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(54) Title: PD-1 ANTIBODIES

(57) Abstract: The present invention provides antibodies that bind human programmed cell death 1 (PD-1). These antibodies may be useful for treating cancer alone and in combination with chemotherapy and other cancer therapeutics.



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## PD-1 Antibodies

The present invention relates to the field of medicine. More particularly, the present invention relates to antibodies that bind human programmed cell death 1 (PD-1),  
5 and may be useful for treating cancer alone and in combination with chemotherapy and other cancer therapeutics.

Tumor cells escape detection and elimination by the immune system through multiple mechanisms. Immune checkpoint pathways are used in self-tolerance maintenance and activated T cell control, but cancer cells can use the pathways to prevent  
10 destruction. The PD-1 / human programmed cell death 1 ligand 1 (PD-L1) pathway is one such immune checkpoint. Human PD-1 is found on T cells, and the binding of PD-L1 and human programmed cell death 1 ligand 2 (PD-L2) to PD-1 inhibits T cell proliferation and cytokine production. Tumor cell production of PD-L1 and PD-L2 can therefore allow escape from T cell surveillance.

15 A fully human IgG4 (S228P) antibody against human PD-1, nivolumab, has been shown to inhibit the binding of PD-1 to PD-L1 and PD-L2, and has been tested in various clinical trials. (Wang et al., Cancer Immunol Res (2014) 2(9):846). A humanized IgG4 (S228P) antibody against PD-1, pembrolizumab (formerly lambrolizumab), has been shown to inhibit the binding of PD-1 to PD-L1 and PD-L2, and has been tested in various  
20 clinical trials. (WO2008156712 and Hamid et al., N Engl J Med (2013) 369:2).

There remains a need to provide alternative antibodies that bind and neutralize human PD-1 interaction with PD-L1 and PD-L2. In particular, there remains a need to provide antibodies that bind human PD-1 with high affinity but with different features than clinically approved PD-1 antibodies, such as binding to mouse and human PD-1.  
25 Further, there remains a need to provide antibodies that bind human PD-1 with affinity similar to clinically approved PD-1 but bind human PD-1 differently. Also, there remains a need to provide antibodies that more effectively block the human PD-1 interaction with PD-L1 and PD-L2 than certain prior art antibodies. Better blocking can translate into greater *in vivo* activity or lower required dosing amounts.

30 Certain antibodies of the present invention block the human PD-1 to PD-L1 and PD-L2 interactions in CHO cells more effectively than nivolumab and pembrolizumab.

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Furthermore, certain antibodies of the present invention bind murine PD-1 on CHO cells whereas nivolumab and pembrolizumab binding to murine PD-1 is not detected.

Accordingly, in some embodiments the present invention provides an antibody that binds human PD-1 (SEQ ID NO: 1), comprising a light chain (LC) and a heavy chain (HC), wherein the light chain comprises light chain complementarity determining regions LCDR1, LCDR2, and LCDR3 consisting of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein the heavy chain comprises heavy chain complementarity determining regions HCDR1, HCDR2, and HCDR3, wherein HCDR1 consists of the amino acid sequence:

- a. KASGYTFTSYMH (SEQ ID NO: 2),
- b. KASGYTFEGYMH (SEQ ID NO: 3),
- c. KASGYTFTAQYMH (SEQ ID NO: 4),
- d. KASGYTFEKYMH (SEQ ID NO: 5),
- e. KASGYTFTSNYMH (SEQ ID NO: 6), or
- f. KASGYTFSAYMH (SEQ ID NO: 7);

wherein HCDR2 consists of the amino acid sequence:

- a. IINPSGGSTSYAQKFQG (SEQ ID NO: 8),
- b. IINPEGGETSYAQKFQG (SEQ ID NO: 9),
- c. IINPSGGETGYAQKFQG (SEQ ID NO: 10),
- d. IINPSEGSTGYAQKFQG (SEQ ID NO: 11), or
- e. IINPDGGSTGYAQKFQG (SEQ ID NO: 12);

and wherein HCDR3 consists of the amino acid sequence AKEGVADGYGLVDV (SEQ ID NO: 13).

In some embodiments, the present invention provides an antibody that binds human PD-1 (SEQ ID NO: 1), wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFTSYMH (SEQ ID NO: 2), IINPSGGSTSYAQKFQG (SEQ ID NO: 8), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.

In some embodiments, the present invention provides an antibody that binds human PD-1 (SEQ ID NO: 1), wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFEGYYMH (SEQ ID NO: 3), IINPEGGETSYAQKFQG (SEQ ID NO: 9), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.

In some embodiments, the present invention provides an antibody that binds human PD-1 (SEQ ID NO: 1), wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFTAQYMH (SEQ ID NO: 4), IINPSGGETGYAQKFQG (SEQ ID NO: 10), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.

In some embodiments, the present invention provides an antibody that binds human PD-1 (SEQ ID NO: 1), wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFEKYYMH (SEQ ID NO: 5), IINPDGGSTGYAQKFQG (SEQ ID NO: 12), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.

In some embodiments, the present invention provides an antibody that binds human PD-1 (SEQ ID NO: 1), wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFTSNYMH (SEQ ID NO: 6), IINPSEGSTGYAQKFQG (SEQ ID NO: 11), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.

In some embodiments, the present invention provides an antibody that binds human PD-1 (SEQ ID NO: 1), wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO:

15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFSAYYMH (SEQ ID NO: 7), IINPDGGSTGYAQKFQG (SEQ ID NO: 12), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.

5 In some embodiments, the present invention provides an antibody, comprising a light chain (LC) and a heavy chain (HC), wherein the light chain comprises a light chain variable region (LCVR) and the heavy chain comprises a heavy chain variable region (HCVR), wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 17, SEQ ID NO: 18, SEQ  
10 ID NO: 19, SEQ ID NO: 20, SEQ ID NO: 21, or SEQ ID NO: 22.

In a further embodiment, the present invention provides an antibody, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 17.

In a further embodiment, the present invention provides an antibody, wherein the  
15 LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 18.

In a further embodiment, the present invention provides an antibody, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 19.

20 In a further embodiment, the present invention provides an antibody, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 20.

In a further embodiment, the present invention provides an antibody, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the  
25 amino acid sequence given in SEQ ID NO: 21.

