ANTIBODIES THAT BIND TO LRP5 PROTEINS AND METHODS OF USE

STATEMENT AS TO FEDERALLY SPONSORED RESEARCH

[0001] None.

REFERENCE TO RELATED APPLICATIONS

[0002] This application claims the benefit of the priority dates of U.S. provisional application 62/886,913, filed August 14, 2019, the contents of which are incorporated herein by reference in its entirety.

SEQUENCE LISTING

[0002.1] 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 November 7, 2022, is named MODM-005-USNTL_SL.txt and is 70,000 bytes in size.

THE NAMES OF THE PARTIES TO A JOINT RESEARCH AGREEMENT

[0003] None.

BACKGROUND

[0004] Wnt signaling has a crucial role in the regulation of various cellular processes such as cell fate determination, proliferation, survival, polarity and migration[1]. Perturbations, either as a result of altered expression or mutations in the Wnt signaling pathway have been implicated in defects in embryonic development as well as in various pathologies such as cancer and osteoporosis[2-4]. Wnt signaling leads to the activation of the canonical and non-canonical signaling pathways[1,5]. The non-canonical pathway activates signaling molecules that do not involve the nucleus or transcription but rather activate cytoplasmic signals that regulate the cytoskeleton and calcium stores. This pathway primarily plays a role in regulating cell polarity or migration. The canonical pathway predominantly controls transcriptional activity by regulating the cytoplasmic levels of β-catenin. In unstimulated conditions, beta-catenin is associated with a destruction complex, comprised of Axin, APC, CK1 and GSK3b, which results in the phosphorylation, ubiquitylation and proteasomal degradation of beta-catenin. Wnt signaling destabilizes this complex, resulting in the accumulation of "free" beta-catenin in the cytosol that translocates to the nucleus and acts as a co-activator for TCF/LEF mediated transcription. Wnts bind to the frizzled family of seven transmembrane domain receptors as well as to either LRP5 or LRP6 leading to the initiation of the canonical signaling pathway[6-8].

[0005] LRP5 and LRP6 are functionally redundant single-pass transmembrane receptors that share approximately 70% homology. Binding of Wnt ligands to Fz and LRP5/6 leads to the recruitment of the destruction complex and Dishevelled (Dsh/Dvl) and the phosphorylation of LRP5/6 on PPPSPxS motifs located in the intracellular domain9. This phosphorylation is mediated by GSK3b and CK1 which in turn results in diminished GSK3b activity, inhibiting beta-catenin phosphorylation and subsequent proteosomal degradation and enhanced TCF/LEF mediated transcriptional activity. LRP5 is widely expressed during embryonic development and in adult tissues. Mutations in LRP5 have been associated with bone mass diseases and several mouse models with LRP5 knockout or mutations exhibit alterations in bone development[10].

[0006] LRP5 expression has also been shown to be elevated in human malignant tissues and human cancer cell lines such as osteosarcoma and Wnt signaling in such cell lines was shown to be diminished with the overexpression of a dominant-negative LRP5[11-13]. Moreover, LRP5 seems to also have an important role in regulating cell invasion capacity and cell motility[14-16]. While studies have shown that LRP6 is more potent than LRP5 in transducing the Wnt signal, recent genetic experiments have shown that some Wnt ligands require both receptors to be present to generate a canonical signal (17). Because of the importance of LRPS in regulating Wnt signaling and its established role in several human diseases, LRPS is becoming an increasingly important target for therapeutic drug development. There is significant biology surrounding LRP5 and its role in Wnt signaling and in the pathogenesis of various diseases that still remains to be discovered and a deep toolbox of synthetic antibodies will help to systematically expose these roles and also provide additional targeted therapeutics.

[0007] Wnt signaling leads to the activation of the canonical and non-canonical signaling pathways. The non-canonical pathway activates signaling molecules that do not involve the nucleus or transcription but rather activate cytoplasmic signals that regulate the cytoskeleton and calcium levels. This pathway primarily plays a role in regulating cell polarity or migration.

[0008] The canonical pathway predominantly controls transcriptional activity by regulating the cytoplasmic levels of β-catenin. In unstimulated conditions, β-catenin is associated with a destruction complex, comprised of Axin, APC, CD1 and GSK β, which results in the phosphorylation, ubiquitylation and proteasomal degradation of β-catenin. Wnt signaling is active when Wnt binds to frizzled (FDZ), a 7-pass transmembrane receptor, and to a co-receptor low density lipoprotein receptor-related protein (either LRP5 or LRP6). This signaling destabilizes the complex, in part by attracting disheveled (Dsh/Dvl) to the plasma membrane, resulting in the accumulation of β-catenin, which then travels to the nucleus and activates TCF/LEF-mediated transcription.

[0009] The invention described below identifies a novel set of synthetic antibodies targeting the extracellular epitopes of LRP5, by taking advantage of state-of-the-art antibody phage display libraries and technology.

SUMMARY

In one embodiment provided herein is an antibody that specifically binds LRP5, comprising a light chain variable region and/or a heavy chain variable region, the heavy chain variable region comprising complementarity determining regions CDR-H1, CDR-H2 and CDR-H3, the light chain variable region comprising complementarity determining region CDR-L1, CDR-L2 and CDR-L3, and with the amino acid sequences of said CDRs comprising or consisting of sequences selected from: CDR sequence sets of anti-LRP5 antibodies: LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6. In one embodiment the amino acid sequences of said CDRs comprise or consist of sequences selected from the sequences as set forth below: CDR-H1 is selected from the group consisting of LSYYYM (SEQ ID NO: 7), ISYSYI, (SEQ ID NO: 5), LSYSSM (SEQ ID NO: 2), ISSYSI (SEQ ID NO: 3), ISYSYI (SEQ ID NO: 5), IYSYSI (SEQ ID NO: 6), LSYYYM (SEQ ID NO: 7), FSSSSI (SEQ ID NO: 4), LYYYYI (SEQ ID NO: 8), LSYSSI (SEQ ID NO: 9), IYSYYI (SEQ ID NO: 10), LLYYSSM (SEQ ID NO: 122) and FSSSSI (SEQ ID NO: 4); CDR-H2 is selected from the group consisting of SIYPYYGYTY (SEQ ID NO: 11), SSSYYGYTY (SEQ ID NO: 12), SISSSYGYTY (SEQ ID NO: 13), SIYSSYGSTS (SEQ ID NO: 14), SIYSSYGYTY (SEQ ID NO: 15), SIYPYSSYTS (SEQ ID NO: 16), SIYSSYGYTY (SEQ ID NO: 15), SIYPSYGYTY (SEQ ID NO: 17), SISPYYGYTS (SEQ ID NO: 18), SISSSYGSTS (SEQ ID NO: 19), SIYSYYGYTY (SEQ ID NO: 20), SISSSYGYTY (SEQ ID NO: 13), SISSSYGYTY (SEQ ID NO: 13) SISSYYGYTS (SEQ ID NO: 53), and YISPYYGYTS (SEQ ID NO: 56); CDR-H3 is selected from the group consisting of HGAM (SEQ ID NO: 21), TVRGSKKPYFSGWAM (SEQ ID NO: 22), SSYYSSVSSSVYAL (SEQ ID NO: 23), TVRGSKKPYFSGWAM (SEQ ID NO: 22), HYSYFFYAM (SEQ ID NO: 24), YAVYFPGYYWGM (SEQ ID NO: 25), WSHVSGHYSGM (SEQ ID NO: 26), WGAYHSSGYGM (SEQ ID NO: 27), GGSGVSHYGSVYYSWWAL (SEQ ID NO: 28), AAPYYGYYYSYAM (SEQ ID NO: 29), SGYGWY AM (SEQ ID NO: 30), GYWAI (SEQ ID NO: 31), SYPAM (SEQ ID NO: 32), SWAM (SEQ ID NO: 33); YWAL (SEQ ID NO: 54), GWGSPASAGYYGL (SEQ ID NO: 57), SSYYSSVSSSVYAL (SEQ ID NO: 23), TVRGSKKPYFSGWAM (SEQ ID NO: 22) and TVRGSKKPYFSGWAM (SEQ ID NO: 22); CDR-L 1 is SVSSA (SEQ ID NO: 34); CDR-L2 is SASSLYS (SEQ ID NO: 35); and CDR-L3 is selected from the group consisting of AWGWGLF (SEQ ID NO: 36), VHYSPYSLI (SEQ ID NO: 37), YQYSGLI (SEQ ID NO: 38), FSHVSLI (SEQ ID NO: 39), ASYSPI (SEQ ID NO: 40), YHYYYLF (SEQ ID NO: 41), ASYAPI (SEQ ID NO:

42), SSSSPI (SEQ ID NO: 43), SSYSLI (SEQ ID NO: 44), GVSLI (SEQ ID NO: 45), YWFLI (SEQ ID NO: 46), PVGHYGYPI (SEQ ID NO: 47), SSYSPI (SEQ ID NO: 48), YWAYYSPI (SEQ ID NO: 49), VSYYPLI (SEQ ID NO: 51), SSYSLI (SEQ ID NO: 44), and VHYSPYSLI (SEQ ID NO: 37). In another embodiment the antibody comprises a heavy chain variable region comprising: i) a heavy chain amino acid sequence as set forth in Table 2; ii) an amino acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the heavy chain amino acid sequence as set forth in Table 2, wherein the CDR sequences are a CDR sequence set as set forth in Table 1, or iii) a conservatively substituted amino acid sequence of i) wherein the CDR sequences are a CDR sequence set as set forth in Table 1. In another embodiment the antibody comprises a light chain variable region comprising: i) a light chain amino acid sequence as set forth in Table 2, ii) an amino acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the light chain amino acid sequence as set forth in Table 2, wherein the CDR sequences are a CDR sequence set as set forth in Table 1, or iii) a conservatively substituted amino acid sequence of i) wherein the CDR sequences are a CDR sequence set as set forth in Table 1. In another embodiment the CDR sequences are a full CDR sequence set selected from the antibodies identified in Table 1. In another embodiment the antibody cross-reacts with LRP6. In another embodiment the CDR sequences comprise a light chain or a heavy chain CDR sequence set selected from the antibodies identified in Table 1. In another embodiment the antibody specifically binds LRP5. In another embodiment the CDR sequences are a CDR sequence set of an antibody selected from antibodies LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6. In another embodiment the antibody blocks binding of a Wnt ligand to a Wnt3a binding site of LRP5. In another embodiment the antibody binding of a Wnt ligand to a non-Wnt3a binding site of LRP5. In another embodiment the antibody the antibody is a monoclonal antibody. In another embodiment the antibody the antibody is a humanized antibody. In another embodiment the antibody the antibody is a single chain antibody. In another embodiment the antibody is an antibody binding fragment selected from Fab, Fab', F(ab')2, scFv, dsFv, ds-scFv, dimers, nanobodies, minibodies, in another embodiment the antibody is a bi-specific antibody that further binds to FZD receptor. In another embodiment antibody comprises a non-natural glycosylation pattern. In another embodiment antibody comprises a cysteine substitution or addition, e.g., in the constant region or a framework region.

[0010] In another aspect provided herein is an immunoconjugate comprising an antibody is provided herein, and a detectable label or cytotoxic agent. In another embodiment the

immuno conjugate comprises a cytotoxic agent selected from maytansinoid, auristatin, dolastatin, tubulysin, cryptophycin, pyrrolobenzodiazepine (PBD) dimer, indolinobenzodiazepine dimer, alpha-amanitin, trichothene, SN-38, duocarmycin, CC1065, calicheamincin, an enediyne antibioatic, taxane, doxorubicin derivatives, anthracycline and stereoisomers, azanofide, isosteres, analogs or derivatives thereof.

[0011] In another aspect provided herein is a nucleic acid molecule encoding and antibody has provided herein. In one embodiment one or more of the CDR sequences is/are encoded by a nucleic acid in Table 2. In another embodiment the antibody comprises a heavy chain variable region encoded by a nucleic acid comprising: i) a heavy chain nucleic acid sequence as set forth in Table 2; ii) a nucleotide sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the heavy chain nucleic acid sequence as set forth in Table 2, wherein the CDR sequences are a CDR sequence set as set forth in Table 1, or iii) a codon degenerate nucleic acid sequence of i) wherein the CDR sequences are a CDR sequence set as set forth in Table 1. In another embodiment the antibody comprises a light chain variable region encoded by a nucleic acid comprising: i) a light chain nucleic acid sequence as set forth in Table 2, ii) a nucleic acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the light chain nucleic acid sequence as set forth in Table 2, wherein the CDR sequences are a CDR sequence set as set forth in n Table 1, or iii) a codon degenerate nucleic acid sequence of i) wherein the CDR sequences are a CDR sequence set as set forth in Table 1.

[0012] In another aspect provided herein is a vector comprising an expression control sequence operatively linked to the nucleic acid as provided herein.

[0013] In another aspect provided herein is a host cell comprising recombinant nucleic acid molecule comprising an expression control sequence operatively linked to the nucleic acid has provided herein. In another embodiment the host cell is a Chinese Hamster Ovary (CHO) cell.

[0014] In another aspect provided herein is a host cell comprising a vector is provided herein.

[0015] In another aspect provided herein is a method for making an anti-LRP5 antibody comprising culturing a host cell as provided herein.

[0016] In another aspect provided herein is a composition comprising antibody, immunoconjugate, a nucleic acid molecule, a vector, or a host cell as provided herein, optionally with a suitable diluent. In one embodiment the composition comprises one or more

antibodies or immunoconjugates, optionally wherein the composition is a pharmaceutical composition.

[0017] In another aspect provided herein is a kit comprising an antibody, immunoconjugate, a nucleic acid molecule, a vector, or a host cell as provided herein.

[0018] In another aspect provided herein is a method of detecting LRP5 expression, the method comprising contacting a sample comprising one or more cells with one or more antibody or immunoconjugate as provided herein under conditions permissive for forming an antibody:cell complex and detecting the presence of any antibody complex. In one embodiment the detection is by immunofluorescence. In another embodiment the detection is by flow cytometry. In another embodiment the method is for detecting LRP4 expression and the antibody or immunoconjugate comprises a CDR sequence set corresponding to an antibody selected from LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[0019] In another aspect provided herein is a method of inhibiting Wnt ligand binding to an LRP5 receptor, disrupting a Wnt signaling pathway, inhibiting Wnt-induced transcriptional activity, inhibiting activation of disheveled, promoting preservation of the beta-catenin destruction complex, promoting accumulation of beta-catenin or inhibiting growth of a cell, the method comprising contacting a cell expressing a LRP5 receptor with an antibody or immunoconjugate as provided herein. In yet another aspect, provided is an antibody or immunoconjugate as provided herein for use in inhibiting Wnt ligand binding to an LRP5 receptor, disrupting a Wnt signaling pathway, inhibiting Wnt-induced transcriptional activity, inhibiting activation of disheveled, promoting preservation of the beta-catenin destruction complex, promoting accumulation of beta-catenin or inhibiting growth of a cell. In a further aspect, provided is a use of an antibody or immunoconjugate as provided herein for inhibiting Wnt ligand binding to an LRP5 receptor, disrupting a Wnt signaling pathway, inhibiting Wnt-induced transcriptional activity, inhibiting activation of disheveled, promoting preservation of the beta-catenin destruction complex, promoting accumulation of beta-catenin or inhibiting growth of a cell. In yet another aspect, provided is a use of an antibody or immunoconjugate as provided herein in the manufacture of a medicament for inhibiting Wnt ligand binding to an LRP5 receptor, disrupting a Wnt signaling pathway, inhibiting Wnt-induced transcriptional activity, inhibiting activation of disheveled, promoting preservation of the beta-catenin destruction complex, promoting accumulation of beta-catenin or inhibiting growth of a cell. In one embodiment the antibody or immunoconjugate blocks binding of a Wnt ligand to a Wnt3a binding site of LRP5. In another embodiment the antibody or immunoconjugate blocks binding of a Wnt ligand to a non-Wnt3a binding site of LRP5. In another embodiment the antibody or

immunoconjugate comprises a CDR sequence set corresponding to an antibody selected LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[0020] In another embodiment provided herein is a method of treating cancer in a subject in need thereof comprising administering to the subject an effective amount of a pharmaceutical composition comprising an antibody or an immunoconjugate as provided herein. In another embodiment is a pharmaceutical composition comprising an antibody or an immunoconjugate as provided herein for use in treating cancer in a subject in need thereof. In a further embodiment is a use of a pharmaceutical composition comprising an antibody or an immunoconjugate as provided herein for treating cancer in a subject in need thereof. In yet another embodiment is a use of a pharmaceutical composition comprising an antibody or an immunoconjugate as provided herein in the manufacture of a medicament for treating cancer in a subject in need thereof. In one embodiment the cancer is selected from colon, lung, breast ovarian, endometrial, pancreas, stomach, liver, adrenocortical carcinoma and osteoblastoma cancer cells. In another embodiment the cancer is selected from acute myeloid leukemia, prostate cancer, glioblastoma, bladder cancer and cervical cancer. In another embodiment the method comprises administering to the subject first and second antibodies or antibody conjugates as provided herein, wherein the first blocks binding of a Wnt ligand to a Wnt3a binding site of LRP5, and the second blocks binding of a Wnt ligand to a non-Wnt3a binding site of LRP5. In another embodiment the first antibody or immunoconjugate comprises a CDR sequence set selected from antibodies of Epitope Group 2. In another embodiment the antibody or immunoconjugate that specifically binds LRP5 in at least one assay, and inhibits Wnt3a- induced signaling in at least one assay, optionally wherein the antibody or immunoconjugate is the antibody or immunoconjugate has provided herein. In another embodiment the antibody or immunoconjugate comprises a CDR sequence set corresponding to an antibody selected from LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[0021] In another aspect provided herein is a method of potentiating the signaling activity of Wnt-ligand binding to a Wnt3a binding site of LRP5 comprising contacting a cell expressing LRP5 with an antibody that blocks binding of Wnt ligands to the non-Wnt3a binding site of LRP5. In one embodiment the method is performed *in vitro*. In another embodiment the method is performed *in vivo*.

[0022] In another aspect provided herein is a method of potentiating the signaling activity of Wnt-ligand binding to a non-Wnt3a binding site of LRP5 by contacting a cell expressing

LRP5 with an antibody that blocks binding of Wnt ligands to the Wnt3a binding site of LRP5. In one embodiment the method is performed *in vitro*. In another embodiment the method is performed *in vivo*.

BRIEF DESCRIPTION OF THE DRAWINGS

[0023] The accompanying drawings, which are incorporated herein and form a part of the specification, illustrate exemplary embodiments and, together with the description, further serve to enable a person skilled in the pertinent art to make and use these embodiments and others that will be apparent to those skilled in the art. The invention will be more particularly described in conjunction with the following drawings wherein:

[0024] **FIGs. 1A and 1B** show sixteen (16) anti-LRP antibodies. (A) The amino acid sequences of the complementarity determining regions (CDR) of the heavy (H) and the light (L) chains are presented. The antibodies were grouped into unique epitopes as determined by competitive ELISAs. (B) Single-point ELISAs were performed on 96-well Maxisorb immune plates coated with ECD's of mouse LRP5, mouse LRP6 and human LRP6 chimeras. The plates were incubated with the purified Fab or IgG1 at the concentrations indicated followed by incubation with horseradish peroxidase (HRP)-conjugated anti-kappa antibody. The wells were washed eight times followed by incubation with 3,3',5,5'-tetramethylbenzidine/H_2O_2 peroxidase (TMB) substrate for 5-10 min. The reaction was stopped by adding 1M H_3PO_4 and the absorbance was measured spectrometrically at 450 nm in a microtiter plate reader. FIG. 1A discloses "SSVA" as SEQ ID NO: 34, and "SASSLYS" as SEQ ID NO: 35, as well as SEQ ID NOS 36-49, 51, 44, 2-4, 4, 5-7, 5, 4, 5, 8-10, 4, 52, 55, 11-16, 15, 15, 13, 17-20, 13, 53, 56, 21-23, 22, 24-33, 54, 57, respectively, in order of columns.

