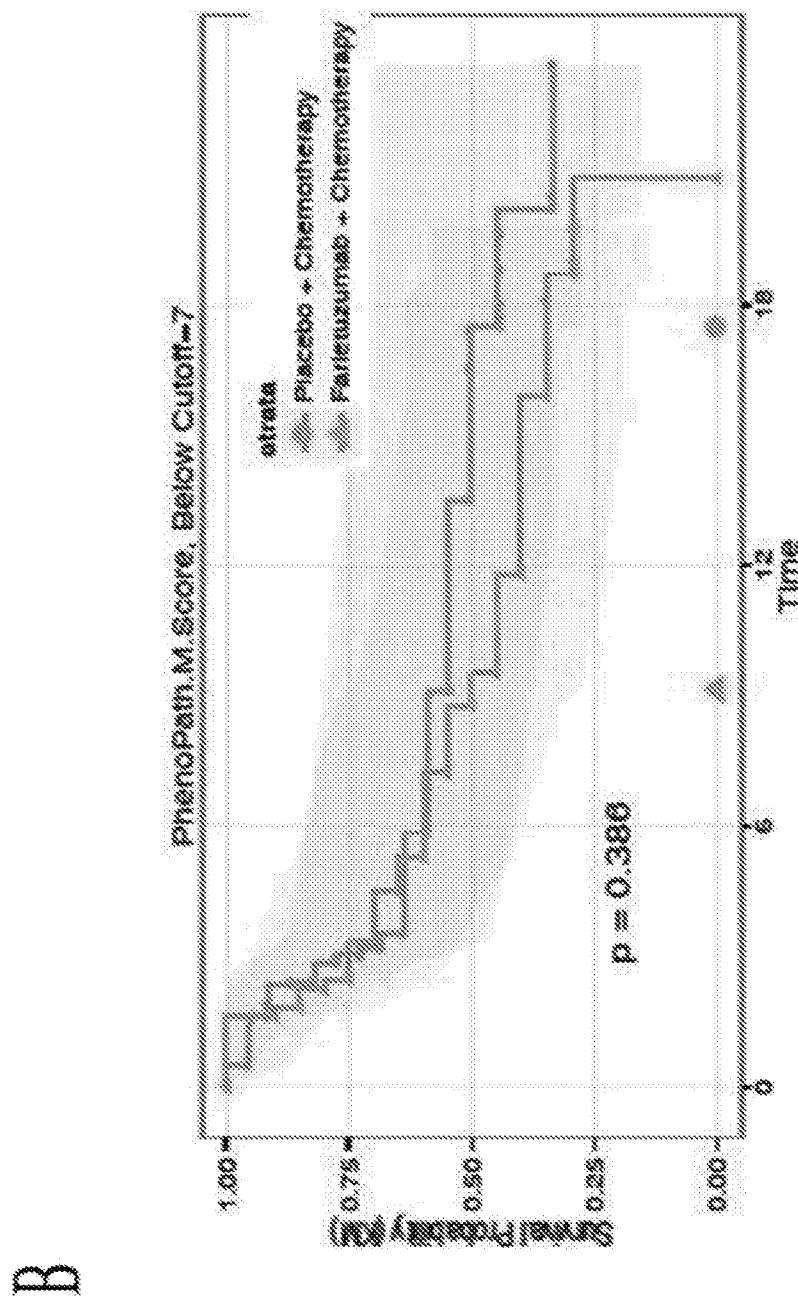
Figure 4



A

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Figure 4

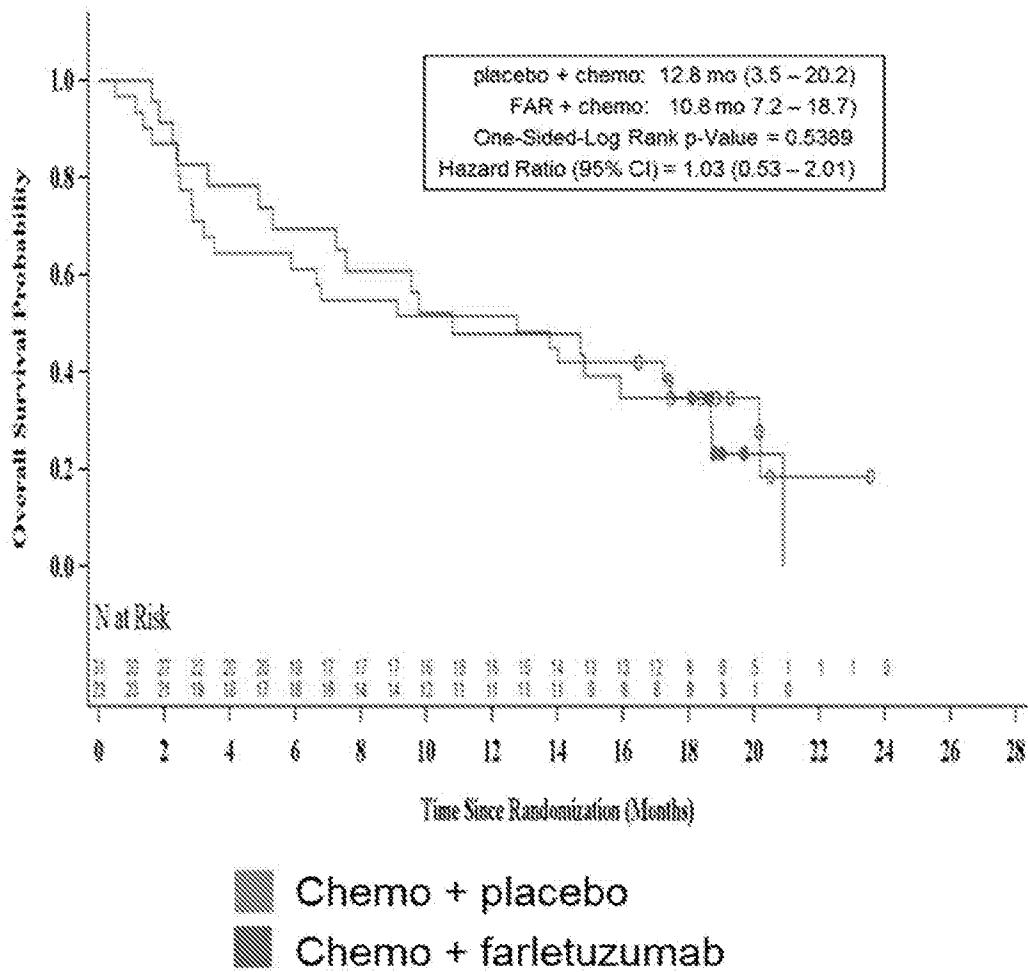


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6/7

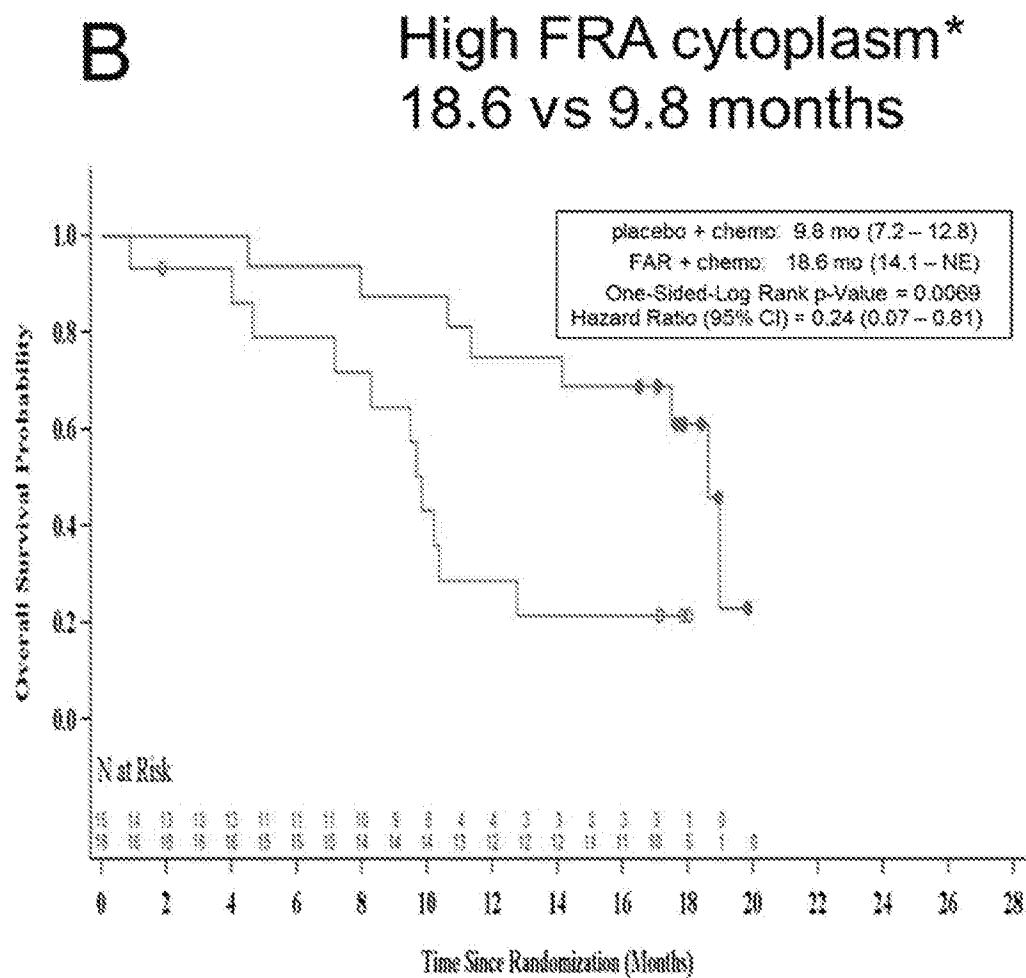
Figure 5

A

Low FRA cytoplasm  
10.8 vs 12.8 months

7/7

Figure 5



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Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp Lys  
210 215 220

Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro  
225 230 235 240

Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser  
245 250 255

### Sequence\_Listing

Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp  
260 265 270

Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn  
275 280 285

Ala Lys Thr Lys Pro Arg Glu Glu Glu Tyr Asn Ser Thr Tyr Arg Val  
290 295 300

Val Ser Val Leu Thr Val Leu His Glu Asp Trp Leu Asn Gly Lys Glu  
305 310 315 320

Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys  
325 330 335

Thr Ile Ser Lys Ala Lys Gly Glu Pro Arg Glu Pro Glu Val Tyr Thr  
340 345 350

Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Glu Val Ser Leu Thr  
355 360 365

Cys Leu Val Lys Glu Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu  
370 375 380

Ser Asn Gly Glu Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu  
385 390 395 400

Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys  
405 410 415

Ser Arg Trp Glu Glu Gly Asn Val Phe Ser Cys Ser Val Met His Glu  
420 425 430

Ala Leu His Asn His Tyr Thr Glu Lys Ser Leu Ser Leu Ser Pro Gly  
435 440 445

Lys

<210> 9  
<211> 962  
<212> DNA  
<213> Homo sapiens

<400> 9  
tcaaggtaa acgacaagga cagacatggc tcagcggatg acaacacagc tgctgctcct 60

tctagtgtgg gtggctgttag taggggaggc tcagacaagg attgcatggg ccaggactga 120

Sequence\_Listing

gcttctcaat	gtctgcata	acgccaagca	ccacaaggaa	aagccaggcc	ccgaggacaa	180	
gttgcata	gagtcgtcga	cctggaggaa	aatgcctgc	tgttctacca	acaccagcca	240	
ggaagccat	aaggatgtt	cctacctata	tagattcaac	tggaaaccact	gtggagagat	300	
ggcacctg	cc tgcaaacggc	at ttcata	ggacacctgc	ctctacgagt	gctccccaa	360	
cttggggccc	tggatccagc	agg tggatca	gagctggcgc	aaagagcggg	tactgaacgt	420	
gcccctgt	gc aa	aaagaggact	gtgagcaatg	gtggaaagat	tgtgcac	480	
caagagca	ac tggcacaagg	gctggaactg	gacttcagg	tttaacaagt	gchgagtgg	540	
agctgc	ctgc caac	tttcc	at ttcactt	ccccacaccc	actgttctgt	600	
ctggactc	ac tc	c tccataaagg	tcagcaacta	cagccgaggg	agtggccgct	660	
gtggttcg	ac cc	ccagcccagg	gcaacccaa	tgaggaggtg	gcgaggttct	720	
catgagtgg	gctggccct	gggcagcctg	gccttcctg	cttagcctgg	cccta atgct	780	
gctgtggc	tg ctc	tcagctgac	ctcc	tttac	cttctgatac	ctggaaatcc	840
cagccccaca	gctcccaact	at tggttcc	tgctccatgg	tcgggcctct	gacagccact	900	
ttgaataaac	cagacaccgc	acatgtgtct	tgagaattat	ttggaaaaaa	aaaaaaaaaa	960	
aa						962	

<210> 10

<211> 257

<212> PRT

<213> Homo sapiens

<400> 10

Met	Ala	Gln	Arg	Met	Thr	Thr	Gln	Leu	Leu	Leu	Leu	Leu	Val	Trp	Val
1				5				10						15	

Ala	Val	Val	Gly	Gl	u	Ala	Gln	Thr	Arg	Ile	Ala	Trp	Ala	Arg	Thr	Gl	u
			20					25						30			

Leu	Leu	Asn	Val	Cys	Met	Asn	Ala	Lys	His	His	Lys	Gl	u	Lys	Pro	Gly
						35		40			45					

Pro	Gl	u	Asp	Lys	Leu	His	Gl	u	Gln	Cys	Arg	Pro	Trp	Arg	Lys	Asn	Ala
	50				55						60						

Cys	Cys	Ser	Thr	Asn	Thr	Ser	Gln	Gl	u	Ala	His	Lys	Asp	Val	Ser	Tyr
	65				70					75					80	

Leu	Tyr	Arg	Phe	Asn	Trp	Asn	His	Cys	Gly	Gl	u	Met	Ala	Pro	Ala	Cys
								85		90						95

Lys Arg His Phe Ile Gln Asp Thr Cys Leu Tyr Glu Cys Ser Pro Asn

## Sequence Listing 105

Leu	Gly	Pro	Trp	Ile	Gln	Gln	Val	Asp	Gln	Ser	Trp	Arg	Lys	Gl u	Arg
115							120					125			
Val	Leu	Asn	Val	Pro	Leu	Cys	Lys	Gl u	Asp	Cys	Gl u	Gln	Trp	Trp	Gl u
130						135					140				
Asp	Cys	Arg	Thr	Ser	Tyr	Thr	Cys	Lys	Ser	Asn	Trp	His	Lys	Gl y	Trp
145					150					155					160
Asn	Trp	Thr	Ser	Gly	Phe	Asn	Lys	Cys	Al a	Val	Gl y	Al a	Al a	Cys	Gl n
				165					170					175	
Pro	Phe	His	Phe	Tyr	Phe	Pro	Thr	Pro	Thr	Val	Leu	Cys	Asn	Gl u	Ile
				180				185					190		
Trp	Thr	His	Ser	Tyr	Lys	Val	Ser	Asn	Tyr	Ser	Arg	Gl y	Ser	Gl y	Arg
				195			200				205				
Cys	Ile	Gln	Met	Trp	Phe	Asp	Pro	Al a	Gln	Gl y	Asn	Pro	Asn	Gl u	Gl u
						215					220				
Val	Al a	Arg	Phe	Tyr	Al a	Al a	Al a	Met	Ser	Gl y	Al a	Gl y	Pro	Trp	Al a
225				230						235					240
Al a	Trp	Pro	Phe	Leu	Leu	Ser	Leu	Al a	Leu	Met	Leu	Leu	Trp	Leu	Leu
									245		250				255