In a further embodiment, the present invention provides an antibody, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 22.

In some embodiments, the present invention provides an antibody, wherein the LC  
30 has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid

sequence given in SEQ ID NO: 24, SEQ ID NO: 25, SEQ ID NO: 26, SEQ ID NO: 27, SEQ ID NO: 28, or SEQ ID NO: 29.

5 In a further embodiment, the present invention provides an antibody, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 24.

In a further embodiment, the present invention provides an antibody, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 25.

10 In a further embodiment, the present invention provides an antibody, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 26.

In a further embodiment, the present invention provides an antibody, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 27.

15 In a further embodiment, the present invention provides an antibody, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 28.

In a further embodiment, the present invention provides an antibody, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 29.

20 In some embodiments, the present invention provides an antibody, comprising two light chains and two heavy chains, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 24, SEQ ID NO: 25, SEQ ID NO: 26, SEQ ID NO: 27, SEQ ID NO: 28, or  
25 SEQ ID NO: 29.

In a further embodiment, the present invention provides an antibody, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 24.

30 In a further embodiment, the present invention provides an antibody, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 25.

In a further embodiment, the present invention provides an antibody, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 26.

In a further embodiment, the present invention provides an antibody, wherein each  
5 light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 27.

In a further embodiment, the present invention provides an antibody, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 28.

10 In a further embodiment, the present invention provides an antibody, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 29.

In an embodiment, the present invention provides an antibody, wherein one of the heavy chains forms an inter-chain disulfide bond with one of the light chains, and the  
15 other heavy chain forms an inter-chain disulfide bond with the other light chain, and one of the heavy chains forms two inter-chain disulfide bonds with the other heavy chain.

In an embodiment, the present invention provides an antibody, wherein the antibody is glycosylated.

In some embodiments, the present invention provides an antibody that binds both  
20 human PD-1 and mouse PD-1. In a further embodiment, the present invention provides an antibody that is an antagonistic antibody and that binds both human PD-1 and mouse PD-1. In a further embodiment, the present invention provides an antibody that binds both human PD-1 and mouse PD-1 with a  $K_d$  for each less than 400 pM. In a further embodiment, the present invention provides an antibody that binds both human PD-1 and  
25 mouse PD-1 with a  $K_d$  for each less than 200 pM.

In an embodiment, the present invention provides a pharmaceutical composition, comprising an antibody of the present invention, and an acceptable carrier, diluent, or excipient.

30 In an embodiment, the present invention provides a method of treating cancer, comprising administering to a patient in need thereof, an effective amount of an antibody of the present invention. In a further embodiment, the present invention provides a

method of treating cancer, comprising administering to a patient in need thereof, an effective amount of an antibody of the present invention, wherein the cancer is melanoma, lung cancer, head and neck cancer, colorectal cancer, pancreatic cancer, gastric cancer, kidney cancer, bladder cancer, prostate cancer, breast cancer, ovarian cancer, or  
5 hepatocellular carcinoma.

In a further embodiment, these methods comprise the administration of an effective amount of the antibody of the present invention in simultaneous, separate, or sequential combination with one or more anti-tumor agents. Non-limiting examples of anti-tumor agents include ramucirumab, necitumumab, olaratumab, galunisertib,  
10 abemaciclib, cisplatin, carboplatin, dacarbazine, liposomal doxorubicin, docetaxel, cyclophosphamide and doxorubicin, navelbine, eribulin, paclitaxel, paclitaxel protein-bound particles for injectable suspension, ixabepilone, capecitabine, FOLFOX (leucovorin, fluorouracil, and oxaliplatin), FOLFIRI (leucovorin, fluorouracil, and irinotecan), and cetuximab.

15 In a further embodiment, these methods comprise the administration of an effective amount of the compound of the present invention in simultaneous, separate, or sequential combination with one or more immuno-oncology agents. Non-limiting examples of immuno-oncology agents include nivolumab, ipilimumab, pidilizumab, pembrolizumab, tremelimumab, urelumab, lirilumab, atezolizumab, and durvalumab.

20 In an embodiment, the present invention provides an antibody of the present invention, for use in therapy. In an embodiment, the present invention provides an antibody of the present invention, for use in the treatment of cancer. In a further embodiment, the present invention provides an antibody of the present invention, for use in the treatment of cancer, wherein the cancer is melanoma, lung cancer, head and neck  
25 cancer, colorectal cancer, pancreatic cancer, gastric cancer, kidney cancer, bladder cancer, prostate cancer, breast cancer, ovarian cancer, or hepatocellular carcinoma. In a further embodiment, the present invention provides the antibody of the present invention for use in simultaneous, separate, or sequential combination with one or more anti-tumor agents. In a further embodiment, the present invention provides the antibody of the present  
30 invention for use in simultaneous, separate, or sequential combination with one or more anti-tumor agents selected from the group consisting of ramucirumab, necitumumab,



olaratumab, galunisertib, abemaciclib, cisplatin, carboplatin, dacarbazine, liposomal doxorubicin, docetaxel, cyclophosphamide and doxorubicin, navelbine, eribulin, paclitaxel, paclitaxel protein-bound particles for injectable suspension, ixabepilone, capecitabine, FOLFOX (leucovorin, fluorouracil, and oxaliplatin), FOLFIRI (leucovorin, 5 fluorouracil, and irinotecan), and cetuximab, in the treatment of cancer.

In a further embodiment, the present invention provides the antibody of the present invention for use in simultaneous, separate, or sequential combination with one or more immuno-oncology agents. In a further embodiment, the present invention provides the antibody of the present invention for use in simultaneous, separate, or sequential 10 combination with one or more immuno-oncology agents selected from the group consisting of nivolumab, ipilimumab, pidilizumab, pembrolizumab, tremelimumab, urelumab, lirilumab, atezolizumab, and durvalumab, in the treatment of cancer.

In a further embodiment, the present invention provides the use of an antibody of the present invention for the manufacture of a medicament for the treatment of cancer. In 15 a further embodiment, the present invention provides the use of an antibody of the present invention for the manufacture of a medicament for the treatment of cancer, wherein the cancer is melanoma, lung cancer, head and neck cancer, colorectal cancer, pancreatic cancer, gastric cancer, kidney cancer, bladder cancer, prostate cancer, breast cancer, ovarian cancer, or hepatocellular carcinoma.