[0025] **FIG. 2** shows an analysis of IgG1 binding to cell-surface LRP5 by flow cytometry. LRP5 IgGs (5 µg/ml) were tested for binding to the NSCLS cancer cell line, H23. Binding of the anti-LRP5 IgG1 proteins was detected using an Alexa488-conjugated secondary antibody against $F(ab')_2$. The stained anti-LRP5 population is shown in green in the secondary only state population is shown in filled blue.

[0026] **FIGs. 3A-3D** show the effect of LRP5 IgG's on transcriptional activity in cancer cells. TCF/LEF reporter assays were performed using the TOPflash (firefly luciferase gene) in (A) MDAMB231, (B) T407D, (C) U2OS and (D) H23 cells. Cells were seeded in white-walled, white bottom-96 well plates in treated with the IgG at the indicated concentration for one hour prior to stimulation with conditioned media. The cells were lysed 16-20 H post stimulation with conditioned media and reporter activity was assessed by measuring the luminescence signal generated by the addition of the firefly luminescence reagent. The values are normalized to the signal observed in cells treated with the negative control antibody, anti-

MBP, and stimulated with ConCM. The data presented is representative of three independent experiments where each condition is an average of three replicates.

[0027] **FIGs. 4A** and **4B** show the effect of LRP5-G2 and LRP5-G10 and Fab on transcriptional activity in cancer cells. TCF/LEF reporter assays were performed using the TOPflash (firefly luciferase gene) in H23 cells pretreated with increasing concentrations of (A) anti-LRP IgG1 or (B) anti-LRP5 Fab. The basal reporter activity was assessed as previously described for Figure 3.

[0028] **FIGs. 5A** and **5B** show the effect of LRP5 IgG1's on β-catenin signaling. The H23 cell line was pretreated with LRP5 IgG's (100 mM) for the time points indicated prior to stimulation with conditioned media. (A) Total whole cell lysates or (B) membrane and cytosolic fractions were generated and separated by SDS/PAGE. The PVDF membrane was cut and immunoblotted for phosphorylated LRP6, LRP5, LRP6, Axin1, Dvl3 and β-catenin. Actin was used as a loading control for whole cell lysates while Na/K ATPase and GAPDH were used as loading control for the membrane and cytosolic fractions, respectively.

[0029] **FIG. 6** shows multi-point competitive ELISA. Dose response curves and the non-linear regression plots are for LRP 5 antibodies.

[0030] **FIG. 7** shows an analysis of IgG1 binding to cell-surface LRP5 by flow cytometry. LRP5 IgG's (5 μg/ml) were tested for binding to the breast cancer cell line, MDAMB231. Binding of the anti-LRP5 IgG1 proteins was detected using an Alexa488-conjugated secondary antibody against $F(ab')_2$. The stained anti-LRP5 population is shown in green in the secondary only state population is shown in filled blue.

[0031] **FIG. 8** shows an analysis of IgG1 binding to cell-surface LRP5 by flow cytometry. LRP5 IgG's (5 μg/ml) were tested for binding to the breast cancer cell line, T47D. Binding of the anti-LRP5 IgG1 proteins was detected using an Alexa488-conjugated secondary antibody against $F(ab')_2$. The stained anti-LRP5 population is shown in green in the secondary only state population is shown in filled blue.

[0032] **FIG. 9** shows a diagram of the Wnt canonical signaling pathway.

DETAILED DESCRIPTION

I. Definitions

[0033] Unless otherwise defined, scientific and technical terms used in connection with the present disclosure shall have the meanings that are commonly understood by those of ordinary skill in the art. Further, unless otherwise required by context, singular terms shall include pluralities and plural terms shall include the singular. For example, the term "a cell"

includes a single cell as well as a plurality or population of cells. Generally, nomenclatures utilized in connection with, and techniques of, cell and tissue culture, molecular biology, and protein and oligonucleotide or polynucleotide chemistry and hybridization described herein are those well-known and commonly used in the art (see, e.g., Green and Sambrook, 2012).

[0034] As used herein, the term “polypeptide” refers to a molecule having a sequence of natural and/or unnatural amino acids connected through peptide bonds. The term “peptide” refers to a short polypeptide, typically no more than 30 amino acids long. The amino acid sequence of a polypeptide is referred to as its “primary structure.” The term “protein” refers to a polypeptide having a secondary, tertiary and/or quaternary structure, e.g., structures stabilized by hydrogen bonds, relationships between secondary structures and structures formed of more than one protein. Proteins can be further modified by other attached moieties such as carbohydrate (glycoproteins), lipids (lipoproteins) phosphate groups (phosphoproteins) and the like.

[0035] As used herein, an amino acid sequence “consists of” only the amino acids in that sequence.

[0036] As used herein, a first amino acid sequence “consists essentially of” a second amino acid sequence if the first amino acid sequence (1) comprises the second amino sequence and (2) is no more than 1, no more than 2 or no more than 3 amino acids longer than the second amino acid sequence.

[0037] As used herein, a first amino acid sequence is a “fragment” of a second amino acid sequence if the second amino acid sequence comprises the first amino acid sequence. In certain embodiments, a first amino acid sequence that is a fragment of a second amino acid sequence may have no more than any of 1, 2, 3, 4, 5, 6, 7, 8, 9 or 10 fewer amino acids than the second amino acid sequence.

[0038] As used herein, a “functional equivalent” of a reference amino acid sequence is a sequence that is not identical to the reference sequence, but that contains minor alterations such as, for example, insertion, deletion or substitution of one or a few amino acids. A functionally equivalent sequence retains the function (*e.g.*, immunogenicity) of the reference sequence to which it is equivalent. If a functionally equivalent amino acid sequence contains substitution of one or more amino acids with respect to the reference sequence, these will generally be conservative amino acid substitutions.

[0039] As used herein, a "conservative amino acid substitution" is one in which one amino acid residue is replaced with another amino acid residue without abolishing the protein's desired properties. Suitable conservative amino acid substitutions can be made by substituting amino acids with similar hydrophobicity, polarity, and R-chain length for one another. See,

e.g., Watson, *et al.*, “Molecular Biology of the Gene,” 4th Edition, 1987, The Benjamin/Cummings Pub. Co., Menlo Park, CA, p. 224. Examples of conservative amino acid substitution include the following (Note, some categories are not mutually exclusive):

Conservative Substitutions	
Type of Amino Acid	Substitutable Amino Acids
Hydrophilic	Ala, Pro, Gly, Glu, Asp, Gln, Asn, Ser, Thr
Sulphydryl	Cys
Aliphatic (non-polar, hydrophobic)	Ala, Val, Ile, Leu, Met, Gly, Pro
Basic	Lys, Arg, His
Aromatic	Phe, Tyr, Trp

[0040] As used herein, the term "substantially identical" refers to identity between a first amino acid sequence that contains a sufficient or minimum number of amino acid residues that are i) identical to, or ii) conservative substitutions of aligned amino acid residues in a second amino acid sequence such that the first and second amino acid sequences have a common structural domain and/or common functional activity and/or common immunogenicity. For example, amino acid sequences that contain a common structural or antigenic domain having at least about 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98% or 99% identity are termed sufficiently or substantially identical. In the context of nucleotide sequence, the term "substantially identical" is used herein to refer to a first nucleic acid sequence that contains a sufficient or minimum number of nucleotides that are identical to aligned nucleotides in a second nucleic acid sequence such that the first and second nucleotide sequences encode a polypeptide having common functional activity, or encode a common structural polypeptide domain or a common functional polypeptide activity, or encode polypeptides having the same immunogenic properties.

[0041] As used herein, the terms “antigen,” “immunogen,” and “antibody target,” refer to a molecule, compound, or complex that is recognized by an antibody, *i.e.*, can be bound by the antibody. The term can refer to any molecule that can be recognized by an antibody, *e.g.*, a polypeptide, polynucleotide, carbohydrate, lipid, chemical moiety, or combinations thereof (*e.g.*, phosphorylated or glycosylated polypeptides, etc.). One of skill will understand that the term does not indicate that the molecule is immunogenic in every context, but simply indicates that it can be targeted by an antibody.

[0042] As used herein, the term “epitope” refers to the localized site on an antigen that is recognized and bound by an antibody. Epitopes can include a few amino acids or portions of a few amino acids, *e.g.*, 5 or 6, or more, *e.g.*, 20 or more amino acids, or portions of those amino acids. In some cases, the epitope includes non-protein components, *e.g.*, from a carbohydrate, nucleic acid, or lipid. In some cases, the epitope is a three-dimensional moiety. Thus, for example, where the target is a protein, the epitope can be comprised of consecutive amino acids, or amino acids from different parts of the protein that are brought into proximity by protein folding (*e.g.*, a discontinuous epitope).

[0043] As used herein, the term “antibody” refers to an immunoglobulin that recognizes and specifically binds to a one or more target antigen(s), such as a protein, polypeptide, peptide, carbohydrate, polynucleotide, lipid or combinations thereof. This binding occurs through at least one antigen recognition site within the variable region of the immunoglobulin at one or more epitopes on the antigen. The variable region is most critical in binding specificity and affinity. As used herein, the term “antibody” encompasses intact polyclonal antibodies, intact monoclonal antibodies, antibody fragments, single chain Fv (scFv) mutants, multispecific antibodies, chimeric antibodies, humanized antibodies, human antibodies, hybrid antibodies, fusion proteins and any other immunoglobulin molecule comprising an antigen recognition site so long as the antibody exhibit the desired biological activity. Antibodies can be of (i) any of the five major classes of immunoglobulins, based on the identity of their heavy-chain constant domains – alpha (IgA), delta (IgD), epsilon (IgE), gamma (IgG) and mu (IgM), or (ii) subclasses (isotypes) thereof (E.g., IgG1, IgG2, IgG3, IgG4, IgA1 and IgA2). The light chains can be either lambda or kappa. Antibodies can be naked or conjugated to other molecules such as toxins, drugs, radioisotopes, chemotherapeutic agents, etc.

[0044] In one embodiment, an “intact antibody” comprises a tetramer composed of two identical pairs of polypeptide chains, each pair having one “light” (about 25 kD) and one “heavy” chain (about 50-70 kD). The heavy chain and light chains are connected through covalent and non-covalent bonds (e.g., disulfide linkage) that vary in number and amount between the various immunoglobulin classes. In one aspect, each chain comprises a variable region and a constant region. The antigen recognition site of the variable region is composed of hypervariable regions or complementarity determining regions (CDRs) and frameworks regions. The framework regions typically do not come into contact with the antigen but provide structural support for the CDRs. The constant region interacts with other immune cells of the body. Between the constant and variable region (IgG, IgD, IgA only but not IgM or IgE) is the hinge region in the center between the two heavy chains that provides flexibility to articulate antigen binding.

[0045] The following are a non-exhaustive list of different antibody forms, all retaining antigen binding activity:

(1) whole immunoglobulins (also referred to as “intact” antibodies) (two light chains and two heavy chains, e.g., a tetramer).

(2) an immunoglobulin polypeptide (a light chain or a heavy chain).

(3) an antibody fragment, such as Fv (a monovalent or bi-valent variable region fragment, and can encompass only the variable regions (*e.g.*, V_L and/or V_H), Fab (V_LC_L V_HC_H), F(ab')2, Fv (V_LV_H), scFv (single chain Fv) (a polypeptide comprising a V_L and V_H joined by a linker, *e.g.*, a peptide linker), (scFv)2, sc(Fv)2, bispecific sc(Fv)2, bispecific (scFv)2, minibody (sc(FV)2 fused to CH3 domain), triabody is trivalent sc(Fv)3 or trispecific sc(Fv)3

(4) a multivalent antibody (an antibody comprising binding regions that bind two different epitopes or proteins, e.g., “scorpion” antibody.

(5) a fusion protein comprising a binding portion of an immunoglobulin fused to another amino acid sequence (such as a fluorescent protein).

[0046] As used herein, the term “antibody fragment" refers to a part or portion of an antibody or antibody chain comprising fewer amino acid residues than an intact or complete antibody or antibody chain and which binds the antigen or competes with intact antibody. Fragments can be obtained via chemical or enzymatic treatment of an intact or complete antibody or antibody chain. Fragments can also be obtained by recombinant means. For example, F(ab')2 fragments can be generated by treating the antibody with pepsin. The resulting F(ab')2 fragment can be treated to reduce disulfide bridges to produce Fab' fragments. Papain digestion can lead to the formation of Fab fragments. Fab, Fab' and F(ab')2, scFv, dsFv, ds-scFv, dimers, minibodies, diabodies, bispecific antibody fragments and other fragments can also be constructed by recombinant expression techniques.

[0047] While various antibody fragments are defined in terms of products of the digestion of an intact antibody, one of skill will appreciate that such fragments may also be synthesized *de novo* chemically or constructed and expressed using recombinant DNA methodology.

[0048] A single chain Fv (scFv) refers to a polypeptide comprising a V_L and V_H joined by a linker, *e.g.*, a peptide linker. ScFvs can also be used to form tandem (or di-valent) scFvs or diabodies. Production and properties of tandem scFvs and diabodies are described, *e.g.*, in Asano *et al.* (2011) *J Biol. Chem.* 286:1812; Kenanova *et al.* (2010) *Prot Eng Design Sel* 23:789; Asano *et al.* (2008) *Prot Eng Design Sel* 21:597.

[0049] Antibody fragments further include Fd (the portion of the heavy chain included in the Fab fragment) and single domain antibodies. A single domain antibody (sdAb) is a variable domain of either a heavy chain or a light chain, produced by recombinant methods.

[0050] The phrase “CDR sequence set” as used herein refers to the 3 heavy chain and/or 3 light chain CDRs of a particular antibody described herein. A “light chain” CDR sequence set refers to the light chain CDR sequences. A “heavy chain” CDR sequence set refers to the heavy chain CDR sequences. A “full” CDR sequence set refers to both heavy chain and light chain CDR sequences. For example, for antibody LRP5-A7, as shown in Table 1, the full CDR sequence set comprises or consists of SVSSA (SEQ ID NO: 34) (CDR L1), SASSLYS (SEQ ID NO: 35) (CDR L2) AWGWGLF (SEQ ID NO: 36) (CDR L3), LYSSSM (SEQ ID NO: 50) (CDR H1), SIYPYYGYTY (SEQ ID NO: 11) (CDR H2) AND HGAM (SEQ ID NO: 21) (CDR H3). The CDR sequence for each CDR can, for example, comprise, consist essentially of, or consist of the CDR in Table 1. CDRs are predicted based on IMGT sequence alignment.

[0051] As used herein, the term “monoclonal antibody” refers to a clonal preparation or composition of antibodies with a single binding specificity and affinity for a given epitope on an antigen (“monoclonal antibody composition”). A “polyclonal antibody” refers to a preparation or composition of antibodies that are raised against a single antigen, but with different binding specificities and affinities (“polyclonal antibody composition”).

[0052] As used herein, the term “chimeric antibody” refers to an antibody having amino acid sequences derived from two or more species. In one embodiment, the variable region of both light and heavy chains correspond to the variable region of antibodies derived from one species of mammal (e.g., mouse, rat, rabbit, etc.) with the desired specificity, affinity and capability, while the constant region are homologous the sequence derived from another species (typically in the subject receiving the therapy, e.g., human) to avoid eliciting an immune response.

[0053] As used herein, the term “humanized antibody” refers to a chimeric antibody in which the CDRs, obtained from the VH and VL regions of a non-human antibody having the desired specificity, affinity and capability are grafted to a human framework sequence. In one embodiment, the framework residues of the humanized antibody is modified to refine and optimize the antibody specificity, affinity and capability. Humanization, i.e., substitution of non-human CDR sequences for the corresponding sequences of a human antibody, can be performed following the methods described in, e.g., U.S. Patent Nos. 5,545,806; 5,569,825; 5,633,425; 5,661,016; Riechmann et al., Nature 332:323-327 (1988); Marks et al., Bio/Technology 10:779-783 (1992); Morrison, Nature 368:812-13 (1994); Fishwild et al., Nature Biotechnology 14:845-51 (1996).

[0054] As used herein, the term “human antibody” refers to an antibody produced by a human or an antibody having an amino acid sequence corresponding thereto made by any technique known in the art.

[0055] As used herein, the term “hybrid antibody” refers to antibody in which pairs of heavy and light chains form antibodies with different antigenic determinant regions are assembled together so that two different epitopes or two different antigens can be recognized and bound by the resulting tetramer. Hybrid antibodies can be bispecific (binding 2 distinct antigens or epitopes) or multispecific (> 1 distinct antigen or epitope).

[0056] As used herein, an antibody is “monospecific” if all of its antigen binding sites bind to the same epitope.

[0057] As used herein, an antibody is “bispecific” if it has at least two different antigen binding sites which each bind to a different epitope or antigen.

[0058] As used herein, an antibody is “polyvalent” if it has more than one antigen binding site. For example, an antibody that is tetravalent has four antigen binding sites.

[0059] The specificity of the binding can be defined in terms of the comparative dissociation constants (Kd) of the antibody (or other targeting moiety) for target, as compared to the dissociation constant with respect to the antibody and other materials in the environment or unrelated molecules in general. A larger (higher) K_d is a K_d that describes a lower affinity interaction. Conversely a smaller (lower) K_d is a K_d that describes a higher affinity interaction or tighter binding. By way of example only, the K_d for an antibody specifically binding to a target may be femtomolar, picomolar, nanomolar, or micromolar and the K_d for the antibody binding to unrelated material may be millimolar or higher. Binding affinity can be in the micromolar range (kD = 10^{-4} to 10^{-6}), nanomole range (kD = 10^{-7} M to 10^{-9} M), picomole range (kD = 10^{-10} M to 10^{-12} M), or femtomole range (kD = 10^{-13} M to 10^{-15} M).

[0060] As used herein, an antibody “binds” or “recognizes” an antigen or epitope if it binds the antigen or epitope with a Kd of less than 10^{-4}M (i.e., in the micromolar range). The term “binds” with respect to a cell type (e.g., an antibody that binds cancer cells), typically indicates that an agent binds a majority of the cells in a pure population of those cells. For example, an antibody that binds a given cell type typically binds to at least 2/3 of the cells in a population of the indicated cells (e.g., 67, 75, 80, 85, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, or 100%). In some cases, binding to a polypeptide can be assayed by comparing binding of the antibody to a cell that presents the polypeptide to binding (or lack thereof) of the antibody to a cell that does not express the polypeptide. One of skill will recognize that some variability will arise depending on the method and/or threshold of determining binding. Affinity of an antibody for a target can be determined according to methods known in the art, *e.g.*, as reviewed in Ernst

et al. Determination of Equilibrium Dissociation Constants, Therapeutic Monoclonal Antibodies (Wiley & Sons ed. 2009).

[0061] As used herein, the term “greater affinity” as used herein refers to a relative degree of antibody binding where an antibody X binds to target Y more strongly (K_{on}) and/or with a smaller dissociation constant (K_{off}) than to target Z, and in this context antibody X has a greater affinity for target Y than for Z. Likewise, the term "lesser affinity" herein refers to a degree of antibody binding where an antibody X binds to target Y less strongly and/or with a larger dissociation constant than to target Z, and in this context antibody X has a lesser affinity for target Y than for Z. The affinity of binding between an antibody and its target antigen, can be expressed as K_A equal to $1/K_D$ where K_D is equal to k_{on}/k_{off}. The k_{on} and k_{off} values can be measured using surface plasmon resonance technology, for example, using a Molecular Affinity Screening System (MASS-1) (Sierra Sensors GmbH, Hamburg, Germany). An antagonist or blocking antibody is an antibody that partially or fully blocks inhibits or neutralizes a biological activity related to the target antigen relative to the activity under similar physiological conditions when the antibody is not present. Antagonists can be competitive, non-competitive or irreversible. A competitive antagonist is a substance that binds to a natural ligand or receptor at the same site as the natural ligand-receptor interaction or binds allosterically in a manner that induces a change to prevent normal binding. A non-competitive antagonist binds at a different site than the natural ligand-receptor interaction, but lower the KD or signal resulting from the interaction. An irreversible inhibitor causes covalent modifications to the receptor preventing any subsequent binding.