<210> 11  
<211> 119  
<212> PRT  
<213> Arti fi ci al Sequence

<220>  
<221> source  
<223> /note="Descri pti on of Arti fi ci al Sequence: Syntheti c  
pol ypepti de"

<220>  
<221> MOD\_RES  
<222> (3)..(3)  
<223> Any ami no aci d

<220>  
<221> MOD\_RES  
<222> (6)..(6)  
<223> Any ami no acid

### Sequence\_Listing

<400> 11  
Gln Val Xaa Leu Gln Xaa Ser Gly Gly Asp Leu Val Lys Pro Gly Gly  
1 5 10 15

Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Gly Tyr  
20 25 30

Gly Leu Ser Trp Val Arg Gln Thr Pro Asp Lys Arg Leu Glu Trp Val  
35 40 45

Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Phe  
65 70 75 80

Leu Gln Met Ser Ser Leu Lys Ser Asp Asp Thr Ala Ile Tyr Ile Cys  
85 90 95

Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly  
100 105 110

Thr Leu Val Thr Val Ser Ala  
115

<210> 12

<211> 110

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic polypeptide"

<400> 12  
Asp Ile Glu Leu Thr Gln Ser Pro Ala Leu Met Ala Ala Ser Pro Gly  
1 5 10 15

Glu Lys Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Asn  
20 25 30

Asn Leu His Trp Tyr Gln Gln Lys Ser Glu Thr Ser Pro Lys Pro Trp  
35 40 45

Ile Tyr Gly Thr Ser Asn Leu Ala Ser Gly Val Pro Leu Arg Phe Arg  
50 55 60

Gly Phe Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser Ser Met Glu  
65 70 75 80

## Sequence\_Listing

Ala Glu Asp Ala Ala Thr Tyr Tyr Cys Glu Glu Trp Ser Ser Tyr Pro  
85 90 95

Tyr Met Tyr Thr Phe Gly Gly Thr Lys Leu Glu Ile Lys  
100 105 110

<210> 13

<400> 13  
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<210> 14  
<211> 5  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 14  
Gly Tyr Phe Met Asn  
1 5

<210> 15  
<211> 17  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 15  
Arg Ile Phe Pro Tyr Asn Gly Asp Thr Phe Tyr Asn Glu Lys Phe Lys  
1 5 10 15

Gly

<210> 16  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 16  
Gly Thr His Tyr Phe Asp Tyr  
1 5

## Sequence\_Listing

<210> 17  
<211> 11  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 17  
Arg Thr Ser Glu Asn Ile Phe Ser Tyr Leu Ala  
1 5 10

<210> 18  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 18  
Asn Ala Lys Thr Leu Ala Glu  
1 5

<210> 19  
<211> 9  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 19  
Gln His His Tyr Ala Phe Pro Trp Thr  
1 5

<210> 20  
<211> 5  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 20  
His Pro Tyr Met His  
1 5