20 In a further embodiment, the present invention provides the use of an antibody of the present invention in the manufacture of a medicament for the treatment of cancer wherein said medicament is to be administered simultaneously, separately, or sequentially with one or more anti-tumor agents. In a further embodiment, the present invention provides the use of an antibody of the present invention in the manufacture of a 25 medicament for the treatment of cancer wherein said medicament is to be administered simultaneously, separately, or sequentially with one or more anti-tumor agents selected from the group consisting of ramucirumab, necitumumab, olaratumab, galunisertib, abemaciclib, cisplatin, carboplatin, dacarbazine, liposomal doxorubicin, docetaxel, cyclophosphamide and doxorubicin, navelbine, eribulin, paclitaxel, paclitaxel protein- 30 bound particles for injectable suspension, ixabepilone, capecitabine, FOLFOX

(leucovorin, fluorouracil, and oxaliplatin), FOLFIRI (leucovorin, fluorouracil, and irinotecan), and cetuximab.

An antibody of the present invention is an engineered, non-naturally occurring polypeptide complex. A DNA molecule of the present invention is a non-naturally occurring DNA molecule that comprises a polynucleotide sequence encoding a polypeptide having the amino acid sequence of one of the polypeptides in an antibody of the present invention.

An antibody of the present invention is designed to have engineered CDRs and have some portions of the antibody (all or parts of the frameworks, hinge regions, and constant regions) to be of human origin that are identical with or substantially identical (substantially human) with frameworks and constant regions derived from human genomic sequences. Fully human frameworks, hinge regions, and constant regions are those human germline sequences as well as sequences with naturally-occurring somatic mutations and those with engineered mutations. An antibody of the present invention may comprise framework, hinge, or constant regions derived from a fully human framework, hinge, or constant region containing one or more amino acid substitutions, deletions, or additions therein. Further, an antibody of the present invention is preferably substantially non-immunogenic in humans.

The antibody of the present invention is an IgG type antibody and has “heavy” chains and “light” chains that are cross-linked via intra- and inter-chain disulfide bonds. Each heavy chain is comprised of an N-terminal HCVR and a heavy chain constant region (“HCCR”). Each light chain is comprised of a LCVR and a light chain constant region (“LCCR”). When expressed in certain biological systems, antibodies having native human Fc sequences are glycosylated in the Fc region. Typically, glycosylation occurs in the Fc region of the antibody at a highly conserved N-glycosylation site. N-glycans typically attach to asparagine. Antibodies may be glycosylated at other positions as well.

Optionally, the antibody of the present invention contains an Fc portion which is derived from human IgG<sub>4</sub> Fc region because of a reduced ability to engage Fc receptor-mediated inflammatory mechanisms or to activate complement resulting in reduced effector function.

Certain antibodies of the present invention contain an IgG<sub>4</sub>-Fc portion that has a serine to proline mutation at position 228. The S228P mutation is a hinge mutation that prevents half-antibody formation (phenomenon of dynamic exchange of half-molecules in IgG<sub>4</sub> antibodies).

5           The HCVR and LCVR regions can be further subdivided into regions of hyper-variability, termed complementarity determining regions (“CDRs”), interspersed with regions that are more conserved, termed framework regions (“FR”). Each HCVR and LCVR is composed of three CDRs and four FRs, arranged from amino-terminus to carboxy-terminus in the following order: FR1, CDR1, FR2, CDR2, FR3, CDR3, FR4.

10       Herein, the three CDRs of the heavy chain are referred to as “HCDR1, HCDR2, and HCDR3” and the three CDRs of the light chain are referred to as “LCDR1, LCDR2 and LCDR3”. The CDRs contain most of the residues which form specific interactions with the antigen. There are currently three systems of CDR assignments for antibodies that are used for sequence delineation. The Kabat CDR definition (Kabat *et al.*, “Sequences of

15       Proteins of Immunological Interest,” National Institutes of Health, Bethesda, Md. (1991)) is based upon antibody sequence variability. The Chothia CDR definition (Chothia *et al.*, “Canonical structures for the hypervariable regions of immunoglobulins”, Journal of Molecular Biology, 196, 901-917 (1987); Al-Lazikani *et al.*, “Standard conformations for the canonical structures of immunoglobulins”, Journal of Molecular Biology, 273, 927-

20       948 (1997)) is based on three-dimensional structures of antibodies and topologies of the CDR loops. The Chothia CDR definitions are identical to the Kabat CDR definitions with the exception of HCDR1 and HCDR2. The North CDR definition (North *et al.*, “A New Clustering of Antibody CDR Loop Conformations”, Journal of Molecular Biology, 406, 228-256 (2011)) is based on affinity propagation clustering with a large number of

25       crystal structures.

          An isolated DNA encoding a HCVR region can be converted to a full-length heavy chain gene by operably linking the HCVR-encoding DNA to another DNA molecule encoding heavy chain constant regions. The sequences of human, as well as other mammalian, heavy chain constant region genes are known in the art. DNA

30       fragments encompassing these regions can be obtained *e.g.*, by standard PCR amplification.

An isolated DNA encoding a LCVR region may be converted to a full-length light chain gene by operably linking the LCVR-encoding DNA to another DNA molecule encoding a light chain constant region. The sequences of human, as well as other mammalian, light chain constant region genes are known in the art. DNA fragments encompassing these regions can be obtained by standard PCR amplification. The light chain constant region can be a kappa or lambda constant region.

The polynucleotides of the present invention will be expressed in a host cell after the sequences have been operably linked to an expression control sequence. The expression vectors are typically replicable in the host organisms either as episomes or as an integral part of the host chromosomal DNA. Commonly, expression vectors will contain selection markers, e.g., tetracycline, neomycin, and dihydrofolate reductase, to permit detection of those cells transformed with the desired DNA sequences.

The antibody of the present invention may readily be produced in mammalian cells such as CHO, NS0, HEK293 or COS cells. The host cells are cultured using techniques well known in the art.

The vectors containing the polynucleotide sequences of interest (e.g., the polynucleotides encoding the polypeptides of the antibody and expression control sequences) can be transferred into the host cell by well-known methods, which vary depending on the type of cellular host.