[0062] As used herein, the term “avidity” refers to the overall stability of the binding complex between the antibody and the target antigen. It is governed by three factors, (i) the intrinsic affinity of the antibody for the antigen, (2) the valency of the antibody, and (3) the geometric arrangement of the interacting components. Affinity is the strength of the interaction between the antibody and a single target, whereas avidity is an accumulated strength of multiple affinities. In one embodiment, the antibodies disclosed herein are divalent.

[0063] As used herein, an antibody “preferentially binds” binds a first antigen relative to a second antigen if it binds the first antigen with greater affinity than it does the second antigen. Preferential binding can be at least any of 2-fold, 5-fold, 9-fold, 10-fold, 20-fold, 30-fold, 40-fold, 50-fold, 100-fold, 500-fold or 1000-fold greater affinity. So, for example, an antibody preferentially binds LRP5 relative to LRP6 if it binds LRP5 with greater affinity than it binds LRP6.

[0064] As used herein, an antibody “specifically binds” or is “specific for” a target antigen or target group of antigens if it binds the target antigen or each member of the target group of

antigens with an affinity of at least any of 1×10^{-6} M, 1×10^{-7}M, 1×10^{-8}M, 1×10^{-9} M, 1×10^{-10} M, 1×10^{-11} M, 1×10^{-12} M, and, for example, binds to the target antigen or each member of the target group of antigens with an affinity that is at least two-fold greater than its affinity for non-target antigens to which it is being compared. Typically, specific binding is characterized by binding the antigen with sufficient affinity that the antibody is useful as a diagnostic to detect the antigen or epitope and/or as a therapeutic agent in targeting the antigen or epitope.

[0065] As used herein, and antibody "blocks" or "antagonizes" the binding of a ligand to receptor when it competitively reduces or prevents interaction all of the ligand with the receptor. In an embodiment, the measured level of reduction can be at least any of 5%, 10%, 25%, 50%, 80%, 90%, 95%, 97.5%, 99%, 99.5%, 99.9% of a control (e.g., untreated) cell. For example, an antibody that antagonizes or blocks the binding of a Wnt ligand to an LRP5 receptor competitively reduces or prevents the interaction of a Wnt protein with an LRP5 receptor. This results in attenuation or blocking of a downstream signaling event associated with Wnt signaling. This includes, for example, activation of disheveled, dissolution of the β-catenin destructive complex, lower cytosolic levels of β-catenin, and/or lower activity of TCF/LEF-mediated transcription.

[0066] The term "captures" with respect to an antibody target (e.g., antigen, analyte, immune complex), typically indicates that an antibody binds a majority of the antibody targets in a pure population (assuming appropriate molar ratios). For example, an antibody that binds a given antibody target typically binds to at least 2/3 of the antibody targets in a solution (e.g., at least any of 67, 75, 80, 85, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, or 100%). One of skill will recognize that some variability will arise depending on the method and/or threshold of determining binding.

[0067] The term "conjugate" refers to a first molecule, e.g., an antibody (an "immunoconjugate"), chemically coupled with a moiety, such as a detectable label or a biologically active moiety, such as a drug, toxin or chemotherapeutic or cytotoxic agent. Accordingly, this disclosure contemplates antibodies conjugated with one or more moieties. Furthermore, an antibody can be "conjugated antibody" or a "non-conjugated antibody" (that is, not conjugated with a moiety.

[0068] As used herein, the term "antibody-drug conjugate" or ("ADC") refers to an antibody conjugated with a drug. Typically, conjugation involves covalent binding through a linker.

[0069] As used herein, the term "labeled" molecule (*e.g.*, nucleic acid, protein, or antibody) refers to a molecule that is bound to a detectable label, either covalently, through a linker or a chemical bond, or noncovalently, through ionic, van der Waals, electrostatic, or hydrogen

bonds, such that the presence of the molecule may be detected by detecting the presence of the detectable label bound to the molecule.

[0070] As used herein, the term “detectable label” refers to a composition detectable by spectroscopic, photochemical, biochemical, immunochemical, chemical, or other physical means. Examples of detectable labels are described herein and include, without limitation, colorimetric, fluorescent, chemiluminescent, enzymatic, and radioactive labels. For the purposes of the present disclosure, a detectable label can also be a moiety that does not itself produce a signal (*e.g.*, biotin), but that binds to a second moiety that is able to produce a signal (*e.g.*, labeled avidin).

[0071] The term “cross-linked” with respect to an antibody refers to attachment of the antibody to a solid or semisolid matrix (*e.g.*, sepharose, beads, microtiter plate), or to another protein or antibody. For example, an antibody can be multimerized to create an antibody complex with multiple (more than 2) antigen-binding sites. The antibody can be multimerized by expressing the antibody as a high-valency isotype (*e.g.*, IgA or IgM, which typically form complexes of 2 or 5 antibodies, respectively). Antibody multimerization can also be carried out by using a cross-linker comprising a reactive group capable of linking proteins (*e.g.*, carbodiimide, NHS esters, etc.). Methods and compositions for cross-linking an antibody to a matrix are described, *e.g.*, in the Abcam and New England Biolab catalogs and websites (available at abcam.com and neb.com). Cross-linker compounds with various reactive groups are described, *e.g.*, in Thermo Fisher Scientific catalog and website (available at piercenet.com).

[0072] As used herein, the term “immunoassay” refers to a method for detecting an analyte by detecting binding between an antibody that recognizes the analyte and the analyte.

[0073] As used herein, the term “expression construct” refers to a polynucleotide comprising an expression control sequence operatively linked with a heterologous nucleotide sequence (i.e., a sequence to which the expression control sequence is not normally connected to in nature) that is to be the subject of expression. As used herein, the term “expression vector” refers to a polynucleotide comprising an expression construct and sequences sufficient for replication in a host cell or insertion into a host chromosome. Plasmids and viruses are examples of expression vectors. As used herein, the term “expression control sequence” refers to a nucleotide sequence that regulates transcription and/or translation of a nucleotide sequence operatively linked thereto. Expression control sequences include promoters, enhancers, repressors (transcription regulatory sequences) and ribosome binding sites (translation regulatory sequences).

[0074] The term "vector" as used herein comprises any intermediary vehicle for a nucleic acid molecule which enables said nucleic acid molecule, for example, to be introduced into prokaryotic and/or eukaryotic cells and/or integrated into a genome, and include plasmids, phagemids, bacteriophages or viral vectors such as retroviral based vectors, Adeno Associated viral vectors and the like. The term "plasmid" as used herein generally refers to a construct of extrachromosomal genetic material, usually a circular DNA duplex, which can replicate independently of chromosomal DNA.

[0075] As used herein, a nucleotide sequence is “operatively linked” with an expression control sequence when the expression control sequence functions in a cell to regulate transcription of the nucleotide sequence. This includes promoting transcription of the nucleotide sequence through an interaction between a polymerase and a promoter.

[0076] As used herein, a “host cell” refers to a recombinant cell comprising an expression construct.

[0077] As used herein, the term “biological sample” refers to a sample containing cells (e.g., tumor cells) or biological molecules derived from cells. A biological sample can be obtained from a subject, e.g., a patient, from an animal, such as an animal model, or from cultured cells, e.g., a cell line or cells removed from a patient and grown in culture for observation. A biological sample can comprise tissue and/or liquid. It can be obtained from any biological source including without limitation blood, a blood fraction (e.g., serum or plasma), cerebrospinal fluid (CSF), lymph, tears, saliva, sputum, buccal swab, milk, urine or feces. A biological sample can be a biopsy, such as a tissue biopsy, such as needle biopsy, fine needle biopsy, surgical biopsy, etc. The sample can comprise a tissue sample harboring a lesion or suspected lesion, although the biological sample may also be derived from another site, e.g., a site of suspected metastasis, a lymph node, or from the blood. A biological sample can be a fraction of a sample taken from a subject. An example of a tissue sample includes a brain tissue sample or a nerve tissue sample. Methods of obtaining such biological samples are known in the art including but not limited to standard blood retrieval procedures.

[0078] As used herein, the term “diagnosis” refers to a relative probability that a subject has a disorder such as cancer. Similarly, the term “prognosis” refers to a relative probability that a certain future outcome may occur in the subject. For example, in the context of the present disclosure, prognosis can refer to the likelihood that an individual will develop cancer, have recurrence, that the cancer will metastasize, that the cancer will be cured, or the likely severity of the disease (e.g., severity of symptoms, rate of functional decline, survival, etc.). The terms are not intended to be absolute, as will be appreciated by any one of skill in the field of medical diagnostics.

[0079] As used herein, the term terms “therapy,” “treatment,” “therapeutic intervention” and “amelioration” refer to any activity resulting in a reduction in the severity of symptoms. In the case of cancer, treatment can refer to, e.g., reducing tumor size, number of cancer cells, growth rate, metastatic activity, reducing cell death of non-cancer cells, reduced nausea and other chemotherapy or radiotherapy side effects, etc. The terms “treat” and “prevent” are not intended to be absolute terms. Treatment and prevention can refer to any delay in onset, amelioration of symptoms, improvement in patient survival, increase in survival time or rate, etc. Treatment and prevention can be complete (undetectable levels of neoplastic cells) or partial, such that fewer neoplastic cells are found in a patient than would have occurred without the present intervention. The effect of treatment can be compared to an individual or pool of individuals not receiving the treatment, or to the same patient prior to treatment or at a different time during treatment. In some aspects, the severity of disease is reduced by at least 10%, as compared, e.g., to the individual before administration or to a control individual not undergoing treatment. In some aspects, the severity of disease is reduced by at least 25%, 50%, 75%, 80%, or 90%, or in some cases, no longer detectable using standard diagnostic techniques.

[0080] As used herein, the terms “effective amount,” “effective dose,” and “therapeutically effective amount,” refer to an amount of an agent, such as an antibody or immunoconjugate, that is sufficient to generate a desired response, such as reduce or eliminate a sign or symptom of a condition or ameliorate a disorder. In some examples, an “effective amount” is one that treats (including prophylaxis) one or more symptoms and/or underlying causes of any of a disorder or disease and/or prevents progression of a disease. For example, for the given parameter, a therapeutically effective amount will show an increase or decrease of therapeutic effect at least any of 5%, 10%, 15%, 20%, 25%, 40%, 50%, 60%, 75%, 80%, 90%, or at least 100%. Therapeutic efficacy can also be expressed as “-fold” increase or decrease. For example, a therapeutically effective amount can have at least any of a 1.2-fold, 1.5-fold, 2-fold, 5-fold, or more effect over a control.

[0081] As used herein, the term “pharmaceutical composition” refers to a composition comprising a pharmaceutical compound (e.g., a drug) and a pharmaceutically acceptable carrier.

[0082] As used herein, the term “pharmaceutically acceptable” refers to a carrier that is compatible with the other ingredients of a pharmaceutical composition and can be safely administered to a subject. The term is used synonymously with “physiologically acceptable” and “pharmacologically acceptable”. Pharmaceutical compositions and techniques for their preparation and use are known to those of skill in the art in light of the present disclosure. For a detailed listing of suitable pharmacological compositions and techniques for their

administration one may refer to texts such as Remington's Pharmaceutical Sciences, 17th ed. 1985; Brunton et al., “Goodman and Gilman’s The Pharmacological Basis of Therapeutics,” McGraw-Hill, 2005; University of the Sciences in Philadelphia (eds.), “Remington: The Science and Practice of Pharmacy,” Lippincott Williams & Wilkins, 2005; and University of the Sciences in Philadelphia (eds.), “Remington: The Principles of Pharmacy Practice,” Lippincott Williams & Wilkins, 2008.

[0083] Pharmaceutically acceptable carriers will generally be sterile, at least for human use. A pharmaceutical composition will generally comprise agents for buffering and preservation in storage, and can include buffers and carriers for appropriate delivery, depending on the route of administration. Examples of pharmaceutically acceptable carriers include, without limitation, normal (0.9%) saline, phosphate-buffered saline (PBS) Hank’s balanced salt solution (HBSS) and multiple electrolyte solutions such as PlasmaLyte ATM (Baxter).

[0084] Acceptable carriers, excipients and/or stabilizers are nontoxic to recipients at the dosages and concentrations employed, and include buffers such as phosphate, citrate, and other organic acids; antioxidants including ascorbic acid, glutathione, cysteine, methionine and citric acid; preservatives (such as ethanol, benzyl alcohol, phenol, m-cresol, p-chlor-m-cresol, methyl or propyl parabens, benzalkonium chloride, or combinations thereof); amino acids such as arginine, glycine, ornithine, lysine, histidine, glutamic acid, aspartic acid, isoleucine, leucine, alanine, phenylalanine, tyrosine, tryptophan, methionine, serine, proline and combinations thereof; monosaccharides, disaccharides and other carbohydrates; low molecular weight (less than about 10 residues) polypeptides; proteins, such as gelatin or serum albumin; chelating agents such as EDTA; sugars such as trehalose, sucrose, lactose, glucose, mannose, maltose, galactose, fructose, sorbose, raffinose, glucosamine, N-methylglucosamine, galactosamine, and neuraminic acid; and/or non-ionic surfactants such as Tween, Pluronics, Triton-X, or polyethylene glycol (PEG).

[0085] The terms “dose” and “dosage” are used interchangeably herein. A dose refers to the amount of active ingredient given to an individual at each administration. For the present invention, the dose can refer to the concentration of the antibody or associated components, e.g., the amount of therapeutic agent or dosage of radiolabel. The dose will vary depending on a number of factors, including frequency of administration; size and tolerance of the individual; severity of the condition; risk of side effects; the route of administration; and the imaging modality of the detectable label (if present). One of skill in the art will recognize that the dose can be modified depending on the above factors or based on therapeutic progress. The term “dosage form” refers to the particular format of the pharmaceutical, and depends on

the route of administration. For example, a dosage form can be in a liquid, e.g., a saline solution for injection.

[0086] As used herein, the term "subject" refers to an individual animal. The term "patient" as used herein refers to a subject under the care or supervision of a health care provider such as a doctor or nurse. Subjects include mammals, such as humans and non-human primates, such as monkeys, as well as dogs, cats, horses, bovines, rabbits, rats, mice, goats, pigs, and other mammalian species. Subjects can also include avians. A patient can be an individual that is seeking treatment, monitoring, adjustment or modification of an existing therapeutic regimen, etc. The term "cancer subject" refers to an individual that has been diagnosed with cancer. Cancer patients can include individuals that have not received treatment, are currently receiving treatment, have had surgery, and those that have discontinued treatment.

[0087] In the context of treating cancer, a subject in need of treatment can refer to an individual that has cancer or a pre-cancerous condition, has had cancer and is at risk of recurrence, is suspected of having cancer, is undergoing standard treatment for cancer, such as radiotherapy or chemotherapy, etc.

[0088] "Cancer", "tumor," "transformed" and like terms include precancerous, neoplastic, transformed, and cancerous cells, and can refer to a solid tumor, or a non-solid cancer (see, e.g., Edge et al. AJCC Cancer Staging Manual (7th ed. 2009); Cibas and Ducatman Cytology: Diagnostic principles and clinical correlates (3rd ed. 2009)). Cancer includes both benign and malignant neoplasms (abnormal growth). "Transformation" refers to spontaneous or induced phenotypic changes, e.g., immortalization of cells, morphological changes, aberrant cell growth, reduced contact inhibition and anchorage, and/or malignancy (see, Freshney, Culture of Animal Cells a Manual of Basic Technique (3rd ed. 1994)). Although transformation can arise from infection with a transforming virus and incorporation of new genomic DNA, or uptake of exogenous DNA, it can also arise spontaneously or following exposure to a carcinogen.

[0089] The term "cancer" can refer to any cancer, including without limitation, leukemias, carcinomas, sarcomas, adenocarcinomas, lymphomas, solid and lymphoid cancers, etc. Examples of different types of cancer include, but are not limited to, lung cancer (e.g., non-small cell lung cancer or NSCLC), breast cancer, prostate cancer, colorectal cancer, bladder cancer, ovarian cancer, leukemia, liver cancer (i.e., hepatocarcinoma), renal cancer (i.e., renal cell carcinoma), thyroid cancer, pancreatic cancer, uterine cancer, cervical cancer, testicular cancer, esophageal cancer, stomach (gastric) cancer, kidney cancer, cancer of the central nervous system, skin cancer, glioblastoma and melanoma.

[0090] As used herein, a chemical entity, such as a polypeptide, is "substantially pure" if it is the predominant chemical entity of its kind (e.g., of polypeptides) in a composition. This

includes the chemical entity representing more than 50%, more than 80%, more than 90%, more than 95%, more than 98%, more than 99%, more than 99.5%, more than 99.9%, or more than 99.99% of the chemical entities of its kind in the composition.

[0091] The phrase "isolated antibody" refers to antibody produced *in vivo* or *in vitro* that has been removed from the source that produced the antibody, for example, an animal, hybridoma or other cell line (such as recombinant insect, yeast or bacterial cells that produce antibody).

[0092] “Substantially pure” or “isolated” means an object species is the predominant species present (i.e., on a molar basis, more abundant than any other individual macromolecular species in the composition), and a substantially purified fraction is a composition wherein the object species comprises at least about 50% (on a molar basis) of all macromolecular species present. Generally, a substantially pure composition means that about 80% to 90% or more of the macromolecular species present in the composition is the purified species of interest. The object species is purified to essential homogeneity (contaminant species cannot be detected in the composition by conventional detection methods) if the composition consists essentially of a single macromolecular species. Solvent species, small molecules (<500 Daltons), stabilizers (e.g., BSA), and elemental ion species are not considered macromolecular species for purposes of this definition.

[0093] The term "sequence identity" as used herein refers to the percentage of sequence identity between two polypeptide sequences or two nucleic acid sequences. To determine the percent identity of two amino acid sequences or of two nucleic acid sequences, the sequences are aligned for optimal comparison purposes (e.g., gaps can be introduced in the sequence of a first amino acid or nucleic acid sequence for optimal alignment with a second amino acid or nucleic acid sequence). The amino acid residues or nucleotides at corresponding amino acid positions or nucleotide positions are then compared. When a position in the first sequence is occupied by the same amino acid residue or nucleotide as the corresponding position in the second sequence, then the molecules are identical at that position. The percent identity between the two sequences is a function of the number of identical positions shared by the sequences (i.e., % identity=number of identical overlapping positions/total number of positions.times.100%). In one embodiment, the two sequences are the same length. The determination of percent identity between two sequences can also be accomplished using a mathematical algorithm. A preferred, non-limiting example of a mathematical algorithm utilized for the comparison of two sequences is the algorithm of Karlin and Altschul, 1990, Proc. Natl. Acad. Sci. U.S.A. 87:2264-2268, modified as in Karlin and Altschul, 1993, Proc. Natl. Acad. Sci. U.S.A. 90:5873-5877. Such an algorithm is incorporated into the NBLAST and XBLAST programs of Altschul et al., 1990, J. Mol. Biol. 215:403. BLAST nucleotide searches can be

performed with the NBLAST nucleotide program parameters set, e.g., for score=100, wordlength=12 to obtain nucleotide sequences homologous to a nucleic acid molecules of the present application. BLAST protein searches can be performed with the XBLAST program parameters set, e.g., to score-50, wordlength=3 to obtain amino acid sequences homologous to a protein molecule described herein. To obtain gapped alignments for comparison purposes, Gapped BLAST can be utilized as described in Altschul et al., 1997, Nucleic Acids Res. 25:3389-3402. Alternatively, PSI-BLAST can be used to perform an iterated search which detects distant relationships between molecules (Id.). When utilizing BLAST, Gapped BLAST, and PSI-Blast programs, the default parameters of the respective programs (e.g., of XBLAST and NBLAST) can be used (see, e.g., the NCBI website). Another preferred, non-limiting example of a mathematical algorithm utilized for the comparison of sequences is the algorithm of Myers and Miller, 1988, CABIOS 4:11-17. Such an algorithm is incorporated in the ALIGN program (version 2.0) which is part of the GCG sequence alignment software package. When utilizing the ALIGN program for comparing amino acid sequences, a PAM120 weight residue table, a gap length penalty of 12, and a gap penalty of 4 can be used. The percent identity between two sequences can be determined using techniques similar to those described above, with or without allowing gaps. In calculating percent identity, typically only exact matches are counted.