<210> 21

Sequence\_Listing

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 21

Arg Ile Asp Pro Ala Asn Gly Asn Thr Lys Tyr Asp Pro Lys Phe Gln  
1 5 10 15

Gly

<210> 22

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 22

Gl u Gl u Val Ala Asp Tyr Thr Met Asp Tyr  
1 5 10

<210> 23

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 23

Arg Ala Ser Gl u Ser Val Asp Thr Tyr Gly Asn Asn Phe Ile His  
1 5 10 15

<210> 24

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 24

Leu Ala Ser Asn Leu Gl u Ser  
1 5

Sequence\_Listing

<210> 25  
<211> 9  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 25  
Gln Gln Asn Asn Gly Asp Pro Trp Thr  
1 5

<210> 26  
<211> 6  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 26  
Ser Gly Tyr Tyr Trp Asn  
1 5

<210> 27  
<211> 16  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 27  
Tyr Ile Lys Ser Asp Gly Ser Asn Asn Tyr Asn Pro Ser Leu Lys Asn  
1 5 10 15

<210> 28  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 28  
Glu Trp Lys Ala Met Asp Tyr  
1 5

<210> 29  
<211> 12  
<212> PRT

Sequence\_Listing

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 29

Arg Ala Ser Ser Thr Val Ser Tyr Ser Tyr Leu His  
1 5 10

<210> 30

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 30

Gly Thr Ser Asn Leu Ala Ser  
1 5

<210> 31

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 31

Gln Gln Tyr Ser Gly Tyr Pro Leu Thr  
1 5

<210> 32

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 32

Ser Tyr Ala Met Ser  
1 5

<210> 33

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

Sequence\_Listing

<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 33  
Glu Ile Gly Ser Gly Gly Ser Tyr Thr Tyr Tyr Pro Asp Thr Val Thr  
1 5 10 15

Gly

<210> 34  
<211> 9  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 34  
Glu Thr Thr Ala Gly Tyr Phe Asp Tyr  
1 5

<210> 35  
<211> 11  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 35  
Ser Ala Ser Gln Gly Ile Asn Asn Phe Leu Asn  
1 5 10

<210> 36  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 36  
Tyr Thr Ser Ser Leu His Ser  
1 5

<210> 37  
<211> 9  
<212> PRT  
<213> Artificial Sequence

Sequence\_Listing

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic peptide"

<400> 37  
Gln His Phe Ser Lys Leu Pro Trp Thr  
1 5

<210> 38  
<211> 110  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic polypeptide"

<400> 38  
Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
1 5 10 15

Asp Arg Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Asn  
20 25 30

Asn Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu  
35 40 45

Ile Tyr Gly Thr Ser Asn Leu Ala Ser Gly Val Pro Ser Arg Phe Ser  
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu Gln  
65 70 75 80

Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Tyr Pro  
85 90 95

Tyr Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys  
100 105 110

<210> 39  
<211> 110  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic polypeptide"

<400> 39  
Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
1 5 10 15

Sequence\_Listing

Asp Arg Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Asn  
20 25 30

Asn Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu  
35 40 45

Ile Tyr Gly Thr Ser Asn Leu Ala Ser Gly Val Pro Ser Arg Phe Ser  
50 55 60

Gly Ser Gly Ser Gly Thr Asp Tyr Thr Phe Thr Ile Ser Ser Leu Gln  
65 70 75 80

Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Tyr Pro  
85 90 95

Tyr Met Tyr Thr Phe Gln Gln Gly Thr Lys Val Glu Ile Lys  
100 105 110

<210> 40

<211> 110

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic polypeptide"

<400> 40

Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
1 5 10 15

Asp Arg Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Asn  
20 25 30

Asn Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Pro Trp  
35 40 45

Ile Tyr Gly Thr Ser Asn Leu Ala Ser Gly Val Pro Ser Arg Phe Ser  
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu Gln  
65 70 75 80

Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Tyr Pro  
85 90 95

Tyr Met Tyr Thr Phe Gln Gln Gly Thr Lys Val Glu Ile Lys  
100 105 110

Sequence\_Listing

<210> 41  
<211> 110  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic polypeptide"