Various methods of protein purification may be employed and such methods are known in the art and described, for example, in Deutscher, *Methods in Enzymology* 182: 83-89 (1990) and Scopes, *Protein Purification: Principles and Practice*, 3rd Edition, Springer, NY (1994).

In another embodiment of the present invention, the antibody, or the nucleic acids encoding the same, is provided in isolated form. As used herein, the term "isolated" refers to a protein, peptide, or nucleic acid which is free or substantially free from any other macromolecular species found in a cellular environment. "Substantially free" as used herein means the protein, peptide, or nucleic acid of interest comprises more than 80% (on a molar basis) of the macromolecular species present, preferably more than 90%, and more preferably more than 95%.

The antibody of the present invention, or pharmaceutical compositions comprising the same, may be administered by parenteral routes (e.g., subcutaneous and intravenous). An antibody of the present invention may be administered to a patient alone with pharmaceutically acceptable carriers, diluents, or excipients in single or multiple doses.

5 Pharmaceutical compositions of the present invention can be prepared by methods well known in the art (e.g., *Remington: The Science and Practice of Pharmacy*, 19<sup>th</sup> ed. (1995), A. Gennaro et al., Mack Publishing Co.) and comprise an antibody, as disclosed herein, and one or more pharmaceutically acceptable carriers, diluents, or excipients.

The term "treating" (or "treat" or "treatment") refers to slowing, interrupting,  
10 arresting, alleviating, stopping, reducing, or reversing the progression or severity of an existing symptom, disorder, condition, or disease.

"Binds" as used herein in reference to the affinity of an antibody for human PD-1 is intended to mean, unless indicated otherwise, a  $K_D$  of less than about  $1 \times 10^{-6}$  M, preferably, less than about  $1 \times 10^{-9}$  M as determined by common methods known in the  
15 art, including by use of MSD essentially as described herein.

For the purposes of the present disclosure, the term "high affinity" refers to a  $K_D$  of less than about 150 pM for human PD-1 as determined by MSD. The  $K_D$  values are established by binding kinetics as described in "Binding kinetics and affinity" in the Assays section.

20 "Effective amount" means the amount of an antibody of the present invention or pharmaceutical composition comprising an antibody of the present invention that will elicit the biological or medical response of or desired therapeutic effect on a tissue, system, animal, mammal or human that is being sought by the researcher, medical doctor, or other clinician. An effective amount of the antibody may vary according to factors  
25 such as the disease state, age, sex, and weight of the individual, and the ability of the antibody to elicit a desired response in the individual. An effective amount is also one in which any toxic or detrimental effect of the antibody is outweighed by the therapeutically beneficial effects.

This invention is further illustrated by the following non-limiting example.

**Example 1: Antibody expression and purification**

The polypeptides of the variable regions of the heavy chain and light chain, the complete heavy chain and light chain amino acid sequences of Antibody A - Antibody F, and the nucleotide sequences encoding the same, are listed below in the section entitled  
5 “Amino Acid and Nucleotide Sequences.” In addition, the SEQ ID NOs for the light chain, heavy chain, light chain variable region, and heavy chain variable region of Antibody A - Antibody F are shown in Table 1.

The antibodies of the present invention, including, but not limited to, Antibody A - Antibody F can be made and purified essentially as follows. An appropriate host cell,  
10 such as HEK 293 or CHO, can be either transiently or stably transfected with an expression system for secreting antibodies using an optimal predetermined HC:LC vector ratio or a single vector system encoding both HC and LC. Clarified media, into which the antibody has been secreted, may be purified using any of many commonly-used techniques. For example, the medium may be conveniently applied to a MabSelect  
15 column (GE Healthcare), or KappaSelect column (GE Healthcare) for Fab fragment, that has been equilibrated with a compatible buffer, such as phosphate buffered saline (pH 7.4). The column may be washed to remove nonspecific binding components. The bound antibody may be eluted, for example, by pH gradient (such as 20 mM Tris buffer pH 7 to 10 mM sodium citrate buffer pH 3.0, or phosphate buffered saline pH 7.4 to 100  
20 mM glycine buffer pH 3.0). Antibody fractions may be detected, such as by SDS-PAGE, and then may be pooled. Further purification is optional, depending on the intended use. The antibody may be concentrated and/or sterile filtered using common techniques. Soluble aggregate and multimers may be effectively removed by common techniques, including size exclusion, hydrophobic interaction, ion exchange, multimodal,  
25 or hydroxyapatite chromatography. The purity of the antibody after these chromatography steps is greater than 95%. The product may be immediately frozen at -70°C or may be lyophilized.

**Table 1: SEQ ID NOs**

	Antibody A	Antibody B	Antibody C	Antibody D	Antibody E	Antibody F
HCVR	17	18	19	20	21	22
LCVR	23	23	23	23	23	23
Heavy chain	24	25	26	27	28	29
Light chain	30	30	30	30	30	30

**Assays****5 Binding kinetics and affinity**

The kinetics and equilibrium dissociation constant ( $K_D$ ) for human PD-1 is determined for antibodies of the present invention using MSD, and bio-layer interferometry (ForteBio) assay methods.

As used herein, nivolumab is a human IgG4 PD-1 antibody transiently expressed in 293 HEK cells that utilizes the heavy chain and light chain sequences from Proposed INN: List 107 (CAS #946414-94-4). As used herein, pembrolizumab is a human IgG4 PD-1 antibody transiently expressed in 293 HEK cells that utilizes the heavy chain and light chain sequences from Proposed INN: List 72.

**15 MSD assay**

Equilibrium affinity measurements are performed as previously described (Estep, P., et al., MAbs, 2013. **5**(2): p. 270-8). Solution equilibrium titrations (SET) are performed in PBS + 0.1% IgG-Free BSA (PBSF) where antigen (b-PD-1 monomer) is held constant at 10-100 pM and is incubated with 3-to 5-fold serial dilutions of Fab or mAbs starting at 5-100 nM (experimental condition is sample dependent). Antibodies diluted at 20 nM in PBS are coated onto standard bind MSD-ECL plates overnight at 4°C or at room temperature for 30 min. Plates are blocked with BSA for 30 min whilst shaking at 700 rpm. Plates are then washed 3x with wash buffer (PBSF + 0.05% Tween 20). SET samples are applied and incubated on the plates for 150s with shaking at 700

rpm followed by one wash. Antigen captured on a plate is detected with 250 ng/mL sulfotag-labeled streptavidin in PBSF by incubation on the plate for 3 min. The plates are washed three times with wash buffer and are then read on the MSD Sector Imager 2400 instrument using 1x Read Buffer T with surfactant. The percent free antigen is plotted as a function of titrated antibody in Prism and fit to a quadratic equation to extract the  $K_D$ . To improve throughput, liquid handling robots are used throughout MSD-SET experiments, including for SET sample preparation.