[0094] For antibodies, percentage sequence identities can be determined when antibody sequences maximally aligned by IMGT. After alignment, if a subject antibody region (e.g., the entire mature variable region of a heavy or light chain) is being compared with the same region of a reference antibody, the percentage sequence identity between the subject and reference antibody regions is the number of positions occupied by the same amino acid in both the subject and reference antibody region divided by the total number of aligned positions of the two regions, multiplied by 100 to convert to percentage.

[0095] Percent amino acid sequence identity may also be determined using the sequence comparison program NCBI-BLAST2 (Altschul et al., Nucleic Acids Res. 25:3389-3402 (1997)). The NCBI-BLAST2 sequence comparison program may be obtained from the National Institute of Health, Bethesda, Md. NCBI-BLAST2 uses several search parameters, wherein all of those search parameters are set to default values including, for example, unmask=yes, strand=all, expected occurrences=10, minimum low complexity length= 15/5, multi-pass e-value=0.01, constant for multi-pass=25, dropoff for final gapped alignment=25 and scoring matrix=BLOSUM62.

[0096] In situations where NCBI-BLAST2 is employed for amino acid sequence comparisons, the % amino acid sequence identity of a given amino acid sequence A to, with, or against a given amino acid sequence B (which can alternatively be phrased as a given

amino acid sequence A that has or comprises a certain % amino acid sequence identity to, with, or against a given amino acid sequence B) is calculated as follows:

100 times the fraction X/Y

[0097] where X is the number of amino acid residues scored as identical matches by the sequence alignment program NCBI-BLAST2 in that program's alignment of A and B, and where Y is the total number of amino acid residues in B. It will be appreciated that where the length of amino acid sequence A is not equal to the length of amino acid sequence B, the % amino acid sequence identity of A to B will not equal the % amino acid sequence identity of B to A. The term “nucleic acid sequence” as used herein refers to a sequence of nucleoside or nucleotide monomers consisting of naturally occurring bases, sugars and intersugar (backbone) linkages and includes cDNA. The term also includes modified or substituted sequences comprising non-naturally occurring monomers or portions thereof. The nucleic acid sequences of the present application may be deoxyribonucleic acid sequences (DNA) or ribonucleic acid sequences (RNA) and may include naturally occurring bases including adenine, guanine, cytosine, thymidine and uracil. The sequences may also contain modified bases. Examples of such modified bases include aza and deaza adenine, guanine, cytosine, thymidine and uracil; and xanthine and hypoxanthine. It is understood that polynucleotides comprising non-transcribable nucleotide bases may be useful as probes in, for example, hybridization assays. The nucleic acid can be either double stranded or single stranded, and represents the sense or antisense strand. Further, the term "nucleic acid" includes the complementary nucleic acid sequences as well as codon optimized or synonymous codon equivalents.

[0098] The term "isolated nucleic acid" as used herein refers to a nucleic acid substantially free of cellular material or culture medium when produced by recombinant DNA techniques, or chemical precursors, or other chemicals when chemically synthesized. An isolated nucleic acid is also substantially free of sequences that naturally flank the nucleic acid (i.e. sequences located at the 5' and 3' ends of the nucleic acid) from which the nucleic acid is derived.

[0099] By “at least moderately stringent hybridization conditions” it is meant that conditions are selected which promote selective hybridization between two complementary nucleic acid molecules in solution. Hybridization may occur to all or a portion of a nucleic acid sequence molecule. The hybridizing portion is typically at least 15 (e.g., 20, 25, 30, 40 or 50) nucleotides in length. Those skilled in the art will recognize that the stability of a nucleic acid duplex, or hybrids, is determined by the Tm, which in sodium containing buffers is a function of the sodium ion concentration and temperature (Tm = 81.5°C – 16.6 (Log10 [Na+]) + 0.41(%(G+C) – 600/l), or similar equation). Accordingly, the parameters in the wash conditions

that determine hybrid stability are sodium ion concentration and temperature. In order to identify molecules that are similar, but not identical, to a known nucleic acid molecule a 1% mismatch may be assumed to result in about a 1°C decrease in Tm, for example, if nucleic acid molecules are sought that have a >95% identity, the final wash temperature will be reduced by about 5°C. Based on these considerations those skilled in the art will be able to readily select appropriate hybridization conditions. In preferred embodiments, stringent hybridization conditions are selected. By way of example the following conditions may be employed to achieve stringent hybridization: hybridization at 5x sodium chloride/sodium citrate (SSC)/5x Denhardt's solution/1.0% SDS at Tm - 5°C based on the above equation, followed by a wash of 0.2x SSC/0.1% SDS at 60°C. Moderately stringent hybridization conditions include a washing step in 3x SSC at 42°C. It is understood, however, that equivalent stringencies may be achieved using alternative buffers, salts and temperatures. Additional guidance regarding hybridization conditions may be found in: Current Protocols in Molecular Biology, John Wiley & Sons, N.Y., 2002, and in: Sambrook et al., Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory Press, 2001.

[00100] The term "treating" or "treatment" as used herein and as is well understood in the art, means an approach for obtaining beneficial or desired results, including clinical results. Beneficial or desired clinical results can include, but are not limited to, alleviation or amelioration of one or more symptoms or conditions, diminishment of extent of disease, stabilized (i.e. not worsening) state of disease, preventing spread of disease, delay or slowing of disease progression, amelioration or palliation of the disease state, diminishment of the reoccurrence of disease, and remission (whether partial or total), whether detectable or undetectable. "Treating" and "Treatment" can also mean prolonging survival as compared to expected survival if not receiving treatment. "Treating" and "treatment" as used herein also include prophylactic treatment. For example, a subject with cancer can be treated to prevent progression can be treated with an antibody, immunoconjugate, nucleic acid or composition described herein to prevent progression.

[00101] As used herein, the term "administration" means to provide or give a subject an agent, such as a composition comprising an effective amount of an antibody by an effective route such as an intratumor or an intravenous administration route.

[00102] As used herein, the term "diluent" refers to a pharmaceutically acceptable carrier which does not inhibit a physiological activity or property of an active compound, such as an antibody, or immunoconjugate, to be administered and does not irritate the subject and does not abrogate the biological activity and properties of the administered compound. Diluents include any and all solvents, dispersion media, coatings, surfactants, antioxidants, preservative salts, preservatives, binders, excipients, disintegration agents, lubricants, such

like materials and combinations thereof, as would be known to one of ordinary skill in the art (see, for example, Remington's Pharmaceutical Sciences, 18th Ed. Mack Printing Company, 1990, pp. 1289-1329, incorporated herein by reference). Except insofar as any conventional carrier is incompatible with the active ingredient, its use in the pharmaceutical compositions is contemplated.

[00103] Compositions or methods "comprising" or "including" one or more recited elements may include other elements not specifically recited. For example, a composition that "comprises" or "includes" an antibody may contain the antibody alone or in combination with other ingredients.

[00104] In understanding the scope of the present disclosure, the term “consisting” and its derivatives, as used herein, are intended to be close ended terms that specify the presence of stated features, elements, components, groups, integers, and/or steps, and also exclude the presence of other unstated features, elements, components, groups, integers and/or steps.

[00105] The recitation of numerical ranges by endpoints herein includes all numbers and fractions subsumed within that range (e.g., 1 to 5 includes 1, 1.5, 2, 2.75, 3, 3.90, 4, and 5). It is also to be understood that all numbers and fractions thereof are presumed to be modified by the term "about." Further, it is to be understood that the singular forms of the articles "a," "an," and "the" include plural references unless the context clearly dictates otherwise. For example, the term "an antibody" or "at least one antibody" can include a plurality of antibodies, including mixtures thereof.

[00106] The terms “Frizzled” and “FZD” refer, depending on context, to any gene or protein member of the Frizzled family. Frizzled proteins are involved in the activation of Disheveled protein in the cytosol. Frizzled refers to any of Frizzled-1, Frizzled-2, Frizzled-3, Frizzled-4, Frizzled-5, Frizzled-6, Frizzled-7, Frizzled-8, Frizzled-9 and Frizzled-10.

[00107] “Lipoprotein receptor-related proteins”, “low density lipoprotein receptor-related proteins” (HGNC) or “prolow-density lipoprotein receptor-related protein” (UniProt), abbreviated “LRP”, are a group of genes and proteins. They include: LRP1, LRP1B, LRP2 (megalin), LRP3, LRP4, LRP5, LRP6, LRP8 (apolipoprotein e receptor), LRP10, LRP11, and LRP12. LRP5 and LRP6 are part of the LRP5/LRP6/Frizzled co-receptor group that is involved in canonical Wnt pathway. LRP5 is also known as LRP5, BMND1, EVR1, EVR4, HBM, LR3, LRP-5, LRP7, OPPG, OPS, OPTA1, VBCH2, and LDL receptor related protein 5. The LRP5 gene has ENTREZ Gene ID: 4041 and the protein has NCBI Reference Sequence: NP_002326. The LRP6 gene has ENTREZ Gene ID: 4040 and the protein has NCBI Reference Sequence: NP_002327. LRP6 is also known as ADCAD2, STHAG7.

II. Disorders Associated with LRP Signaling Dysregulation

[00108] Wnt binding to LRP5 or LRP6 destabilizes a β-catenin binding complex causing β-catenin degradation. The result is increased levels of intracellular β-catenin. Accordingly, provided herein are methods of blocking Wnt binding to LRP family proteins such as LRP5 or LRP6.

[00109] An "LRP-associated disorder" (e.g., an "LRP5-associated disorder" or an "LRP6-associated disorder") refers to a condition or disease correlated with dysregulation of the particular LRP receptor referred to. Dysregulation refers to abnormal decreases or increases in signaling that affect normal β-catenin mediated transcriptional changes or any other intracellular signaling pathways governed by these receptors. So, for example, abnormal LRP-associated increases in signaling, e.g., through the canonical Wnt signaling pathway (Wnt3/Wnt3A), are associated with certain cancers and with increases in bone density, while abnormal LRP-associated decreases in signaling are associated with decreases in bone density.

[00110] Antibodies that block the binding of Wnt to LRP5 or LRP6 are useful in the treatment of cancer. In particular, method of blocking Wnt binding to LRP5 is useful in the treatment of brain cancer, breast cancer, colon cancer, endometrial cancer, esophageal cancer, kidney cancer, liver cancer, lung cancer, ovarian cancer, skin cancer, stomach cancer and testicular cancer.

III. Anti-LRP5 antibodies

A. Antibodies

[00111] LRP5 and LRP6 each have two Wnt binding sites. The majority of Wnt activators of β-catenin signaling, including Wnt 1, 2, 2b, 6, 8a, 9a, 9b,10b etc., associate with β propeller 1 and 2 regions of LRP5 and LRP6. This site is referred to herein as the "non-Wnt3a binding site". Wnt proteins Wnt3 and Wnt3a are known to associate with a distinct site on β propeller regions 3 and 4 of LRP5 and LRP6. This site is referred to herein as the "Wnt3a binding site". Antibody binding to LRP5 or LRP6 that blocks binding of Wnt ligands to a particular binding site decreases activity of the signaling pathway associated with Wnt ligand binding to that site. Such blocking activity also may, in certain circumstances, potentiate activity of the signaling pathway associated with Wnt ligand binding to the other binding site. So, for example, an antibody that blocks binding of a Wnt ligand to the Wnt3a binding site inhibits activity of the Wnt3a signaling pathway and may potentiate activity of the non-Wnt3a signaling pathway. Similarly, an antibody that blocks binding of a Wnt ligand to the non-Wnt3a binding site inhibits activity of the non-Wnt3a signaling pathway and may potentiate activity of the Wnt3a signaling pathway.

[00112] Antibodies against LRP5 receptors are described herein. Certain of these antibodies block binding of Wnt ligands to the Wnt3a binding site of LRP5. Others of these antibodies block binding of Wnt ligands to the non-Wnt3a binding site. These antibodies block ligand Wnt binding and modulate activity of the Wnt signaling pathway. These antibodies also have anti-proliferative effects have therapeutic potential for treating cancer and other diseases where the LRP receptors are dysregulated.

[00113] Accordingly, an aspect of the disclosure includes an isolated antibody that specifically binds LRP5 receptor. The antibodies comprise a light chain variable region and a heavy chain variable region, the heavy chain variable region comprising complementarity determining regions CDR-H1, CDR-H2 and CDR-H3, the light chain variable region comprising complementarity determining region CDR-L1, CDR-L2 and CDR-L3, and with the amino acid sequences of said CDRs comprising, consisting essentially of, or consisting of sequences selected from sequences in Table 1 or 2.

[00114] In an embodiment, the antibody comprises a CDR sequence set selected from the CDR sequence sets in Table 1, that is, for clones LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[00115] Also described herein are heavy chain and light chain variable regions. Table 2 provides exemplary variable domain sequences for the Fab heavy and light chains, from clones LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6. Antibodies comprising the sequences in Table 2 or sequences substantially identical thereto, wherein the CDRs are a CDR sequence set identified in Tables 1 are also contemplated. In another embodiment, the antibody comprises a heavy chain variable region comprising: i) a heavy chain amino acid sequence as set forth in Table 2; ii) an amino acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the heavy chain amino acid sequence as set forth in Table 2, wherein the CDR sequences are a CDR sequence set as set forth in Table 1, or iii) a conservatively substituted amino acid sequence of i) wherein the CDR sequences are a CDR sequence set as set forth in Table 1.

[00116] In another embodiment, the antibody comprises a light chain variable region comprising i) a light chain amino acid sequence as set forth in Table 2, ii) an amino acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the light chain amino acid sequence as set forth in Table 2, wherein the CDR sequences are a CDR sequence set as set forth in

Table 1, or iii) a conservatively substituted amino acid sequence of i) wherein the CDR sequences are a CDR sequence set as set forth in Table 1.

[00117] In Table 1, below antibodies are assigned to epitope groups based on their binding characteristics. Antibodies in Epitope Group 1 potentiate the activity of Wnt ligands binding to a non-Wnt3A binding site. However, they do not inhibit Wnt3A ligand activity. Antibodies in Epitope Group 2 inhibit activity of Wnt ligand binding to non-Wnt3a binding sites. Furthermore, antibody LRP5-G 10 potentiates activity of went ligands binding to Wnt3a binding sites. Antibodies belonging to Epitope Groups 3 and 4bind LRP5 but do not inhibit or potentiate activity of Wnt ligands. Epitope groups of other antibodies were not determined.

[00118] Amino acid sequences for CDRs of these antibodies are provided in Table 1.

[00119] Table 1: CDR sequences of anti-LRP5 antibodies

Clone ID	**CDR-L1**	**SEQ. ID NO:**	**CDR-L2**	**SEQ. ID NO:**	**CDR-L3**	**SEQ. ID NO:**	**Epitope Group**
LRP5 – A7	SVSSA	34	SASSLYS	35	AWGWGLF	36	1
LRP5 – A9	SVSSA	34	SASSLYS	35	VHYSPYSLI	37	ND
LRP5 – C5	SVSSA	34	SASSLYS	35	YQYSGLI	38	1
LRP5 – C12	SVSSA	34	SASSLYS	35	FSHVSLI	39	2
LRP5 – D9	SVSSA	34	SASSLYS	35	ASYSPI	40	1
LRP5 – E5	SVSSA	34	SASSLYS	35	YHYYYLF	41	4
LRP5 – G2	SVSSA	34	SASSLYS	35	ASYAPI	42	1
LRP5 – G9	SVSSA	34	SASSLYS	35	SSSSPI	43	1
LRP5 – G10	SVSSA	34	SASSLYS	35	SSYSLI	44	2
LRP5 – G11	SVSSA	34	SASSLYS	35	GVSLI	45	1
LRP5 – H3	SVSSA	34	SASSLYS	35	YWFLI	46	3
LRP5 – H5	SVSSA	34	SASSLYS	35	PVGHYGYPI	47	2
LRP5 – H9	SVSSA	34	SASSLYS	35	SSYSPI	48	1
LRP5 – R3O_D3	SVSSA	34	SASSLYS	35	YWAYYSPI	49	ND
LRP5 – R3_E8	SVSSA	34	SASSLYS	35	VSYYPLI	51	ND
LRP5 – R3O_G6	SVSSA	34	SASSLYS	35	SSYSLI	44	ND

Clone ID	CDR-H1	SEQ ID NO:	CDR-H2	SEQ ID NO:	CDR-H3	SEQ ID NO:	Epitope Group
LRP5 – A7	LSYSSM	2	SIYPYYGYTY	11	HGAM	21	1

LRP5 – A9	ISSYSI	3	SSSYYGYTY	12	TVRGSKKPYFSGWAM	22	ND
LRP5 – C5	FSSSSI	4	SISSSYGYTY	13	SSYYSSVSSSVYAL	23	1
LRP5 – C12	FSSSSI	4	SIYSSYGSTS	14	TVRGSKKPYFSGWAM	22	2
LRP5 – D9	ISYSYI	5	SIYSSYGYTY	15	HYSYFFYAM	24	1
LRP5 – E5	IYSYSI	6	SIYPYSSYTS	16	YAVYFPGYYWGM	25	4
LRP5 – G2	LSYYYM	7	SIYSSYGYTY	15	WSHVSGHYSGM	26	1
LRP5 – G9	ISYSYI	5	SIYSSYGYTY	15	WGAYHSSGYGM	27	1
LRP5 – G10	FSSSSI	4	SISSSYGYTY	13	GGSGVSHYGSVYYSWWAL	28	2
LRP5 – G11	ISYSYI	5	SIYPSYGYTY	17	AAPYYGYYYSYAM	29	1
LRP5 – H3	LYYYYI	8	SISPYYGYTS	18	SGYGWYAM	30	3
LRP5 – H5	LSYSSI	9	SISSSYGSTS	19	GYWAI	31	2
LRP5 – H9	IYSYYI	10	SIYSYYGYTY	20	SYPAM	32	1
LRP5 – R3O_D3	FSSSSI	4	SISSSYGYTY	13	SWAM	33	ND
LRP5 – R3_E8	LYYSSM	52	SISSYYGYTS	53	YWAL	54	ND
LRP5 – R3O_G6	ISSYSM	55	YISPYYGYTS	56	GWGSPASAGYYGL	57	ND

[00120] Amino acid and nucleotide sequences for the variable heavy and variable light domains are provided in Table 2:

[00121] **Table 2** – heavy chain and light chain DNA and amino acid sequences of LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[00122]

LRP5-A7

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACCTCTCTTATTCTTCTATGCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTATCC
TTATTATGGCTATACTTATTATGCCGATAGCGTCAAGGGCCGTTTCACTATAA
GCGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAG
CTGAGGACACTGCCGTCTATTATTGTGCTCGCCATGGTGCTATGGACTACTGG
GGTCAAGGAACCCTGGTCACCGTCTCCTCG (SEQ ID NO.: 58)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNLSYSSMHWVRQAPGKGLEWVASIYP
YYGYTYYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARHGAMDYW
GQGTLVTVSS (SEQ ID NO.: 59)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAAGCTTGGGGTTGGGGTCTGTTCACGTTCGGACAGGGTACCAAGGT
GGAGATCAAACGT (SEQ ID NO.: 60)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQAWGWGLFTFGQGTKVEIKR

[00123] (SEQ ID NO.: 61)