<400> 41  
Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
1 5 10 15

Asp Arg Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Asn  
20 25 30

Asn Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Pro Trp  
35 40 45

Ile Tyr Gly Thr Ser Asn Leu Ala Ser Gly Val Pro Ser Arg Phe Ser  
50 55 60

Gly Ser Gly Ser Gly Thr Asp Tyr Thr Phe Thr Ile Ser Ser Leu Gln  
65 70 75 80

Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Tyr Pro  
85 90 95

Tyr Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys  
100 105 110

<210> 42  
<211> 119  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source  
<223> /note="Description of Artificial Sequence: Synthetic polypeptide"

<400> 42  
Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg Pro Ser Gln  
1 5 10 15

Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly Phe Thr Phe Ser Gly Tyr  
20 25 30

Gly Leu Ser Trp Val Arg Gln Pro Pro Gly Arg Gly Leu Glu Trp Val  
35 40 45

Sequence Listing

Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Val Thr Met Leu Arg Asp Thr Ser Lys Asn Gln Phe Ser  
65 70 75 80

Leu Arg Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly  
100 105 110

Ser Leu Val Thr Val Ser Ser  
115

<210> 43

<211> 119

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic polypeptide"

<400> 43

Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg Pro Ser Gln  
1 5 10 15

Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly Phe Thr Phe Ser Gly Tyr  
20 25 30

Gly Leu Ser Trp Val Arg Gln Pro Pro Gly Arg Gly Leu Glu Trp Val  
35 40 45

Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ser Lys Asn Gln Phe Ser  
65 70 75 80

Leu Arg Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly  
100 105 110

Ser Leu Val Thr Val Ser Ser  
115

Sequence\_Listing

<210> 44  
<211> 119  
<212> PRT  
<213> Artificial Sequence

<220>  
<221> source

<223> /note="Description of Artificial Sequence: Synthetic polypeptide"

<400> 44

Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg Pro Ser Gln  
1 5 10 15

Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly Phe Thr Phe Ser Gly Tyr  
20 25 30

Gly Leu Ser Trp Val Arg Gln Pro Pro Gly Arg Gly Leu Glu Trp Val  
35 40 45

Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Val Thr Met Leu Arg Asp Thr Ser Lys Asn Ser Leu Phe  
65 70 75 80

Leu Arg Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly  
100 105 110

Thr Thr Val Thr Val Ser Ser  
115

<210> 45  
<211> 119  
<212> PRT  
<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic polypeptide"

<400> 45

Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Arg Pro Ser Gln  
1 5 10 15

Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly Phe Thr Phe Ser Gly Tyr  
20 25 30

Gly Leu Ser Trp Val Arg Gln Pro Pro Gly Arg Gly Leu Glu Trp Val

35

40

Sequence\_Listing

45

Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Val Thr Met Leu Arg Asp Thr Ser Lys Asn Gln Phe Ser  
65 70 75 80

Leu Arg Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Ile Tyr Ile Cys  
85 90 95

Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly  
100 105 110

Ser Leu Val Thr Val Ser Ser  
115

<210> 46

<211> 119

<212> PRT

<213> Artificial Sequence

<220>

<221> source

<223> /note="Description of Artificial Sequence: Synthetic polypeptide"

<400> 46

Gl u Val Gln Leu Val Gl u Ser Gly Gly Gly Val Val Gln Pro Gly Arg  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ser Ala Ser Gly Phe Thr Phe Ser Gly Tyr  
20 25 30

Gly Leu Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Gl u Trp Val  
35 40 45

Ala Met Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Phe  
65 70 75 80

Leu Gln Met Asp Ser Leu Arg Pro Glu Asp Thr Gly Val Tyr Phe Cys  
85 90 95

Ala Arg His Gly Asp Asp Pro Ala Trp Phe Ala Tyr Trp Gly Gln Gly  
100 105 110

Thr Pro Val Thr Val Ser Ser