In experiments performed essentially as described in this assay, Antibody C and Antibody E, in an IgG1 format and expressed in yeast, bind human PD-1 with a  $K_D$  of 120 pM and 91 pM, respectively. Pembrolizumab and nivolumab bind PD-1 with a  $K_D$  of 130 pM and 640 pM respectively. Avidity measurements for Antibody C and Antibody E result in a  $K_D$  of approximately 9 pM and 22 pM, respectively, to human PD-1. Pembrolizumab and nivolumab bind human PD-1 with a  $K_D$  of approximately 3 pM and 5 pM respectively. Antibody C and Antibody E, in an IgG1 format and expressed in yeast, bind murine PD-1 with a  $K_D$  of 1900 pM and 1100 pM, respectively. Avidity measurements for Antibody C and Antibody E result in a  $K_D$  of approximately 130 pM and 330 pM, respectively, to murine PD-1.

**Table 1: Binding by MSD of antibodies of the invention in IgG1 format (P.F.: poor fit; N.D.: none detected)**

Name	Monovalent $K_D$ (M) against human PD-1	Avid $K_D$ (M) against human PD-1	Monovalent $K_D$ (M) against murine PD-1	Avid $K_D$ (M) against murine PD-1
Antibody A	4.90E-09	4.90E-11	9.60E-09	P.F.
Antibody B	1.30E-10	1.40E-11	2.70E-09	3.80E-10
Antibody C	1.20E-10	9.30E-12	1.90E-09	1.30E-10
Antibody D	3.00E-10	3.50E-11	2.30E-09	P.F.
Antibody E	9.10E-11	2.20E-11	1.10E-09	3.30E-10
Antibody F	1.70E-10	1.80E-11	1.20E-09	4.40E-10
Pembrolizumab	1.30E-10	3.00E-12	N.D.	N.D.
Nivolumab	6.40E-10	5.10E-12	N.D.	N.D.



Bio-layer interferometry

ForteBio affinity measurements were performed generally as previously described (Estep, P., et al., *High throughput solution-based measurement of antibody-antigen affinity and epitope binning*, MAbs, 2013. **5**(2): p. 270-8.). Briefly, ForteBio affinity measurements were performed by loading IgGs online onto AHQ sensors. Sensors were equilibrated off-line in assay buffer for 30 min and then monitored on-line for 60 seconds for baseline establishment. Sensors with loaded IgGs were exposed to 100 nM antigen for 5 min, afterwards they were transferred to assay buffer for 5 min for off-rate measurement. Kinetics was analyzed using the 1:1 binding model.

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**Table 2: Binding by Bio-light interferometry of antibodies of the invention in IgG1 format (N.D.: none detected)**

	Monovalent $K_D$ (M) Fab in solution, human PD-1_Fc on sensor tip	Monovalent $K_D$ (M) human PD-1_HIS in solution, IgG on sensor tip	Monovalent $K_D$ (M) Fab in solution, cyno PD-1_Fc on sensor tip	Monovalent $K_D$ (M) Fab in solution, murine PD-1_Fc on sensor tip
Antibody A	1.60E-08	4.30E-09	1.40E-08	3.30E-08
Antibody B	2.70E-09	1.40E-09	3.70E-09	1.70E-08
Antibody C	2.00E-09	1.00E-09	2.40E-09	1.00E-08
Antibody D	3.40E-09	1.90E-09	4.40E-09	7.80E-09
Antibody E	2.70E-09	1.20E-09	3.30E-09	7.60E-09
Antibody F	3.00E-09	1.60E-09	3.60E-09	6.10E-09
Pembrolizumab	2.00E-09	2.00E-09	4.70E-10	N.D.
Nivolumab	1.70E-09	4.10E-09	1.20E-09	N.D.

In experiments performed essentially as described in this assay, Antibody C and Antibody E, in an IgG1 format and expressed in yeast, bind human PD-1\_Fc with a  $K_D$  similar to nivolumab and pembrolizumab when PD-1\_Fc was on the sensor tip. When the antibody was on the sensor tip, Antibody C and Antibody E, in an IgG1 format and expressed in yeast, bind human PD-1\_Fc with a  $K_D$  similarly to nivolumab and pembrolizumab. Antibody C and Antibody E, in an IgG1 format and expressed in yeast,

15

bind cyno PD-1\_Fc with a similar  $K_D$  to nivolumab and pembrolizumab. Antibody C and Antibody E, in an IgG1 format and expressed in yeast, bind murine PD-1\_Fc whereas no binding was detected with nivolumab and pembrolizumab.

## 5 Binding to human and murine PD-1 on CHO cells

The binding of an antibody of the present invention to human PD-1 may be measured in a flow cytometry assay.

CHO cells ( $0.2 \times 10^6$ ) are incubated with the experimental antibody from 100 nM titrated 14x by a factor of 2 to the lowest concentration of 6.1 pM for 30 min in PBS 1% BSA on ice. Cells are then washed 3x, and are incubated with a secondary antibody (PE-labelled, at final concentration of 5  $\mu\text{g/ml}$ ) in PBS 1% BSA for 30 min on ice (protected from light). Cells are washed 3x and analyzed via flow cytometry. Flow cytometry is performed on an Accuri C6 system (BD Biosciences) and MFIs are calculated on the C6 software. EC50s are calculated on Graphpad software.

In experiments performed essentially as described in this assay, Antibody A, Antibody C, and Antibody E, in an IgG1 format and expressed in yeast bind human PD-1 in a dose-dependent manner, with an EC50 value (n=1) of 10.22 nM, 4.115 nM and 4.587 nM, respectively, and pembrolizumab binds human PD-1 with an EC50 value (n=1) of 0.6921 nM and nivolumab with an EC50 value (n=1) of 0.8057 nM.