LRP5-D9

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACATCTCTTATTCTTATATCCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTATTCT
TCTTATGGCTATACTTATTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAG
CGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAGCT
GAGGACACTGCCGTCTATTATTGTGCTCGCCATTACTCTTACTTCTTCTACGCT
ATGGACTACTGGGGTCAAGGAACCCTGGTCACCGTCTCCTCG (SEQ ID NO.: 62)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNISYSYIHWVRQAPGKGLEWVASIYSSY
GYTYYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARHYSYFFYAMD
YWGQGTLVTVSS (SEQ ID NO.: 63)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAAGCTTCTTACTCTCCGATCACGTTCGGACAGGGTACCAAGGTGGA
GATCAAACGT (SEQ ID NO.: 64)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQASYSPITFGQGTKVEIKR
[00124] (SEQ ID NO.: 65)

LRP5-E5

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACATCTATTCTTATTCTATCCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTATCC
TTATTCTAGCTATACTTCTTATGCCGATAGCGTCAAGGGCCGTTTCACTATAA
GCGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAG
CTGAGGACACTGCCGTCTATTATTGTGCTCGCTACGCTGTTTACTTCCCGGGT
TACTACTGGGGTATGGACTACTGGGGTCAAGGAACCCTGGTCACCGTCTCCTC
G (SEQ ID NO.: 66)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNIYSYSIHWVRQAPGKGLEWVASIYPYS
SYTSYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARYAVYFPGYYW
GMDYWGQGTLVTVSS (SEQ ID NO.: 67)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAATACCATTACTACTACCTGTTCACGTTCGGACAGGGTACCAAGGT
GGAGATCAAACGT (SEQ ID NO.: 68)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQYHYYYLFTFGQGTKVEIKR
(SEQ ID NO.: 69)

[00125]

LRP5-G2

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACCTCTCTTATTATTATATGCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTATTCT
TCTTATGGCTATACTTATTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAG
CGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAGCT
GAGGACACTGCCGTCTATTATTGTGCTCGCTGGTCTCATGTTTCTGGTCATTA
CTCTGGTATGGACTACTGGGGTCAAGGAACCCTGGTCACCGTCTCCTCG
(SEQ ID NO.: 70)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNLSYYYMHWVRQAPGKGLEWVASIYS
SYGYTYYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARWSHVSGHY
SGMDYWGQGTLVTVSS (SEQ ID NO.: 71)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAAGCTTCTTACGCTCCGATCACGTTCGGACAGGGTACCAAGGTGGA
GATCAAACGT (SEQ ID NO.: 72)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQASYAPITFGQGTKVEIKR

[00126] (SEQ ID NO.: 73)

LRP5-G9

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACATCTCTTATTCTTATATCCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTATTCT
TCTTATGGCTATACTTATTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAG
CGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAGCT
GAGGACACTGCCGTCTATTATTGTGCTCGCTGGGGTGCTTACCATTCTTCTGG
TTACGGTATGGACTACTGGGGTCAAGGAACCCTGGTCACCGTCTCCTCG
VH-AA (SEQ ID NO.: 74)

EVQLVESGGGLVQPGGSLRLSCAASGFNISYSYIHWVRQAPGKGLEWVASIYSSY
GYTYYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARWGAYHSSGY
GMDYWGQGTLVTVSS (SEQ ID NO.: 75)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAATCTTCTTCTTCTCCGATCACGTTCGGACAGGGTACCAAGGTGGAG
ATCAAACGT (SEQ ID NO.: 76)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQSSSSPITFGQGTKVEIKR
(SEQ ID NO.: 77)

[00127]

LRP5-G10

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACTTTTCTTCTTCTTCTATACACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTCTTCT
TCTTATGGCTATACTTATTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAG
CGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAGCT
GAGGACACTGCCGTCTATTATTGTGCTCGCGGTGGTTCTGGTGTTTCTCATTA
CGGTTCTGTTTACTACTCTTGGTGGGCTTTGGACTACTGGGGTCAAGGAACCC
TGGTCACCGTCTCCTCG (SEQ ID NO.: 78)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNFSSSSIHWVRQAPGKGLEWVASISSSY
GYTYYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARGGSGVSHYGS
VYYSWWALDYWGQGTLVTVSS (SEQ ID NO.: 79)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAATCTTCTTATTCTCTGATCACGTTCGGACAGGGTACCAAGGTGGAG
ATCAAACGT (SEQ ID NO.: 80)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQSSYSLITFGQGTKVEIKR
(SEQ ID NO.: 81)

[00128]

LRP5-G11

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACATCTCTTATTCTTATATCCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTATCC
TTCTTATGGCTATACTTATTATGCCGATAGCGTCAAGGGCCGTTTCACTATAA
GCGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAG
CTGAGGACACTGCCGTCTATTATTGTGCTCGCGCTGCTCCGTACTACGGTTAC
TACTACTCTTACGCTATGGACTACTGGGGTCAAGGAACCCTGGTCACCGTCTC
CTCG (SEQ ID NO.: 82)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNISYSYIHWVRQAPGKGLEWVASIYPSY
GYTYYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARAAPYYGYYYS
YAMDYWGQGTLVTVSS (SEQ ID NO.: 83)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAAGGTGTTTCTCTGATCACGTTCGGACAGGGTACCAAGGTGGAGAT
CAAACGT (SEQ ID NO.: 84)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQGVSLITFGQGTKVEIKR
(SEQ ID NO.: 85)

[00129]

LRP5-H3

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACCTCTATTATTATTATATCCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTCTCCT
TATTATGGCTATACTTCTTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAG
CGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAGCT
GAGGACACTGCCGTCTATTATTGTGCTCGCTCTGGTTACGGTTGGTACGCTAT
GGACTACTGGGGTCAAGGAACCCTGGTCACCGTCTCCTCG (SEQ ID NO.: 86)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNLYYYYIHWVRQAPGKGLEWVASISPY
YGYTSYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARSGYGWYAM
DYWGQGTLVTVSS (SEQ ID NO.: 87)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAATACTGGTTCCTGATCACGTTCGGACAGGGTACCAAGGTGGAGAT
CAAACGT(SEQ ID NO.: 88)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQYWFLITFGQGTKVEIKR
(SEQ ID NO.: 89)

[00130]

LRP5-H5

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACCTCTCTTATTCTTCTATCCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTCTTCT
TCTTATGGCTCTACTTCTTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAG
CGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAGCT
GAGGACACTGCCGTCTATTATTGTGCTCGCGGTTACTGGGCTATTGACTACTG
GGGTCAAGGAACCCTGGTCACCGTCTCCTCG (SEQ ID NO.: 90)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNLSYSSIHWVRQAPGKGLEWVASISSSY
GSTSYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARGYWAIDYWGQ
GTLVTVSS (SEQ ID NO.: 91)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAACCGGTTGGTCATTACGGTTACCCGATCACGTTCGGACAGGGTAC
CAAGGTGGAGATCAAACGT (SEQ ID NO.: 92)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQPVGHYGYPITFGQGTKVEIKR
(SEQ ID NO.: 93)

[00131]

LRP5-H9

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACATCTATTCTTATTATATCCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTATTCT
TATTATGGCTATACTTATTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAG
CGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAGCT
GAGGACACTGCCGTCTATTATTGTGCTCGCTCTTACCCGGCTATGGACTACTG
GGGTCAAGGAACCCTGGTCACCGTCTCCTCG (SEQ ID NO.: 94)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNIYSYYIHWVRQAPGKGLEWVASIYSY
YGYTYYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARSYPAMDYW
GQGTLVTVSS (SEQ ID NO.: 95)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAATCTTCTTACTCTCCGATCACGTTCGGACAGGGTACCAAGGTGGA
GATCAAACGT (SEQ ID NO.: 96)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQSSYSPITFGQGTKVEIKR
(SEQ ID NO.: 97)

[00132]

LRP5-R3O_D3

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACTTTTCTTCTTCTTCTATACACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTCTTCT
TCTTATGGCTATACTTATTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAG
CGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAGCT
GAGGACACTGCCGTCTATTATTGTGCTCGCTCTTGGGCTATGGACTACTGGGG
TCAAGGAACCCTGGTCACCGTCTCCTCG (SEQ ID NO.: 98)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNFSSSSIHWVRQAPGKGLEWVASISSSY
GYTYYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARSWAMDYWGQ
GTLVTVSS (SEQ ID NO.: 99)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAATACTGGGCTTACTACTCTCCGATCACGTTCGGACAGGGTACCAA
GGTGGAGATCAAACGT (SEQ ID NO.: 100)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQYWAYYSPITFGQGTKVEIKR
(SEQ ID NO.: 101)

[00133]

LRP5-R3_E8

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACCTCTATTATTCTTCTATGCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTCTTCT
TATTATGGCTATACTTCTTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAG
CGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAGCT
GAGGACACTGCCGTCTATTATTGTGCTCGCTACTGGGCTTTGGACTACTGGGG
TCAAGGAACCCTGGTCACCGTCTCCTCG (SEQ ID NO.: 102)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNLYYSSMHWVRQAPGKGLEWVASISS
YYGYTSYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARYWALDYW
GQGTLVTVSS (SEQ ID NO.: 103)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAAGTTTCTTACTACCCGCTGATCACGTTCGGACAGGGTACCAAGGT
GGAGATCAAACGT (SEQ ID NO.: 104)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQVSYYPLITFGQGTKVEIKR
(SEQ ID NO.: 105)

[00134]

LRP5-R3O_G6

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCAC
TCCGTTTGTCCTGTGCAGCTTCTGGCTTCAACATCTCTTCTTATTCTATGCACT
GGGTGCGTCAGGCCCCGGGTAAGGGCCTGGAATGGGTTGCATATATTTCTCC
TTATTATGGCTATACTTCTTATGCCGATAGCGTCAAGGGCCGTTTCACTATAA
GCGCAGACACATCCAAAAACACAGCCTACCTACAAATGAACAGCTTAAGAG
CTGAGGACACTGCCGTCTATTATTGTGCTCGCGGTTGGGGTTCTCCGGCTTCT
GCTGGTTACTACGGTTTGGACTACTGGGGTCAAGGAACCCTGGTCACCGTCTC
CTCG (SEQ ID NO.: 106)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNISSYSMHWVRQAPGKGLEWVAYISPY
YGYTSYADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARGWGSPASAG
YYGLDYWGQGTLVTVSS (SEQ ID NO.: 107)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATA
GGGTCACCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTG
GTATCAACAGAAACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCC
AGCCTCTACTCTGGAGTCCCTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGA
TTTCACTCTGACCATCAGCAGTCTGCAGCCGGAAGACTTCGCAACTTATTACT
GTCAGCAATCTTCTTATTCTCTGATCACGTTCGGACAGGGTACCAAGGTGGAG
ATCAAACGT (SEQ ID NO.: 108)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLY
SGVPSRFSGSRSGTDFTLTISSLQPEDFATYYCQQSSYSLITFGQGTKVEIKR
(SEQ ID NO.: 109)

[00135] LRP5-A9

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCACTCCGT
TTGTCCTGTGCAGCTTCTGGCTTCAACATCTCTTCTTATTCTATCCACTGGGTGCGTCAG
GCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTTCTTCTTATTATGGCTATACTTATTA
TGCCGATAGCGTCAAGGGCCGTTTCACTATAAGCGCAGACACATCCAAAAACACAGCC
TACCTACAAATGAACAGCTTAAGAGCTGAGGACACTGCCGTCTATTATTGTGCTCGCAC
TGTTCGTGGATCCAAAAAACCGTACTTCTCTGGTTGGGCTATGGACTACTGGGGTCAAG
GAACCCTGGTCACCGTCTCCTCG (SEQ ID NO.: 114)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNISSYSIHWVRQAPGKGLEWVASSSYYGYTYYA

DSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARTVRGSKKPYFSGWAMDYWGQGT LVTVSS (SEQ ID NO.: 115)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATAGGGTCA CCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTGGTATCAACAGAA ACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCCAGCCTCTACTCTGGAGTCC CTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGATTTCACTCTGACCATCAGCAGTCTG CAGCCGGAAGACTTCGCAACTTATTACTGTCAGCAAGTGCACTACAGCCCCTACAGCCT GATC ACGTTCGGACAGGGTACCAAGGTGGAGATCAAACGA (SEQ ID NO.: 116)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLYSGVPSR FSGSRSGTDFTLTISSLQPEDFATYYCQQ VHYSPYSLITFGQGTKVEIKR (SEQ ID NO.: 117)

[00136] LRP5-C5

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCACTCCGT TTGTCCTGTGCAGCTTCTGGCTTCAACTTTTCTTCTTCTTCTATACACTGGGTGCGTCAG GCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTCTTCTTCTTATGGCTATACTTA TTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAGCGCAGACACATCCAAAAACACAG CCTACCTACAAATGAACAGCTTAAGAGCTGAGGACACTGCCGTCTATTATTGTGCTCGC TCTTCTTACTACTCTTCTGTTTCTTCTTCTGTTTACGCTTTGGACTACTGGGGTCAAGGA ACCCTGGTCACCGTCTCCTCG (SEQ ID NO.: 110)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNFSSSSIHWVRQAPGKGLEWVASISSSYGYTYY ADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARSSYYSSVSSSVYALDYWGQGTL VTVSS (SEQ ID NO.: 111)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATAGGGTCA CCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTGGTATCAACAGAA ACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCCAGCCTCTACTCTGGAGTCC CTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGATTTCACTCTGACCATCAGCAGTCTG CAGCCGGAAGACTTCGCAACTTATTACTGTCAGCAATACCAGTACTCTGGTCTGATCAC GTTCGGACAGGGTACCAAGGTGGAGATCAAACGA (SEQ ID NO.: 112)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLYSGVPSR FSGSRSGTDFTLTISSLQPEDFATYYCQQ YQYSGLITFGQGTKVEIKR (SEQ ID NO.: 113)

[00137] LRP5-C12

VH-DNA

GAGGTTCAGCTGGTGGAGTCTGGCGGTGGCCTGGTGCAGCCAGGGGGCTCACTCCGT
TTGTCCTGTGCAGCTTCTGGCTTCAACTTTTCTTCTTCTTCTATACACTGGGTGCGTCAG
GCCCCGGGTAAGGGCCTGGAATGGGTTGCATCTATTTATTCTTCTTATGGCTCTACTTC
TTATGCCGATAGCGTCAAGGGCCGTTTCACTATAAGCGCAGACACATCCAAAAACACAG
CCTACCTACAAATGAACAGCTTAAGAGCTGAGGACACTGCCGTCTATTATTGTGCTCGC
ACTGTTCGTGGATCCAAAAAACCGTACTTCTCTGGTTGGGCTATGGACTACTGGGGTCA
AGGAACCCTGGTCACCGTCTCCTCG (SEQ ID NO.: 118)

VH-AA

EVQLVESGGGLVQPGGSLRLSCAASGFNFSSSSIHWVRQAPGKGLEWVASIYSSYGSTSY
ADSVKGRFTISADTSKNTAYLQMNSLRAEDTAVYYCARTVRGSKKPYFSGWAMDYWGQG
TLVTVSS (SEQ ID NO.: 119)

VL-DNA

GATATCCAGATGACCCAGTCCCCGAGCTCCCTGTCCGCCTCTGTGGGCGATAGGGTCA
CCATCACCTGCCGTGCCAGTCAGTCCGTGTCCAGCGCTGTAGCCTGGTATCAACAGAA
ACCAGGAAAAGCTCCGAAGCTTCTGATTTACTCGGCATCCAGCCTCTACTCTGGAGTCC
CTTCTCGCTTCTCTGGTAGCCGTTCCGGGACGGATTTCACTCTGACCATCAGCAGTCTG
CAGCCGGAAGACTTCGCAACTTATTACTGTCAGCAATTCAGCCACGTGAGCCTGATCAC
GTTCGGACAGGGTACCAAGGTGGAGATCAAACGA (SEQ ID NO.: 120)

VL-AA

DIQMTQSPSSLSASVGDRVTITCRASQSVSSAVAWYQQKPGKAPKLLIYSASSLYSGVPSR
FSGSRSGTDFTLTISSLQPEDFATYYCQQ FSHVSLITFGQGTKVEIKR (SEQ ID NO.: 121)

[00138] In some embodiments, the variable domain sequences are at least 95%, 96%, 97%, 98%, or 99% similar outside of the CDR regions and the CDR sequence set is 100% identical to the amino acid sequences provided in Table 1.

[00139] Also provided in another embodiment, is a competing antibody that competes for binding with an antibody comprising a CDR sequence set described herein. For example, the competing antibody in one embodiment reduces binding of the antibody comprising the CDR sequence set to LRP5 CDR by at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99%.

[00140] As demonstrated herein, the antibodies described herein have high affinity for LRP5. For example, the antibodies in one embodiment, have a binding affinity measured by surface plasmon resonance of between about 1 nM and about 50 nM.

[00141] The antibody can be a humanized antibody as described herein or a chimeric antibody.

[00142] In some embodiments, the antibody is a single chain antibody which can be obtained for example, by fusing the heavy chain and light chain or parts thereof together.

[00143] In some embodiments, the antibody is an antibody binding fragment selected from Fab, Fab', F(ab')2, scFv, dsFv, ds-scFv, dimers, nanobodies, minibodies, diabodies, and multimers thereof.

[00144] In some other embodiments the antibody is the binding fragment Fab. For some embodiments, the binding fragment is preferable, e.g., *in vitro* uses.

[00145] It may be preferable in other embodiments to have a multivalent antibody or an antibody comprising an Ig portion.

[00146] As demonstrated in the Examples, a Fab fragment can be combined with an Ig such as an IgG. In an embodiment, the IgG is IgG1, IgG2, IgG3 or IgG4.

B. Detectably Labeled Antibodies

[00147] Detectable labels can include peptide sequences (such a myc tag, HA-tag, V5-tag or NE-tag), fluorescent or luminescent proteins (e.g., green fluorescent protein or luciferase) that can be appended to or introduced into an antibody described herein and which is capable of producing, either directly or indirectly, a detectable signal. For example, the label may be radio-opaque, positron-emitting radionuclide (for example, for use in PET imaging), or a radioisotope, such as ^{3}H, ^{13}N, ^{14}C, ^{18}F, ^{32}P, ^{35}S, ^{123}I, ^{125}I, ^{131}I; a fluorescent (fluorophore) or chemiluminescent (chromophore) compound, such as fluorescein isothiocyanate, rhodamine or luciferin; an enzyme, such as alkaline phosphatase, beta-galactosidase or horseradish peroxidase; an imaging agent; or a metal ion.

C. Antibody-Drug Conjugates

[00148] A further aspect includes an immunoconjugate comprising an antibody described herein and a detectable label or cytotoxic agent.

[00149] A chemotherapeutic (anti-cancer) agent can be any agent capable of reducing cancer growth, interfering with cancer cell replication, directly or indirectly killing cancer cells, reducing metastasis, reducing tumor blood supply, etc. Chemotherapeutic agents thus include cytotoxic agents. Cytotoxic agents include but are not limited to saporin, taxanes, vinca alkaloids, anthracycline, and platinum-based agents. Classes of chemotherapeutic agents include but are not limited to alkylating agents, antimetabolites, *e.g.*, methotrexate, plant alkaloids, *e.g.*, vincristine, and antitumor antibiotics such as anthracyclines, *e.g.*, doxorubicin as well as miscellaneous drugs that do not fall in to a particular class such as hydroxyurea. Platinum-based drugs, exemplified by cisplatin and oxaliplatin, represent a major class of chemotherapeutics. These drugs bind to DNA and interfere with replication. Taxanes, exemplified by taxol, represent another major class of chemotherapeutics. These compounds act by interfering with cytoskeletal and spindle formation to inhibit cell division, and thereby

prevent growth of rapidly dividing cancer cells. Other chemotherapeutic drugs include hormonal therapy. Chemotherapeutics also include agents that inhibit tubulin assembly or polymerization such as maytansine, mertansine, and auristatin. Chemotherapeutic agents also include DNA damage agents such as calicheamicin.

[00150] Chemotherpeutic agents can include maytansinoid, auristatin, dolastatin, tubulysin, cryptophycin, pyrrolobenzodiazepine (PBD) dimer, indolinobenzodiazepine dimer, alpha-amanitin, trichothene, SN-38, duocarmycin, CC1065, calicheamincin, an enediyne antibioatic, taxane, doxorubicin derivatives, anthracycline and stereoisomers, azanofide, isosteres, analogs or derivatives thereof.

IV. Nucleic Acids

[00151] Further aspects include nucleic acid molecules or polynucleotides, recombinant nucleic acid molecules, expression constructs, and vectors as described herein.

A. Nucleic Acid Molecules

[00152] A further aspect includes a nucleic acid molecule as set forth in Table 2, as well as a polynucleotide that hybridizes to one of said sequences, for example, under stringent hybridization conditions. The CDR and variable domain nucleic sequences can be used for example, to prepare expression constructs.

B. Expression Constructs and Vectors

[00153] The nucleic acid molecules may be incorporated in a known manner into an appropriate expression construct or expression vector which ensures expression of the protein. Expression constructs can comprise an expression control sequence, e.g., a promoter, operatively linked with a polynucleotide comprising a nucleotide sequence encoding an antibody of this disclosure. Possible expression vectors include but are not limited to cosmids, plasmids, or modified viruses (e.g., replication defective retroviruses, adenoviruses and adeno-associated viruses). The vector should be compatible with the host cell used. The expression vectors are "suitable for transformation of a host cell", which means that the expression vectors contain a nucleic acid molecule encoding the peptides corresponding to epitopes or antibodies described herein.

[00154] In an embodiment, the vector is suitable for expressing for example, single chain antibodies by gene therapy. In an embodiment, the vector comprises an IRES and allows for expression of a light chain variable region and a heavy chain variable region. Such vectors can be used to deliver antibody *in vivo*.

[00155] Suitable regulatory sequences may be derived from a variety of sources, including bacterial, fungal, viral, mammalian, or insect genes.

[00156] Examples of such regulatory sequences include: a transcriptional promoter and enhancer or RNA polymerase binding sequence, a ribosomal binding sequence, including a translation initiation signal. Additionally, depending on the host cell chosen and the vector employed, other sequences, such as an origin of replication, additional DNA restriction sites, enhancers, and sequences conferring inducibility of transcription may be incorporated into the expression vector.

[00157] In an embodiment, the regulatory sequences direct or increase expression in neural tissue and/or cells.

[00158] The vector can be any vector, including vectors suitable for producing an antibody described herein.

[00159] In an embodiment, the vector is a viral vector.

[00160] The recombinant expression vectors may also contain a marker gene which facilitates the selection of host cells transformed, infected or transfected with a vector for expressing an antibody or epitope peptide described herein.

[00161] The recombinant expression vectors may also contain expression cassettes which encode a fusion moiety (i.e. a “fusion protein”) which provides increased expression or stability of the recombinant peptide; increased solubility of the recombinant peptide; and aid in the purification of the target recombinant peptide by acting as a ligand in affinity purification, including for example, tags and labels described herein. Further, a proteolytic cleavage site may be added to the target recombinant protein to allow separation of the recombinant protein from the fusion moiety subsequent to purification of the fusion protein. Typical fusion expression vectors include pGEX (Amrad Corp., Melbourne, Australia), pMAL (New England Biolabs, Beverly, MA) and pRIT5 (Pharmacia, Piscataway, NJ) which fuse glutathione S-transferase (GST), maltose E binding protein, or protein A, respectively, to the recombinant protein.

[00162] Systems for the transfer of genes both *in vitro* and *in vivo* include vectors based on viruses, most notably Herpes Simplex Virus, Adenovirus, Adeno-associated virus (AAV) and retroviruses including lentiviruses. Alternative approaches for gene delivery include the use of naked, plasmid DNA as well as liposome–DNA complexes.

[00163] In an aspect the disclosure includes a method for making an antibody described herein, the method comprising synthesizing a nucleic acid molecule that comprises an antibody framework and a CDR sequence set described herein.

V. Recombinant Cells

[00164] A further aspect is a recombinant host cell expressing an antibody described herein.

[00165] Antibodies as described herein can be made by recombinant expression of nucleic acids encoding the antibody sequences.

[00166] Antibodies as disclosed herein can be made by culturing cells engineered to express nucleic acid constructs encoding immunoglobulin polypeptides.

[00167] The recombinant host cell can be generated using any cell suitable for producing a polypeptide, for example, suitable for producing an antibody. For example, to introduce a nucleic acid (e.g., a vector) into a cell, the cell may be transfected, transformed or infected, depending upon the vector employed.

[00168] Suitable host cells include a wide variety of prokaryotic and eukaryotic host cells. For example, the proteins described herein may be expressed in bacterial cells such as E. coli, insect cells (using baculovirus), yeast cells or mammalian cells.

[00169] In an embodiment, the cell is a eukaryotic cell selected from a yeast, plant, worm, insect, avian, fish, reptile and mammalian cell.

[00170] In another embodiment, the mammalian cell is a CHO cell, a myeloma cell, a spleen cell, or a hybridoma cell.

[00171] Yeast and fungi host cells suitable for expressing an antibody include, but are not limited to Saccharomyces cerevisiae, Schizosaccharomyces pombe, the genera Pichia or Kluyveromyces and various species of the genus Aspergillus. Examples of vectors for expression in yeast S. cerivisiae include pYepSec1, pMFa, pJRY88, and pYES2 (Invitrogen Corporation, San Diego, CA). Protocols for the transformation of yeast and fungi are well known to those of ordinary skill in the art.

[00172] Mammalian cells that may be suitable include, among others: COS (e.g., ATCC No. CRL 1650 or 1651), BHK (e.g., ATCC No. CRL 6281), CHO (ATCC No. CCL 61), HeLa (e.g., ATCC No. CCL 2), 293 (ATCC No. 1573) and NS-1 cells. Suitable expression vectors for directing expression in mammalian cells generally include a promoter (e.g., derived from viral material such as polyoma, Adenovirus 2, cytomegalovirus and Simian Virus 40), as well as other transcriptional and translational control sequences. Examples of mammalian expression vectors include pCDM8 and pMT2PC.

VI. Pharmaceutical Compositions

[00173] A further aspect is a composition comprising an antibody, immunoconjugate, nucleic acid molecule, vector or recombinant cell described herein, optionally with a suitable diluent, e.g., a pharmaceutically acceptable carrier.

[00174] The composition can for example, comprise one or more antibodies or immunoconjugates.

[00175] Suitable diluents for polypeptides, including antibodies and/or cells include but are not limited to saline solutions, pH buffered solutions and glycerol solutions or other solutions suitable for freezing polypeptides and/or cells.

[00176] Suitable diluents for nucleic acids include but are not limited to water, saline solutions and ethanol.

[00177] In an embodiment, the composition is a pharmaceutical composition comprising any of the antibodies, nucleic acids or vectors disclosed herein, and optionally comprising a pharmaceutically acceptable vehicle such as a diluent or carrier.

[00178] The compositions described herein can be prepared by per se known methods for the preparation of pharmaceutically acceptable compositions that can be administered to subjects, such that an effective quantity of the active substance is combined in a mixture with a pharmaceutically acceptable vehicle.

[00179] Pharmaceutical compositions include, without limitation, lyophilized powders or aqueous or non-aqueous sterile injectable solutions or suspensions, which may further contain antioxidants, buffers, bacteriostats and solutes that render the compositions substantially compatible with the tissues or the blood of an intended recipient. Other components that may be present in such compositions include water, surfactants (such as Tween), alcohols, polyols, glycerin and vegetable oils, for example. Extemporaneous injection solutions and suspensions may be prepared from sterile powders, granules, tablets, or concentrated solutions or suspensions. The composition may be supplied, for example, but not by way of limitation, as a lyophilized powder which is reconstituted with sterile water or saline prior to administration to the patient.

[00180] Pharmaceutical compositions may comprise a pharmaceutically acceptable carrier. Suitable pharmaceutically acceptable carriers include essentially chemically inert and nontoxic compositions that do not interfere with the effectiveness of the biological activity of the pharmaceutical composition. Examples of suitable pharmaceutical carriers include, but are not limited to, water, saline solutions, glycerol solutions, ethanol, N-(1(2,3-dioleyloxy)propyl)N,N,N-trimethylammonium chloride (DOTMA), diolesylphosphotidyl-

ethanolamine (DOPE), and liposomes. Such compositions should contain a therapeutically effective amount of the compound, together with a suitable amount of carrier so as to provide the form for direct administration to the patient.

[00181] The composition may be in the form of a pharmaceutically acceptable salt which includes, without limitation, those formed with free amino groups such as those derived from hydrochloric, phosphoric, acetic, oxalic, tartaric acids, etc., and those formed with free carboxyl groups such as those derived from sodium, potassium, ammonium, calcium, ferric hydroxides, isopropylamine, triethylamine, 2-ethylarnino ethanol,

[00182] In an embodiment, the composition comprises an antibody described herein. In another embodiment, the composition comprises an antibody described herein and a diluent. In an embodiment, the composition is a sterile composition.

[00183] A further aspect includes an antibody complex comprising an antibody described herein bound to an LRP protein, e.g., LRP5 or LRP6. The complex may be in solution or comprised in a tissue, optionally *in vitro*.

[00184] Also provided are methods for making and using the regents described herein.

VII. Kits

[00185] Another aspect is a kit or package comprising any of the antibodies, immunoconjugates, nucleic acid molecules, vectors, recombinant cells and/or compositions comprised herein. The antibodies, immunoconjugates, nucleic acid molecules, vectors, recombinant cells and/or compositions can be comprised in a vial such as a sterile vial or other housing. As used herein, the term “kit” refers to a collection of items intended for use together. The kit can optionally include a reference agent and/or instructions for use thereof. A kit can further include a shipping container adapted to hold a container, such as a vial, that contains a composition as disclosed herein.

VIII. Methods of Using Antibodies

[00186] Antibodies described herein can be used in a number of *in vitro* and *in vivo* methods.

A. Methods of Detecting Expression of LRP5

[00187] As demonstrated herein, the antibodies can be used to detect LRP5 expression.

[00188] Accordingly, the disclosure provides in one aspect, a method of detecting LRP5 expression, the method comprising contacting a sample comprising one or more cells with one or more antibody or immunoconjugates described herein under conditions permissive for forming an antibody: LRP5 complex and detecting the presence of any antibody complex.

Typically, the antibody is part of an immunoconjugate comprising an antibody coupled to a detectable label.

[00189] The sample can comprise viable cells or a cell extract. The antibody: LRP5 complex can be detected immunoassays such as immunofluorescence, flow cytometry, Western blots, ELISA, SPR and immunoprecipitation followed by SDS-PAGE immunocytochemistry. In some embodiments, the detection is by immunofluorescence. In some embodiments, the detection is by flow cytometry.

[00190] As demonstrated herein, a number of the antibodies identified preferentially recognize LRP5. Accordingly, in embodiments wherein the method is for detecting LRP5 expression, the antibody or immunoconjugate comprises a CDR sequence set corresponding to an antibody selected from LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

B. Methods of inhibiting WNT binding to LRP5

[00191] Antibodies disclosed herein inhibit binding of Wnt to LRP proteins, in particular, to LRP5. Without wishing to be limited by theory, inhibition of Wnt binding to LRP5 impacts signal transduction induced by the binding of the particular Wnt ligand. For example, antibody binding to LRP5 receptors inhibits LRP5 promotion of beta-catenin phosphorylation. Because phosphorylated beta-catenin is marked for destruction in a cell, the non-phosphorylated form builds up. Accumulation of beta-catenin is associated with malignancy.

[00192] It can be desirable to reduce or inhibit Wnt ligand signaling through LRP5. Accordingly, another aspect is a method of inhibiting Wnt ligand binding to a LRP5 or Wnt induced transcriptional activity comprising contacting one or more cells expressing one or more LRP5 polypeptides with an effective amount of an antibody or immunoconjugate described herein.

[00193] In an embodiment, the antibody or immunoconjugate comprises a CDR sequence set (full, light chain or heavy chain) corresponding to an antibody as described herein. As demonstrated herein, a number of the antibodies identified preferentially recognize LRP5. Accordingly, in embodiments wherein the method is for detecting LRP5 expression, the antibody or immunoconjugate comprises a CDR sequence set corresponding to an antibody selected from LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[00194] The contacting can for example, be done *in vivo* by administering an antibody or immunoconjugate to the subject. Such inhibition may be desirable particularly where Wnt signaling is dysregulated as in cancer cells.

C. Methods of potentiating WNT signaling pathways

[00195] Certain antibodies that block binding of Wnt ligands to the Wnt3a binding site of LRP5 or LRP6 potentiate the signaling activity of Wnt ligands binding to the non-Wnt3a binding site. Similarly, certain antibodies that block binding of Wnt ligands to the non-Wnt3a binding site of LRP5 or LRP6 potentiate the signaling activity of Wnt ligands binding to the Wnt3a binding site. The blocking activity may be competitive or allosteric. Accordingly, provided herein are methods of potentiating the signaling activity of Wnt-ligand binding to a Wnt3a binding site of LRP5 or LRP6 by contacting LRP5 or LRP6 with an antibody that blocks binding of Wnt ligands to the non-Wnt3a binding site of LRP5 or LRP6. Also provided herein are provided herein are methods of potentiating the signaling activity of Wnt-ligand binding to a non-Wnt3a binding site of LRP5 or LRP6 by contacting LRP5 or LRP6 with an antibody that block binding of Wnt ligands to the Wnt3a binding site of LRP5 or LRP6. Contacting can be performed *in vitro* or *in vivo*. *In vivo* contacting can comprise administering to the subject the appropriate anti-LRP5 or anti-LRP6 antibody.

[00196] For example, antibodies comprising CDR sequence sets from antibodies of Epitope Group 1 potentiate the activity of Wnt ligands binding to a non-Wnt3A binding site.

D. Methods and Uses for Treating Cancer

[00197] Methods of treating cancer comprise use of or administering to a subject in need thereof a pharmaceutical composition comprising an antibody of this disclosure that binds to LRP5. The subject in thereof can be a subject, e.g., a person, suffering from cancer, or at risk of cancer, such as recurrence of cancer.

[00198] In one embodiment, the cancer is selected from acute myeloid leukemia, prostate cancer, glioblastoma, bladder cancer and cervical cancer.

[00199] In another embodiment, the cancer cells are selected from brain cancer, breast cancer, colon cancer, endometrial cancer, esophageal cancer, kidney cancer, liver cancer, lung cancer, ovarian cancer, skin cancer, stomach cancer and testicular cancer.

[00200] Without wishing to be limited by theory, such therapy may function by inhibiting activation of the canonical Wnt pathway, for example, by inhibiting Wnt binding to LRP5, by inhibiting Wnt-induced transcriptional activity, by inhibiting activation of disheveled, by inhibiting inhibition of the beta-catenin destruction complex and by promoting accumulation of beta-catenin.

[00201] As LRP5 and/or LRP6 are often upregulated in cancer, the disclosure in another aspect includes a method for treating cancer, the method comprising administering an effective amount of an antibody or immunoconjugate that specifically binds LRP5 or LRP6 in at least one assay, and inhibits Wnt3a-induced signalling in at least one assay to a subject in need thereof. The disclosure also includes use of an effective amount of an antibody or immunoconjugate that specifically binds LRP5 or LRP6 in at least one assay, and inhibits Wnt3a-induced signalling in at least one assay for treating cancer or in the manufacture of a medicament for treating cancer. The disclosure further includes an effective amount of an antibody or immunoconjugate that specifically binds LRP5 or LRP6 in at least one assay, and inhibits Wnt3a-induced signalling in at least one assay for use in treating cancer.

[00202] In an embodiment, antibody or immunoconjugate, e.g., an antibody-drug conjugate, is comprised in a pharmaceutical composition.

[00203] In an embodiment, the cancer is selected from brain cancer, breast cancer, colon cancer, endometrial cancer, esophageal cancer, kidney cancer, liver cancer, lung cancer, ovarian cancer, skin cancer, stomach cancer and testicular cancer. In an embodiment, the antibody or immunoconjugate comprises a CDR sequence set corresponding to an antibody as described herein. A number of the antibodies identified preferentially recognize LRP5. Accordingly, in embodiments wherein the method is for detecting LRP5 expression, the antibody or immunoconjugate comprises a CDR sequence set corresponding to an antibody selected from LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[00204] As demonstrated herein, the antibodies are also able to inhibit cancer cell proliferation. Accordingly, also provided is a method for inhibiting cancer cell proliferation comprising contacting one or more cancer cells expressing an LRP5 with an effective amount of an antibody or immunoconjugate that specifically binds LRP5 in at least one assay, and inhibits Wnt3a-induced signalling in at least one assay.

[00205] In one embodiment, a method of treating cancer comprises administering antibodies or immunoconjugate comprising antibody, wherein a first antibody blocks binding of a Wnt ligand to a Wnt3a binding site of LRP5 or LRP6, and a second antibody blocks binding of a Wnt ligand to a non-Wnt3a binding site of the LRP5 or LRP6 to a subject in need thereof. The disclosure also provided antibodies or immunoconjugate comprising antibody, wherein a first antibody blocks binding of a Wnt ligand to a Wnt3a binding site of LRP5 or LRP6, and a second antibody blocks binding of a Wnt ligand to a non-Wnt3a binding site of the LRP5 or LRP6 for use in treating cancer. The disclosure further provides a use of antibodies or

immunoconjugate comprising antibody, wherein a first antibody blocks binding of a Wnt ligand to a Wnt3a binding site of LRP5 or LRP6, and a second antibody blocks binding of a Wnt ligand to a non-Wnt3a binding site of the LRP5 or LRP6 for treating cancer. The disclosure yet also provides a use of antibodies or immunoconjugate comprising antibody, wherein a first antibody blocks binding of a Wnt ligand to a Wnt3a binding site of LRP5 or LRP6, and a second antibody blocks binding of a Wnt ligand to a non-Wnt3a binding site of the LRP5 or LRP6 in the manufacture of a medicament for treating cancer.

[00206] In an embodiment, the antibody or immunoconjugate is the antibody or immunoconjugate described herein, for example, an antibody or immunoconjugate that comprises a CDR sequence set (full, light chain or heavy chain) corresponding to an antibody selected from LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

IX. Methods of Administration

[00207] The anti-LRP antibodies of the invention can efficiently deliver a therapeutic composition to cells undergoing Wnt signaling *in vivo*. In some embodiments, the method of treatment comprises administering to an individual an effective amount of a therapeutic anti-LRP conjugate, e.g., an anti-LRP antibody attached to a therapeutic agent. In some embodiments, the individual has been diagnosed with cancer. In some embodiments, the individual is receiving or has received cancer therapy, e.g., surgery, radiotherapy, or chemotherapy. In some embodiments, the individual has been diagnosed, but the cancer is in remission.

[00208] In some embodiments, the anti-LRP conjugate includes a liposome. In some embodiments, the method further comprises monitoring the individual for progression of the cancer. In some embodiments, the dose of the anti-LRP conjugate for each administration is determined based on the therapeutic progress of the individual, e.g., where a higher dose of chemotherapeutic is administered if the individual is not responding sufficiently to therapy.

[00209] In some embodiments, the invention can include an antibody or antibody-targeted composition and a physiologically (i.e., pharmaceutically) acceptable carrier. The term “carrier” refers to a typically inert substance used as a diluent or vehicle for a diagnostic or therapeutic agent. The term also encompasses a typically inert substance that imparts cohesive qualities to the composition. Physiologically acceptable carriers can be liquid, e.g., physiological saline, phosphate buffer, normal buffered saline (135-150 mM NaCl), water, buffered water, 0.4% saline, 0.3% glycine, glycoproteins to provide enhanced stability (e.g., albumin, lipoprotein, globulin, etc.), and the like. Since physiologically acceptable carriers are

determined in part by the particular composition being administered as well as by the particular method used to administer the composition, there are a wide variety of suitable formulations of pharmaceutical compositions of the present invention (See, e.g., Remington's Pharmaceutical Sciences, 17th ed., 1989).