Antibody A, B, C, D, E, and F, in an IgG1 format and expressed in yeast bind to murine PD-1 in a dose dependent manner with an EC50 of 11.01 nM, 6.135 nM, 2.884 nM, 9.259 nM, and 5.044 nM, and 5.855 nM, respectively, whereas no binding was detected for nivolumab or pembrolizumab.

## 25 Blocking of human PD-1 to PD-L1 and PD-L2 in CHO cells.

The ability of an antibody of the present invention to block binding of human PD-1 to PD-L1 and PD-L2 may be measured by flow cytometry.

CHO cells  $0.2 \times 10^6$  are incubated with the experimental antibody (100 nM) for 30 min in PBS 1% BSA on ice. Cells are then washed 3X, and are incubated with PD-L2 linked with NHS-Fluorescein (Promega) in PBS 1% BSA for 30 min on ice (protected from light). Cells are washed 3x and analyzed via flow cytometry. Flow cytometry is

performed on an Accuri C6 system (BD Biosciences) and MFIs are calculated on the C6 software. EC50s are calculated on Graphpad software.

In experiments performed essentially as described in this assay, Antibody C and Antibody E (IgG1 format expressed in yeast) blocked human PD-L2-FITC binding, resulting in an MFI of 30,123.4 and 38,682.1, respectively, as compared to control IgG which resulted in an MFI of 182,959.1. Pembrolizumab and nivolumab resulted in MFI's of 46,245.9 and 54,509.8 respectively.

**Table 3: Blocking of human PD-1 on CHO cells**

Test Sample	MFI (PD-L2-FITC)
Cells only	33,449.7
No IgG	199,716.0
IgG Control	182,959.1
Nivolumab	54,509.8
Pembrolizumab	46,245.9
Antibody A	90,866.5
Antibody C	30,123.4
Antibody E	38,682.1

### Mixed Lymphocyte Reaction

The blocking of PD-1 signals by antibodies of the present invention may be evaluated by measuring the release of inhibitory signals during T cell activation.

$2 \times 10^6$  PBMC are plated per well in a 6 well tissue culture plate or T25 tissue culture flask in complete T cell media. Cells are incubated for 2-3 hours, to allow for adherence of monocytes. If adherence is insufficient, serum free media is used. Unattached cells are removed by gently swirling the flask with fresh media 3X.

Immature myeloid DCs are generated by culturing monocytes ( $1 \times 10^6$  cells/ml) from PBMC in X-VIVO 15 media containing 1% AB serum, 10 mM HEPES, 50  $\mu$ M  $\beta$ -Me, IL-4 (1000 U/ml) and GM-CSF (1000 U/ml), or 25-50 ng/ml of each. After 2 days fresh medium supplemented with IL-4 and GM-CSF is added. On Day 5, cells are either frozen or maturation is induced by adding a stimulation cocktail containing rTNFa (1000U/ml), IL-1b (5 ng/ml), IL-6 (10ng/ml) and 1  $\mu$ M PGE<sub>2</sub> for 2 days at a cell density of  $3 \times 10^5$  cells/ml.

T cell Isolation is performed as per manufacturer's instructions in the Untouched CD4<sup>+</sup> T cell isolation kit (Invitrogen). A magnet fitted with a 1.5 ml tube rack is used to remove unwanted magnetic beads (QIAGEN). 100,000-200,000 isolated T cells are mixed with 10,000-20,000 allogeneic moDCs in a total volume of 200  $\mu$ l in 96-round bottom tissue culture plates for 4-5 days at 37°C. T cells are stimulated using anti-CD3/CD28 DynaBeads at a ratio of 3:1 (cells:beads) as a positive control; beads are prepared as per the manufacturer's instructions. Test antibodies are added at the beginning of the MLR and incubated throughout the culture period. Detection of IL-2 and IFN- $\gamma$  is carried out as per manufacturer's instructions (eBioscience). OD measurements are determined on a Multiskan FC system (Thermo).

In experiments performed essentially as described in this assay, Antibody C and Antibody E increase IL-2 and IFN $\gamma$  secretion by activated T cells in the mixed lymphocyte reaction with equivalent potency as compared with nivolumab and pembrolizumab.

**Table 4: IL-2 secretion fold change vs. IgG control**

IgG	Concentrations of IgG				
	100 nM	10 nM	1 nM	0.1 nM	0.01 nM
Pembrolizumab	2.03	2.49	2.04	1.47	1.06
Nivolumab	2.37	2.44	1.72	1.26	1.09
Antibody A	2.28	1.87	1.23	1.15	1.37
Antibody C	2.62	2.29	1.83	1.15	1.01
Antibody E	3.20	2.61	1.95	1.21	1.13

	Cytokine in supernatant	Concentrations of IgG				
		100 nM	10 nM	1 nM	0.1 nM	0.01 nM
IgG control	IL-2 (pg/ml)	619.16	521.57	500.33	508.82	515.02
	Stdev. (pg/ml)	18.53	4.13	25.45	18.92	18.33
Pembrolizumab	IL-2 (pg/ml)	1257.59	1299.83	1021.63	749.33	545.48
	Stdev. (pg/ml)	27.37	13.42	37.56	33.53	11.21
Nivolumab	IL-2 (pg/ml)	1469.85	1274.69	858.20	641.13	562.30

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	Stdev. (pg/ml)	55.01	68.79	44.24	23.40	33.04
Antibody A	IL-2 (pg/ml)	1410.46	975.38	617.86	585.50	707.77
	Stdev. (pg/ml)	41.38	27.88	49.01	8.39	27.93
Antibody C	IL-2 (pg/ml)	1622.02	1194.17	913.62	587.10	518.69
	Stdev. (pg/ml)	49.88	46.17	30.18	42.17	12.79
Antibody E	IL-2 (pg/ml)	1983.95	1361.71	977.51	614.19	580.24
	Stdev. (pg/ml)	119.32	57.80	20.95	15.81	12.83

**Table 5: IFN $\gamma$  secretion fold change vs. IgG control**

IgG	Concentrations of IgG				
	100 nM	10 nM	1 nM	0.1 nM	0.01 nM
Pembrolizumab	1.78	1.77	1.76	1.99	1.03
Nivolumab	1.97	1.88	1.58	1.53	0.84
Antibody A	1.52	1.37	1.11	1.49	1.49
Antibody C	1.84	1.79	1.82	1.50	0.97
Antibody E	1.76	2.01	1.86	1.26	0.96