[00210] The compositions of the present invention may be sterilized by conventional, well-known sterilization techniques or may be produced under sterile conditions. Aqueous solutions can be packaged for use or filtered under aseptic conditions and lyophilized, the lyophilized preparation being combined with a sterile aqueous solution prior to administration. The compositions can contain pharmaceutically acceptable auxiliary substances as required to approximate physiological conditions, such as pH adjusting and buffering agents, tonicity adjusting agents, wetting agents, and the like, e.g., sodium acetate, sodium lactate, sodium chloride, potassium chloride, calcium chloride, sorbitan monolaurate, and triethanolamine oleate. Sugars can also be included for stabilizing the compositions, such as a stabilizer for lyophilized antibody compositions.

[00211] Dosage forms can be prepared for mucosal (e.g., nasal, sublingual, vaginal, buccal, or rectal), parenteral (e.g., subcutaneous, intravenous, intramuscular, or intraarterial injection, either bolus or infusion), oral, or transdermal administration to a patient. Examples of dosage forms include, but are not limited to: dispersions; suppositories; ointments; cataplasms (poultices); pastes; powders; dressings; creams; plasters; solutions; patches; aerosols (e.g., nasal sprays or inhalers); gels; liquid dosage forms suitable for oral or mucosal administration to a patient, including suspensions (e.g., aqueous or non-aqueous liquid suspensions, oil-in-water emulsions, or a water-in-oil liquid emulsions), solutions, and elixirs; liquid dosage forms suitable for parenteral administration to a patient; and sterile solids (e.g., crystalline or amorphous solids) that can be reconstituted to provide liquid dosage forms suitable for parenteral administration to a patient.

[00212] Injectable (e.g., intravenous) compositions can comprise a solution of the antibody or antibody-targeted composition suspended in an acceptable carrier, such as an aqueous carrier. Any of a variety of aqueous carriers can be used, e.g., water, buffered water, 0.4% saline, 0.9% isotonic saline, 0.3% glycine, 5% dextrose, and the like, and may include glycoproteins for enhanced stability, such as albumin, lipoprotein, globulin, etc. Often, normal buffered saline (135-150 mM NaCl) will be used. The compositions can contain pharmaceutically acceptable auxiliary substances to approximate physiological conditions, such as pH adjusting and buffering agents, tonicity adjusting agents, wetting agents, e.g., sodium acetate, sodium lactate, sodium chloride, potassium chloride, calcium chloride, sorbitan monolaurate, triethanolamine oleate, etc. In some embodiments, the antibody-targeted composition can be formulated in a kit for intravenous administration.

[00213] Formulations suitable for parenteral administration, such as, for example, by intraarticular (in the joints), intravenous, intramuscular, intratumoral, intradermal, intraperitoneal, and subcutaneous routes, include aqueous and non-aqueous, isotonic sterile injection solutions, which can contain antioxidants, buffers, bacteriostats, and solutes that render the formulation isotonic with the blood of the intended recipient, and aqueous and non-aqueous sterile suspensions that can include suspending agents, solubilizers, thickening agents, stabilizers, and preservatives. Injection solutions and suspensions can also be prepared from sterile powders, granules, and tablets. In the practice of the present invention, compositions can be administered, for example, by intravenous infusion, topically, intraperitoneally, intravesically, or intrathecally. Parenteral administration and intravenous administration are the preferred methods of administration. The formulations of targeted compositions can be presented in unit-dose or multi-dose sealed containers, such as ampoules and vials.

[00214] The targeted delivery composition of choice, alone or in combination with other suitable components, can be made into aerosol formulations ("nebulized") to be administered via inhalation. Aerosol formulations can be placed into pressurized acceptable propellants, such as dichlorodifluoromethane, propane, and nitrogen.

[00215] The pharmaceutical preparation can be packaged or prepared in unit dosage form. In such form, the preparation is subdivided into unit doses containing appropriate quantities of the active component, e.g., according to the dose of the therapeutic agent or concentration of antibody. The unit dosage form can be a packaged preparation, the package containing discrete quantities of preparation. The composition can, if desired, also contain other compatible therapeutic agents.

[00216] The antibody (or antibody- targeted composition) can be administered by injection or infusion through any suitable route including but not limited to intravenous, subcutaneous, intramuscular or intraperitoneal routes. An example of administration of a pharmaceutical composition includes storing the antibody at 10 mg/ml in sterile isotonic aqueous saline solution for injection at 4°C, and diluting it in either 100 ml or 200 ml 0.9% sodium chloride for injection prior to administration to the patient. The antibody is administered by intravenous infusion over the course of 1 hour at a dose of between 0.2 and 10 mg/kg. In other embodiments, the antibody is administered by intravenous infusion over a period of between 15 minutes and 2 hours. In still other embodiments, the administration procedure is via sub-cutaneous bolus injection.

[00217] The dose of antibody is chosen in order to provide effective therapy for the patient and is in the range of less than 0.1 mg/kg body weight to about 25 mg/kg body weight or in

the range 1 mg- 2 g per patient. In some cases, the dose is in the range 1- 100 mg/kg, or approximately 50 mg- 8000 mg / patient. The dose may be repeated at an appropriate frequency which may be in the range once per day to once every three months, depending on the pharmacokinetics of the antibody (e.g., half-life of the antibody in the circulation) and the pharmacodynamic response (e.g., the duration of the therapeutic effect of the antibody). In some embodiments, the *in vivo* half-life of between about 7 and about 25 days and antibody dosing is repeated between once per week and once every 3 months.

[00218] Administration can be periodic. Depending on the route of administration, the dose can be administered, e.g., once every 1, 3, 5, 7, 10, 14, 21, or 28 days or longer (e.g., once every 2, 3, 4, or 6 months). In some cases, administration is more frequent, e.g., 2 or 3 times per day. The patient can be monitored to adjust the dosage and frequency of administration depending on therapeutic progress and any adverse side effects, as will be recognized by one of skill in the art.

[00219] Thus, in some embodiments, additional administration is dependent on patient progress, e.g., the patient is monitored between administrations. For example, after the first administration or round of administrations, the patient can be monitored for rate of tumor growth, recurrence (e.g., in the case of a post-surgical patient), or general disease-related symptoms such as weakness, pain, nausea, etc.

[00220] In therapeutic use for the treatment of cancer, an antibody-targeted composition (e.g., including a therapeutic and/or diagnostic agent) can be administered at the initial dosage of about 0.001 mg/kg to about 1000 mg/kg daily and adjusted over time. A daily dose range of about 0.01 mg/kg to about 500 mg/kg, or about 0.1 mg/kg to about 200 mg/kg, or about 1 mg/kg to about 100 mg/kg, or about 10 mg/kg to about 50 mg/kg, can be used. The dosage is varied depending upon the requirements of the patient, the severity of the condition being treated, and the targeted composition being employed. For example, dosages can be empirically determined considering the type and stage of cancer diagnosed in a particular patient. The dose administered to a patient, in the context of the present invention, should be sufficient to affect a beneficial therapeutic response in the patient over time. The size of the dose will also be determined by the existence, nature, and extent of any adverse side-effects that accompany the administration of a particular targeted composition in a particular patient, as will be recognized by the skilled practitioner.

[00221] The above disclosure generally describes the present disclosure. A more complete understanding can be obtained by reference to the following specific examples. These examples are described solely for the purpose of illustration and are not intended to limit the scope of the application. Changes in form and substitution of equivalents are contemplated as

circumstances might suggest or render expedient. Although specific terms have been employed herein, such terms are intended in a descriptive sense and not for purposes of limitation.

EXEMPLARY EMBODIMENTS

[00222] 1. An antibody that specifically binds LRP5, comprising a light chain variable region and/or a heavy chain variable region, the heavy chain variable region comprising complementarity determining regions CDR-H1, CDR-H2 and CDR-H3, the light chain variable region comprising complementarity determining region CDR-L1, CDR-L2 and CDR-L3, wherein the amino acid sequences of said CDRs comprise or consist of CDR sequences selected from: CDR sequence sets of anti-LRP5 antibodies: LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[00223] 2. The antibody of embodiment 1 wherein the amino acid sequences of said CDRs comprise or consist of sequences selected from the sequences as set forth below:

CDR-H1 is selected from the group consisting of LSYYYM (SEQ ID NO: 7), ISYSYI (SEQ ID NO: 5), LSYSSM (SEQ ID NO: 2), ISSYSI (SEQ ID NO: 3), ISYSYI (SEQ ID NO: 5), IYSYSI (SEQ ID NO: 6), LSYYYM (SEQ ID NO: 7), FSSSSI (SEQ ID NO: 4), LYYYYI (SEQ ID NO: 8), LSYSSI (SEQ ID NO: 9), IYSYYI (SEQ ID NO: 10), LLYYSSM (SEQ ID NO: 122) and FSSSSI (SEQ ID NO: 4);

CDR-H2 is selected from the group consisting of SIYPYYGYTY (SEQ ID NO: 11), SSSYYGYTY (SEQ ID NO: 12), SISSSYGYTY (SEQ ID NO: 13), SIYSSYGSTS (SEQ ID NO: 14), SIYSSYGYTY (SEQ ID NO: 15), SIYPYSSYTS (SEQ ID NO: 16), SIYSSYGYTY (SEQ ID NO: 15), SIYPSYGYTY (SEQ ID NO: 17), SISPYYGYTS (SEQ ID NO: 18), SISSSYGSTS (SEQ ID NO: 19), SIYSYYGYTY (SEQ ID NO: 20), SISSSYGYTY (SEQ ID NO: 13), SISSSYGYTY (SEQ ID NO: 13) SISSYYGYTS (SEQ ID NO: 53), and YISPYYGYTS (SEQ ID NO: 56);

CDR-H3 is selected from the group consisting of HGAM (SEQ ID NO: 21), TVRGSKKPYFSGWAM (SEQ ID NO: 22), SSYYSSVSSSVYAL (SEQ ID NO: 23), TVRGSKKPYFSGWAM (SEQ ID NO: 22), HYSYFFYAM (SEQ ID NO: 24), YAVYFPGYYWGM (SEQ ID NO: 25), WSHVSGHYSGM (SEQ ID NO: 26), WGAYHSSGYGM (SEQ ID NO: 27), GGSGVSHYGSVYYSWWAL (SEQ ID NO:

28), AAPYYGYYYSYAM (SEQ ID NO: 29), SGYGWYAM (SEQ ID NO: 30), GYWAI (SEQ ID NO: 31), SYPAM (SEQ ID NO: 32), SWAM (SEQ ID NO: 33); YWAL (SEQ ID NO: 54), GWGSPASAGYYGL (SEQ ID NO: 57), SSYYSSVSSSVYAL (SEQ ID NO: 23), TVRGSKKPYFSGWAM (SEQ ID NO: 22) and TVRGSKKPYFSGWAM (SEQ ID NO: 22);

CDR-L1 is SVSSA (SEQ ID NO: 34);

CDR-L2 is SASSLYS (SEQ ID NO: 35); and

[00224] CDR-L3 is selected from the group consisting of AWGWGLF (SEQ ID NO: 36), VHYSPYSLI (SEQ ID NO: 37), YQYSGLI (SEQ ID NO: 38), FSHVSLI (SEQ ID NO: 39), ASYSPI (SEQ ID NO: 40), YHYYYLF (SEQ ID NO: 41), ASYAPI (SEQ ID NO: 42), SSSSPI (SEQ ID NO: 43), SSYSLI (SEQ ID NO: 44), GVSLI (SEQ ID NO: 45), YWFLI (SEQ ID NO: 46), PVGHYGYPI (SEQ ID NO: 47), SSYSPI (SEQ ID NO: 48), YWAYYSPI (SEQ ID NO: 49), VSYYPLI (SEQ ID NO: 51), SSYSLI (SEQ ID NO: 44), and VHYSPYSLI (SEQ ID NO: 37).

[00225] 3. The antibody of embodiment 2, wherein the antibody comprises a heavy chain variable region comprising:

i) a heavy chain amino acid sequence as set forth in Table 2;

ii) an amino acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the heavy chain amino acid sequence as set forth in Table 2, wherein the CDR sequences are a CDR sequence set as set forth in Table 1, or

iii) a conservatively substituted amino acid sequence of i) wherein the CDR sequences are a CDR sequence set as set forth in Table.

[00226] 4. The antibody of any one of embodiments 2 to 4, wherein the antibody comprises a light chain variable region comprising:

i) a light chain amino acid sequence as set forth in Table 2,

ii) an amino acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the light chain amino acid sequence as set forth in Table 2, wherein the CDR sequences are a CDR sequence set as set forth in Table 1, or

iii) a conservatively substituted amino acid sequence of i) wherein the CDR sequences are a CDR sequence set as set forth in Table 1.

[00227] 5. The antibody of any one of embodiments 1 to 4, wherein the CDR sequences are a full CDR sequence set selected from an antibody identified in Table 1.

[00228] 6. The antibody of any one of embodiments 1 to 5, wherein the antibody cross-reacts with LRP6.

[00229] 7. The antibody of embodiment 1, wherein the CDR sequences comprise a light chain CDR sequence set or a heavy chain CDR sequence set selected from an antibody identified in Table 1.

[00230] 8. The antibody of any of embodiments 1 to 7, wherein the antibody specifically binds LRP5.

[00231] 9. The antibody of embodiment 8, wherein the CDR sequences are a CDR sequence set of an antibody selected from antibodies LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[00232] 10. The antibody of embodiments 1 to 9, which blocks binding of a Wnt ligand to a Wnt3a binding site of LRP5.

[00233] 11. The antibody of embodiments 1 to 9, blocks binding of a Wnt ligand to a non-Wnt3a binding site of LRP5.

[00234] 12. The antibody of any one of embodiments 1 to 11, wherein the antibody is a monoclonal antibody.

[00235] 13. The antibody of any one of embodiments 1 to 12, wherein the antibody is a humanized antibody.

[00236] 14. The antibody of any one of embodiments 1 to 13, wherein the antibody is a single chain antibody.

[00237] 15. The antibody of any one of embodiments 1 to 14, wherein the antibody is an antibody binding fragment selected from Fab, Fab', F(ab')2, scFv, dsFv, ds-scFv, dimers, nanobodies, minibodies, diabodies, and multimers thereof.

[00238] 16. The antibody of any one of embodiments 1 to 14, wherein the antibody is a bi-specific antibody.

[00239] 17. The antibody of any one of embodiments 1 to 14, wherein the antibody is a bi-specific antibody that further binds to FZD receptor.

[00240] 18. The antibody of any one of embodiments 1 to 17, comprising a non-natural glycosylation pattern.

[00241] 19. The antibody of any one of embodiments 1 to 17, comprising a cysteine substitution or addition, e.g., in the constant region or a framework region.

[00242] 20. An immunoconjugate comprising the antibody of any one of embodiments 1 to 17, and a detectable label or cytotoxic agent.

[00243] 21. The immunoconjugate of embodiment 20, comprising a cytotoxic agent selected from maytansinoid, auristatin, dolastatin, tubulysin, cryptophycin, pyrrolobenzodiazepine (PBD) dimer, indolinobenzodiazepine dimer, alpha-amanitin, trichothene, SN-38, duocarmycin, CC1065, calicheamincin, an enediyne antibioatic, taxane, doxorubicin derivatives, anthracycline and stereoisomers, azanofide, isosteres, analogs or derivatives thereof.

[00244] 22. A nucleic acid molecule encoding the antibody of any one of embodiments 1 to 17.

[00245] 23. The nucleic acid molecule of embodiment 22, wherein one or more of the CDR sequences is/are encoded by a nucleic acid in Table 2.

[00246] 24. The nucleic acid molecule of embodiment 22, wherein the antibody comprises a heavy chain variable region encoded by a nucleic acid comprising:

i) a heavy chain nucleic acid sequence as set forth in Table 2;

ii) a nucleotide sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the heavy chain nucleic acid sequence as set forth in Table 2, wherein the CDR sequences are a CDR sequence set as set forth in Table 1, or

iii) a codon degenerate nucleic acid sequence of i) wherein the CDR sequences are a CDR sequence set as set forth in Table 1.

[00247] 25. The nucleic acid molecule of embodiment 22, wherein the antibody comprises a light chain variable region encoded by a nucleic acid comprising:

i) a light chain nucleic acid sequence as set forth in Table 2,

ii) a nucleic acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, or at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the light chain nucleic acid sequence as set forth in Table 2, wherein the CDR sequences are a CDR sequence set as set forth in Table 1, or

iii) a codon degenerate nucleic acid sequence of i) wherein the CDR sequences are a CDR sequence set as set forth in Table 1.

[00248] 26. A vector comprising an expression control sequence operatively linked to the nucleic acid of any one of embodiments 22 to 25.

[00249] 27. A host cell comprising recombinant nucleic acid molecule comprising an expression control sequence operatively linked to the nucleic acid of any of embodiments 22-26.

[00250] 28. The host cell of embodiment 27, that is a Chinese Hamster Ovary (CHO) cell.

[00251] 29. A host cell comprising the vector of embodiment 26.

[00252] 30. A method for making an anti-LRP5 antibody comprising culturing a host cell of any one of embodiments 27 to 29.

[00253] 31. A composition comprising the antibody of any one or more of embodiments 1 to 17, immunoconjugate of embodiments 20-21, the nucleic acid molecule of embodiments 22-25, the vector of embodiment 26, or host cell of embodiment 29, optionally with a suitable diluent.

[00254] 32. The composition of embodiment 31, wherein the composition comprises one or more antibodies or immunoconjugates, optionally wherein the composition is a pharmaceutical composition.

[00255] 33. A kit comprising the antibody of any one or more of embodiments 1 to 17, immunoconjugate of embodiments 20-21, the nucleic acid molecule of embodiments 22-25, the vector of embodiment 26, or host cell of embodiments 29-29.

[00256] 34. A method of detecting LRP5 expression, the method comprising contacting a sample comprising one or more cells with one or more antibody or immunoconjugate of any one of embodiments 1 to 21 under conditions permissive for forming an antibody:cell complex and detecting the presence of any antibody complex.

[00257] 35. The method of embodiment 34, wherein the detection is by immunofluorescence.

[00258] 36. The method of embodiment 34, wherein the detection is by flow cytometry.

[00259] 37. The method of any one of embodiments 34 to 36, wherein the method is for detecting LRP4 expression and the antibody or immunoconjugate comprises a CDR sequence set corresponding to an antibody selected from LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[00260] 38. A method of inhibiting Wnt ligand binding to an LRP5 receptor, disrupting a Wnt signaling pathway, inhibiting Wnt-induced transcriptional activity, inhibiting activation of

disheveled, promoting preservation of the beta-catenin destruction complex, promoting accumulation of beta-catenin or inhibiting growth of a cell, the method comprising contacting a cell expressing a LRP5 receptor with an antibody or immunoconjugate of any one of embodiments 1 to 21.

[00261] 39. The method of embodiment 38, wherein the antibody or immunoconjugate blocks binding of a Wnt ligand to a Wnt3a binding site of LRP5.

[00262] 40. The method of embodiment 38, wherein the antibody or immunoconjugate blocks binding of a Wnt ligand to a non-Wnt3a binding site of LRP5.

[00263] 41. The method of embodiment 38, wherein the antibody or immunoconjugate comprises a CDR sequence set corresponding to an antibody selected from the group consisting of LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[00264] 42. A method of treating cancer in a subject in need thereof comprising administering to the subject an effective amount of a pharmaceutical composition comprising an antibody or an immunoconjugate of any one of embodiments 1 to 21.

[00265] 43. The method of embodiment 42, wherein the cancer is selected from colon, lung, breast ovarian, endometrial, pancreas, stomach, liver, adrenocortical carcinoma and osteoblastoma cancer cells.

[00266] 44. The method of embodiment 42, wherein the cancer is selected from acute myeloid leukemia, prostate cancer, glioblastoma, bladder cancer and cervical cancer.

[00267] 45. The method of embodiment 42, comprising administering to the subject first and second antibodies or antibody conjugates of any one of embodiments 1 to 21, wherein the first blocks binding of a Wnt ligand to a Wnt3a binding site of LRP5, and the second blocks binding of a Wnt ligand to a non-Wnt3a binding site of LRP5.

[00268] 46. The method of embodiment 45, wherein the first antibody or immunoconjugate comprises a CDR sequence set selected from antibodies of Epitope Group 2.

[00269] 47. The method of embodiment 42, wherein the antibody or immunoconjugate that specifically binds LRP5 in at least one assay, and inhibits Wnt3a- induced signaling in at least one assay, optionally wherein the antibody or immunoconjugate is the antibody or immunoconjugate of any one of embodiments 1 to 21.

[00270] 48. The method of embodiment 42, wherein the antibody or immunoconjugate comprises a CDR sequence set corresponding to an antibody selected from the group

consisting of LRP5 - A7, LRP5 - A9, LRP5 - C5, LRP5 - C12, LRP5 - D9, LRP5 - E5, LRP5 - G2, LRP5 - G9, LRP5 - G10, LRP5 - G11, LRP5 - H3, LRP5 - H5, LRP5 - H9, LRP5 - R3O_D3, LRP5 - R3_E8, LRP5 - R3O_G6.

[00271] 49. A method of potentiating the signaling activity of Wnt-ligand binding to a Wnt3a binding site of LRP5 comprising contacting a cell expressing LRP5 with an antibody that blocks binding of Wnt ligands to the non-Wnt3a binding site of LRP5.

[00272] 50. A method of potentiating the signaling activity of Wnt-ligand binding to a non-Wnt3a binding site of LRP5 by contacting a cell expressing LRP5 with an antibody that blocks binding of Wnt ligands to the Wnt3a binding site of LRP5.

[00273] 51. The method of any of embodiments 49 or 50, performed *in vitro*.

[00274] 52. The method of any of embodiments 49 or 50, performed *in vivo*.

EXAMPLES

[00275] The following non-limiting examples are illustrative of the present disclosure:

[00276] Example 1

[00277] Phage display antibody library

[00278] Phage-displayed antibody libraries are a powerful technology for the generation of therapeutic antibodies[17,18]. Highly complex libraries of > 10^{10} independent antibody fragments are displayed on phage particles as coat protein fusion molecules and screened to isolate antibodies that recognize antigens of interest. The inventors have established synthetic antibody libraries with antigen-binding sites constructed entirely from engineered sequences. The resulting antibodies use an optimized human framework, and are thus minimally immunogenic when used as potential therapeutics. The synthetic antibodies are highly stable, and their human framework and antigen combining sites can be tailored to optimize affinity, specificity and efficacy.

[00279] Characterization and optimization of anti-LRP5 antibodies

[00280] Example 2

[00281] Novel synthetic antibodies targeting LRP5, a highly optimized Fab-phage library constructed in the lab was used to generate specific anti-LRP5 antibodies by directly selecting on immobilized recombinant LRP5 ECD. Fab-phage clones from rounds 3 and 4 were screened for binding by ELISA and the amino acid sequences of the CDRs of the positive binders are present in Figure 1A. The complete DNA and amino acid sequences of the variable region (heavy and light chains) of the lgGs are presented in the specification above. Epitope

mapping by competitive ELISA reveal that the antibodies bind to four unique epitopes on the ECO of LRP5.

[00282] The full length lgG1s and the antibody fragments (Fabs) were purified and a single-point ELISA was done to determine the specificity of the antibodies. As shown in Figure 1B, all the LRP5 antibodies bind to the recombinant mouse LRP5-His chimera. Interestingly, LRP5-H5 and LRP5-R30D3 antibodies show partial binding to the recombinant mouse LRP6-His chimera. LRP5-R30D3 antibody also shows partial binding to the recombinant human LRP6-Fc chimera. LRP5-R3OD3 antibody also shows partial binding to the recombinant human LRP6-Fc chimera suggesting that this antibody could cross-react with both LRP5 and LRP6. A multipoint competitive ELISA was used to determine the relative affinities of the antibodies to the recombinant antigen. The IC50 was determined by non-linear regression analysis and is summarized in Table 3. The complete dose response curves and the non-linear regression plots are presented in Figure 6.

[00283] Table 3 reports the IC50 of LRP5 antibodies as determined by competitive ELISA. The log of the recombinant human LRP6-Fc chimera concentration (x-axis) was plotted against the OD450 reading of the antibody (y-axis). The IC50 was determined by non-linear regression analysis.

Table 3

IgG1 antibody	IC_{50} (nM)
LRP5-A7	4.68
LRP5-D9	25.11
LRP5-G2	24.21
LRP5-G9	26.44
LRP5-G10	3.02
LRP5-G11	25.11
LRP5-H5	4.44
LRP5-H9	37.46

[00284] Western blot analysis of whole cell lysates has shown that LRP5 is highly expressed in the NSCLC cell line, H23, the triple negative breast cancer cell line, MOAMB231, and in the breast cancer cell line, T470 (data not shown). The LRP5 lgG1s label the surfaces

of H23 (Figure 2), MOAMB231 (Appendix 3) and T470 (Appendix 4) cells as assessed by FACS analysis and demonstrate that the LRP5 antibodies can bind to the full-length receptor expressed on the surface of these cell lines. As mentioned above, Wnt signaling initiates the canonical pathway, which primarily regulates beta-catenin stability and function.

[00285] Upon Wnt stimulation, beta-catenin levels in the cytosol and nucleus rise resulting in an increase in TCF/LEF (transcription factors) mediated transcription. The TOPflash reporter assay is a widely used tool to assess beta-catenin activity *in vitro*. The vector consists of several TCF/LEF binding sites that drive the expression of the firefly luciferase reporter gene. Several cancer cell lines expressing this reporter were generated in order to assess the functional activity of the LRP5 antibodies in regulating the Wnt-canonical pathway.

[00286] As shown in Figure 3A, conditioned media expressing Wnt3a (Wnt3aCM) induces a 120-fold increase in luciferase reporter activity compared to treatment with control conditioned media (ConCM) in MOAMB231 cells pre-treated with the negative control lgG1, anti-MBP. LRP5-G10 and LRP5-H5 antibodies significantly potentiated the Wnt3aCM-induced reporter activity. A similar result was observed in the breast cancer cell line, T47D (Figure 38) and the osteosarcoma cell line, U20S (Figure 3C). In the U20S cell line, both the ConCM- and Wnt3aCM-induced reporter activity was potentiated by LRP5-G10 and LRP5-H5 antibodies (compared to M8P).

[00287] Structural and mutagenesis studies have revealed that Wnt3a binds to a domain that is unique from the binding site of the remaining Wnt ligands. Moreover, recent studies have shown that certain Wnt ligands require both LRP5 and LRP6 to initiate the canonical pathway. To address the hypothesis that epitope specific LRP5 antibodies will regulate beta-catenin mediated transcriptional activity in a Wnt-dependent manner, we assessed reporter activity in the NSCLC cell line, H23. Previous studies (20) have shown that Wnt2 primarily drives the basal TCF/LEF mediated transcriptional activity and these cells do not express Wnt3a. As expected, LRP5-G10 and LRP5-H5 antibodies potentiated the Wnt3aCM induced reporter activity (compared to M8P; Figure 3D). Interestingly, these antibodies (as well as LRP5-A7, another member of this epitope group) inhibited the basal (ConCM-induced) reporter activity (compared to M8P). Moreover, LRP5-D9 and LRP5-G2 antibodies significantly potentiate the ConCM-induced reporter activity (compared to MBP). Our results show that LRP5-G2 and LRP5-G10 antibodies induce the most potent effect on reporter activity. Consistent with this observation, LRP5-G2 antibody potentiates the ConCM induced reporter activity in a dose-dependent manner (Figure 4A). Interestingly, LRP5-G10 antibody inhibits ConCM-induced reporter activity at all the indicated concentrations. LRP5-G10 Fab also inhibits ConCM-induced reporter activity in at the indicated concentrations while LRP5-G2 Fab fails to recapitulate the effect of the lgG1 on reporter activity (Figure 4B).

[00288] The effects of LRP5-G2 and LRP5-G10 antibodies on the proximal events of Wnt signaling were assessed by western blot analysis (Figure 5A). LRP5-G10 but not LRP5-G2 antibody significantly inhibited both ConCM- and Wnt3aCM-induced LRP6 phosphorylation in H23 cells. LRP5-G10 antibody also significantly reduced total LRP6 phosphorylation in H23 cells. A similar effect on LRP6 phosphorylation is observed in membrane fractions isolated from H23 cells treated with LRP5-G2 and LRP5-G10 antibodies prior to stimulation with conditioned media (Figure 5B). LRP5-G10 antibody also significantly reduced the Axin1 protein levels in Wnt3aCM-stimulated cells. Since Axin1 is a component of the destruction complex that regulates the "free" pool of beta-catenin level, its diminished expression upon treatment with LRP5-G10 antibody prior to Wnt3aCM stimulation may explain why this antibody effectively potentiates the Wnt3aCM-induced reporter activity (Figures 3).

[00289] To get a better understanding of the effects of the antibodies on beta-catenin levels, cytosolic fractions of H23 cells treated with LRP5-G2 and LRP5-G10 antibodies prior to stimulation with conditioned media were isolated. As shown in Figure 5B, LRP5-G2 antibody significantly upregulates beta-catenin levels in ConCM-treated cells consistent with its effect on reporter activity. Similarly, LRP5-G10 antibody slightly decreases and increases beta-catenin levels in ConCM- treated cells.

[00290] The NSCLC cell line, H23, serves as an excellent system to explore the effects of co-treatment of the LRP5 antibodies on TOPflash reporter activity. Our previous results show that LRP5-G2 antibody potentiates the ConCM-induced reporter activity. In contrast, LRP5-G10 antibody inhibits and potentiates the ConCM-induced and Wnt3aCM-induced reporter activity, respectively. As shown in Figure 6, the potentiation of the ConCM-induced reporter activity by LRP5-G2 is significantly inhibited in the presence of LRP5-G10 antibody. Our observations suggest that the co-treatment of LRP5-G2 and LRP5-G10 antibodies can have a potent inhibitory effect on Wnt-stimulated TCF/LEF-mediated transcription. Thus, by taking advantage of advances in library design strategies, we have been able to discover novel LRP5 antibodies that can exert both antagonistic and potentiating activities on proximal and distal events related to beta-catenin signaling. These activities also depend on the different interactions between Wnt ligands and LRP5. We are actively investigating these antibodies for their therapeutic potential *in vitro* and *in vivo*.

[00291] Abbreviations:

Abbreviation	Complete term
APC	Adenomatous polyposis coli
CDR	Complementarity determining region

CK1	Casein kinase 1
Dsh/Dvl	Dishevelled
ECD	Extracellular domain
ELISA	Enzyme-linked immunosorbent assay
Fab	Fragment antigen-binding
FACS	Fluorescence-activated cell sorting
Fc	Fragment crystallizable region
GSK3b	Glycogen synthase kinase 3 beta
IC_{50}	Half maximal inhibitory concentration
IgG1	Immunoglobulin G 1
LRP	Low-density lipoprotein receptor-related protein
OD	Optical density
NSCLC	Non-small cell lung carcinoma

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[00312] As used herein, the following meanings apply unless otherwise specified. The word "may" is used in a permissive sense (i.e., meaning having the potential to), rather than the mandatory sense (i.e., meaning must). The words "include", "including", and "includes" and the like mean including, but not limited to. The singular forms "a," "an," and "the" include plural referents. Thus, for example, reference to "an element" includes a combination of two or more elements, notwithstanding use of other terms and phrases for one or more elements, such as "one or more." The phrase "at least one" includes "one or more", "one or a plurality" and "a plurality". The term "or" is, unless indicated otherwise, non-exclusive, i.e., encompassing both "and" and "or." The term "any of" between a modifier and a sequence means that the modifier modifies each member of the sequence. So, for example, the phrase "at least any of 1, 2 or 3" means "at least 1, at least 2 or at least 3". The term "consisting essentially of" refers to the inclusion of recited elements and other elements that do not materially affect the basic and novel characteristics of a claimed combination.

[00313] It should be understood that the description and the drawings are not intended to limit the invention to the particular form disclosed, but to the contrary, the intention is to cover all modifications, equivalents, and alternatives falling within the spirit and scope of the present invention as defined by the appended claims. Further modifications and alternative embodiments of various aspects of the invention will be apparent to those skilled in the art in view of this description. Accordingly, this description and the drawings are to be construed as illustrative only and are for the purpose of teaching those skilled in the art the general manner of carrying out the invention. It is to be understood that the forms of the invention shown and described herein are to be taken as examples of embodiments. Elements and materials may be substituted for those illustrated and described herein, parts and processes may be reversed or omitted, and certain features of the invention may be utilized independently, all as would be apparent to one skilled in the art after having the benefit of this description of the invention. Changes may be made in the elements described herein without departing from the spirit and scope of the invention as described in the following claims. Headings used herein are for organizational purposes only and are not meant to be used to limit the scope of the description.

[00314] All publications, patents, and patent applications mentioned in this specification are herein incorporated by reference to the same extent as if each individual publication, patent, or patent application was specifically and individually indicated to be incorporated by reference.

WHAT IS CLAIMED IS:

1. An antibody that specifically binds LRP5, comprising a light chain variable region and/or a heavy chain variable region, the heavy chain variable region comprising complementarity determining regions CDR-H1, CDR-H2 and CDR-H3, the light chain variable region comprising complementarity determining region CDR-L1, CDR-L2 and CDR-L3, wherein the amino acid sequences of said CDRs comprise or consist of CDR sequences as follows:

CDR-H1 is SEQ ID NO: 2,
CDR-H2 is SEQ ID NO: 11,
CDR-H3 is SEQ ID NO: 21,
CDR-L1 is SEQ ID NO: 34,
CDR-L2 is SEQ ID NO: 35, and
CDR-L3 is SEQ ID NO: 36.

2. The antibody of claim **1**, wherein the antibody comprises a heavy chain variable region comprising:

i) a heavy chain amino acid sequence of SEQ ID NO: 59;

ii) an amino acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the heavy chain amino acid sequence of any one of SEQ ID NOs: 59, 63, 67, 71, 75, 79, 83, 87, 91, 95, 99, 103, 107, 111, 115 and 119, wherein the CDR sequences are those of claim 1, or

iii) a conservatively substituted amino acid sequence of i), wherein the CDR sequences are those of claim 1.

3. The antibody of claim **1 or 2**, wherein the antibody comprises a light chain variable region comprising:

i) a light chain amino acid sequence of SEQ ID NO: 61,

ii) an amino acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the light chain amino acid sequence of any one of SEQ ID NOs: 61, 65, 69, 73, 77, 81, 85, 89, 93, 97, 101, 105, 109, 113, 117 and 121, wherein the CDR sequences are those of claim 1, or

iii) a conservatively substituted amino acid sequence of i), wherein the CDR sequences are those of claim 1.

4. The antibody of any one of claims **1** to **3**, wherein the antibody cross-reacts with LRP6.

5. The antibody of any one of claims **1** to **4**, wherein the antibody specifically binds LRP5.

6. The antibody of any one of claims **1** to **5**, blocks binding of a Wnt ligand to a non-Wnt3a binding site of LRP5.

7. The antibody of any one of claims **1** to **6,** wherein the antibody is a monoclonal antibody.

8. The antibody of any one of claims **1** to **7**, wherein the antibody is a humanized antibody.

9. The antibody of any one of claims **1** to **8**, wherein the antibody is a single chain antibody.

10. The antibody of any one of claims **1** to **9**, wherein the antibody is an antibody binding fragment selected from Fab, Fab', F(ab')2, scFv, dsFv, ds-scFv, dimers, nanobodies, minibodies, diabodies, and multimers thereof.

11. The antibody of any one of claims **1** to **9**, wherein the antibody is a bi-specific antibody.

12. The antibody of any one of claims **1** to **9**, wherein the antibody is a bi-specific antibody that further binds to FZD receptor.

13. The antibody of any one of claims **1** to **12**, comprising a non-natural glycosylation pattern.

14. The antibody of any one of claims **1** to **12**, comprising a cysteine substitution or addition, e.g., in the constant region or a framework region.

15. An immunoconjugate comprising the antibody of any one of claims **1** to **12**, and a detectable label or cytotoxic agent.

16. The immunoconjugate of claim **15**, comprising a cytotoxic agent selected from maytansinoid, auristatin, dolastatin, tubulysin, cryptophycin, pyrrolobenzodiazepine (PBD) dimer, indolinobenzodiazepine dimer, alpha-amanitin, trichothene, SN-38, duocarmycin, CC1065, calicheamincin, an enediyne antibioatic, taxane, doxorubicin derivatives, anthracycline and stereoisomers, azanofide or isosteres thereof.

17. A nucleic acid molecule encoding the antibody of any one of claims **1** to **12**.

18. The nucleic acid molecule of claim **17**, wherein one or more of the CDR sequences is/are encoded by a nucleic acid in Table 2.

19. The nucleic acid molecule of claim **17**, wherein the antibody comprises a heavy chain variable region encoded by a nucleic acid comprising:

i) a heavy chain nucleic acid sequence of SEQ ID NO: 58,

ii) a nucleotide sequence with at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the heavy chain nucleic acid sequence of any one of SEQ ID NOs: 58, 62, 66, 70, 74, 78, 82, 86, 90, 94, 98, 102, 106, 110, 114 and 118, wherein the CDR polypeptide sequences are that of claim 1, or

iii) a codon degenerate nucleic acid sequence of i) wherein the CDR polypeptide sequences are that of claim 1.

20. The nucleic acid molecule of claim **17 or claim 19**, wherein the antibody comprises a light chain variable region encoded by a nucleic acid comprising:

i) a light chain nucleic acid sequence of SEQ ID NO: 60,

ii) a nucleic acid sequence with at least 50%, at least 60%, at least 70%, at least 80%, or at least 90%, at least 95%, at least 98% or at least 99% sequence identity to the light chain nucleic acid sequence of any one of SEQ ID NOs: 60, 64, 68, 72, 76, 80, 84, 88, 92, 96, 100, 104, 108, 112, 116 and 120, wherein the CDR polypeptide sequences are that of claim 1, or

iii) a codon degenerate nucleic acid sequence of i) wherein the CDR polypeptide sequences are that of claim 1.

21. A vector comprising an expression control sequence operatively linked to the nucleic acid of any one of claims **17** to **20**.

22. A host cell comprising recombinant nucleic acid molecule comprising an expression control sequence operatively linked to the nucleic acid of any of claims **17** to **21**.

23. The host cell of claim **22**, that is a Chinese Hamster Ovary (CHO) cell.

24. A host cell comprising the vector of claim **21**.

25. A method for making an anti-LRP5 antibody comprising culturing a host cell of any one of claims **22** to **24**.

26. A composition comprising the antibody of any one or more of claims **1** to **12**, immunoconjugate of any one of claims **15** to **16**, the nucleic acid molecule of any one of claims **17** to **20**, the vector of claim **21**, or host cell of claim **24**, optionally with a suitable diluent.

27. The composition of claim **26**, wherein the composition comprises one or more antibodies or immunoconjugates, optionally wherein the composition is a pharmaceutical composition.

28. A kit comprising the antibody of any one or more of claims **1** to **12**, immunoconjugate of any one of claims **15** to **16**, the nucleic acid molecule of any one of claims **17** to **20**, the vector of claim **21**, or host cell of any one of claims **22** to **24**.

29. A method of detecting LRP5 expression, the method comprising contacting a sample comprising one or more cells with one or more antibody or immunoconjugate of any one of claims **1** to **16** under conditions permissive for forming an antibody:cell complex and detecting the presence of any antibody complex.

30. The method of claim **29**, wherein the detection is by immunofluorescence.

31. The method of claim **29**, wherein the detection is by flow cytometry.

32. Use of an antibody or immunoconjugate of any one of claims **1** to **16** in the manufacture of a medicament for treating cancer.

33. The use of claim **32**, wherein the antibody or immunoconjugate blocks binding of a Wnt ligand to a non-Wnt3a binding site of LRP5.