	Cytokine in supernatant	Concentrations of IgG				
		100 nM	10 nM	1 nM	0.1 nM	0.01 nM
IgG control	IFN $\gamma$ (pg/ml)	14936.03	13497.03	11603.92	11007.43	14881.91
	Stdev. (pg/ml)	331.65	912.08	815.69	1400.66	453.67
Pembrolizumab	IFN $\gamma$ (pg/ml)	26598.52	23903.29	20390.76	21894.58	15326.75
	Stdev. (pg/ml)	1143.84	409.35	1274.53	1038.25	280.19
Nivolumab	IFN $\gamma$ (pg/ml)	29482.91	25333.89	18296.62	16820.33	12487.21
	Stdev. (pg/ml)	3935.40	3201.10	203.28	725.93	613.05
Antibody A	IFN $\gamma$ (pg/ml)	22631.06	18442.75	12915.14	16368.19	22121.86
	Stdev.	712.64	3029.49	1683.58	580.07	160.46

	(pg/ml)					
Antibody C	IFN $\gamma$ (pg/ml)	27443.11	24094.79	21136.75	16460.20	14382.09
	Stdev. (pg/ml)	1036.81	888.47	936.17	914.50	929.68
Antibody E	IFN $\gamma$ (pg/ml)	26262.69	27124.92	21575.50	13825.35	14331.43
	Stdev. (pg/ml)	898.53	1884.76	508.65	513.84	1708.45

### Tumor model

The antibodies of the present invention can be measured for *in vivo*

- 5 immunomodulatory activity with the MC38 *in vivo* tumor mode. C57Bl/6 mice are inoculated subcutaneously with MC38 murine colon cancer cells  $2 \times 10^6$  per mouse to establish the tumor-bearing model. Mice are selected 10 days after tumor inoculation with tumor volumes between  $34.81 \text{ mm}^3 \sim 148.24 \text{ mm}^3$  and then divided into five groups. These groups include a control group, RMP1-14 (Bio X Cell), Compound C-2.5,
- 10 Compound C-5, and Compound C-10 (n=10). RMP1-14 group is given a dose of 10 mg/kg. Compound C-2.5, Compound C-5, and Compound C-10 groups are given 2.5 mg/kg, 5 mg/kg and 10 mg/kg of 11430 respectively (Compound C S228P IgG4 produced in HEK293 cells).

- The control group receives an equal volume of saline, and all groups are dosed via
- 15 intraperitoneal injection, dose frequency of 2 times weekly for 4 weeks. Weekly monitoring of animal body weight, and tumor volume (using formula  $V=L \times W^2/2$ ) is conducted. After the experiment, tumors are photographed and weighed to calculate tumor weight and to calculate relative tumor inhibition.

- In experiments performed essentially as described in this assay, the results show
- 20 that administration of RMP1-14 and Compound C has no effect on animal weight. RMP1-14 and Compound C show significant anti-tumor effect after one week of administration. The three Compound C dose groups have higher anti-tumor (tumor volume reducing) effects compared to the RMP1-14 group. In groups Compound C-2.5 and Compound C-10, 1 and 2 mice, respectively, had complete tumor regression. After
- 25 treatment, tumor weight is significantly reduced by the RMP1-14 and three Compound C

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groups as compared to the control group. RMP1-14 group had a relative tumor inhibition value of 55.03%, with Compound C -2.5, Compound C -5, and Compound C -10 groups with relative tumor inhibition values of 69.70%, 76.53%, and 81.04%, respectively, showing a dose-dependent manner of Compound C on tumor weight. Thus, Compound C

5 has significant anti-tumor efficacy in tumor-bearing mice, and has better effect than the positive control clone RMP1-14.

**Amino Acid and Nucleotide Sequences**

SEQ ID NO: 1 (human PD-1)

MQIPQAPWPVVWAVLQLGWRPGWFLDSPDRPWNPPTFSPALLVVTEGDNATFT  
5 CSFSNTSESFVLNWYRMSPSNQTDKLAAFPEDRSQPGQDCRFRVTQLPNGRDFH  
MSVVRARRNDSTGYLCGAISLAPKAQIKESLRAELRVTERRAEVPTAHPSPSPRPA  
GQFQTLVVG VVGGLLGSLLVWVLAVICSRAARGTIGARRTGQPLKEDPSAVP  
VFSVDYGELDFQWREKTPEPPVPCVPEQTEYATIVFPSGMGTSSPARRGSADGPR  
SAQPLRPEDGHCSWPL

10

SEQ ID NO: 2 (HCDR1 of Antibody A)

KASGYTFTSYMH

SEQ ID NO: 3 (HCDR1 of Antibody B)

15 KASGYTFEGYYMH

SEQ ID NO: 4 (HCDR1 of Antibody C)

KASGYTFTAQYMH

20 SEQ ID NO: 5 (HCDR1 of Antibody D)

KASGYTFEKYYMH

SEQ ID NO: 6 (HCDR1 of Antibody E)

KASGYTFTSNYMH

25

SEQ ID NO: 7 (HCDR1 of Antibody F)

KASGYTFSAYMH

SEQ ID NO: 8 (HCDR2 of Antibody A)

30 IINPSGGSTSYAQKFQG



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SEQ ID NO: 9 (HCDR2 of Antibody B)

IINPEGGETSYAQKFQG

SEQ ID NO: 10 (HCDR2 of Antibody C)

5 IINPSGGETGYAQKFQG

SEQ ID NO: 11 (HCDR2 of Antibody E)

IINPSEGSTGYAQKFQG

10 SEQ ID NO: 12 (HCDR2 of Antibody D and F)

IINPDGGSTGYAQKFQG

SEQ ID NO: 13 (HCDR3 of Antibody A-F)

AKEGVADGYGLVDV

15

SEQ ID NO: 14 (LCDR1 of Antibody A-F)

RASQSVSSYLA

SEQ ID NO: 15 (LCDR2 of Antibody A-F)

20 YDASKRAT

SEQ ID NO: 16 (LCDR3 of Antibody A-F)

DQRNNWPLT

25 SEQ ID NO: 17 (HCVR of Antibody A)

QVQLVQSGAEVKKPGASVKVSCKASGYTFTSYMHWVRQAPGQGLEWMGIINP  
SGGSTSYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDVAVYYCAKEGVADGYG  
LVDVWGQGTMVTTIS

30 SEQ ID NO: 18 (HCVR of Antibody B)

QVQLVQSGAEVKKPGASVKVSCKASGYTFEGYYMHWVRQAPGQGLEWMGIIN  
PEGGETSYAQKFQGRVTMTRDTSISTAYMELSLRSDDTAVYYCAKEGVADGY  
GLVDVWGQGTMVTVSS

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SEQ ID NO: 19 (HCVR of Antibody C)

QVQLVQSGAEVKKPGASVKVSCKASGYTFTAQYMHWRQAPGQGLEWMGIIN  
 PSGGETGYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDVAVYYCAKEGVADGY  
 5 GLVDVWGQGTMTVTVSS

SEQ ID NO: 20 (HCVR of Antibody D)

QVQLVQSGAEVKKPGASVKVSCKASGYTFEKYYMHWRQAPGQGLEWMGIIN  
 PDGGSTGYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDVAVYYCAKEGVADGY  
 10 GLVDVWGQGTMTVTVSS

SEQ ID NO: 21 (HCVR of Antibody E)

QVQLVQSGAEVKKPGASVKVSCKASGYTFTSNYMHWRQAPGQGLEWMGIINP  
 SEGSTGYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDVAVYYCAKEGVADGYG  
 15 LVDVWGQGTMTVTVSS

SEQ ID NO: 22 (HCVR of Antibody F)

QVQLVQSGAEVKKPGASVKVSCKASGYTFSAYYMHWRQAPGQGLEWMGIINP  
 DGGSTGYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDVAVYYCAKEGVADGYG  
 20 LVDVWGQGTMTVTVSS

SEQ ID NO: 23 (LCVR of Antibody A-F)

EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASKRAT  
 GIPARFSGSGSGTDFTLTISLEPEDFAVYYCDQRNNWPLTFGGGGTKVEIK  
 25

SEQ ID NO: 24 (HC of Antibody A – S228P IgG4)

QVQLVQSGAEVKKPGASVKVSCKASGYTFTSYMHWRQAPGQGLEWMGIINP  
 SGGSTSYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDVAVYYCAKEGVADGYG  
 LVDVWGQGTMTVTVSSASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVS  
 30 WNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTKTYTCNVDHKPSNTKVD  
 KRVESKYGPPCPPCPAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDP  
 EVQFNWYVDGVEVHNAKTKPREEQFNSTYRVVSVLTVLHQDWLNGKEYKCKV  
 SNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAV  
 EWESNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMHEALH  
 35 NHYTQKSLSLSLGK

SEQ ID NO: 25 (HC of Antibody B – S228P IgG4)

QVQLVQSGAEVKKPGASVKVSCKASGYTFEGYYMHWRQAPGQGLEWMGIIN  
 PEGGETSYAQKFQGRVTMTRDTSISTAYMELSLRSDDVAVYYCAKEGVADGY  
 40 GLVDVWGQGTMTVTVSSASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTV  
 SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTKTYTCNVDHKPSNTKV

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DKR VESKYGPPCPPCPAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQED  
 PEVQFNWYVDGVEVHNAKTKPREEQFNSTYRVVSVLTVLHQDWLNGKEYKCK  
 VSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIA  
 VEWESNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMHEAL  
 5 HNHYTQKSLSLSLGK

SEQ ID NO: 26 (HC of Antibody C – S228P IgG4)

QVQLVQSGAEVKKPGASVKVSCKASGYTFTAQYMHWRQAPGQGLEWMGIIN  
 PSGGETGYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCAKEGVADGY  
 10 GLVDVWGQGTMTVTSSASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTV  
 SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTKTYTCNVDHKPSNTKV  
 DKR VESKYGPPCPPCPAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQED  
 PEVQFNWYVDGVEVHNAKTKPREEQFNSTYRVVSVLTVLHQDWLNGKEYKCK  
 VSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIA  
 15 VEWESNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMHEAL  
 HNHYTQKSLSLSLGK

SEQ ID NO: 27 (HC of Antibody D – S228P IgG4)

QVQLVQSGAEVKKPGASVKVSCKASGYTFEKYYMHWRQAPGQGLEWMGIIN  
 20 PDGGSTGYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCAKEGVADGY  
 GLVDVWGQGTMTVTSSASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTV  
 SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTKTYTCNVDHKPSNTKV  
 DKR VESKYGPPCPPCPAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQED  
 PEVQFNWYVDGVEVHNAKTKPREEQFNSTYRVVSVLTVLHQDWLNGKEYKCK  
 25 VSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIA  
 VEWESNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMHEAL  
 HNHYTQKSLSLSLGK

SEQ ID NO: 28 (HC of Antibody E – S228P IgG4)

QVQLVQSGAEVKKPGASVKVSCKASGYTFTSNYMHWRQAPGQGLEWMGIINP  
 30 SEGSTGYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCAKEGVADGYG  
 LVDVWGQGTMTVTSSASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVS  
 WNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTKTYTCNVDHKPSNTKVD  
 KR VESKYGPPCPPCPAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDP  
 35 EVQFNWYVDGVEVHNAKTKPREEQFNSTYRVVSVLTVLHQDWLNGKEYKCKV  
 SNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAV  
 EWESNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMHEALH  
 NHYTQKSLSLSLGK

40 SEQ ID NO: 29 (HC of Antibody F – S228P IgG4)

QVQLVQSGAEVKKPGASVKVSCKASGYTFSAYYMHWRQAPGQGLEWMGIINP  
 DGGSTGYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCAKEGVADGYG  
 LVDVWGQGTMTVTSSASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVS

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WNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGKTYTCNVDPHKPSNTKVD  
 KRVESKYGPPCPPCPAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDP  
 EVQFNWYVDGVEVHNAKTKPREEQFNSTYRVVSVLTVLHQDWLNGKEYKCKV  
 SNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAV  
 5 EWESNGQPENNYKTTTPVLDSGDSFFLYSRLTVDKSRWQEGNVFSCSVMHEALH  
 NHYTQKSLSLSLGK

SEQ ID NO: 30 (LC of Antibody A-F)

EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASKRAT  
 10 GIPARFSGSGSGTDFTLTISLEPEDFAVYYCDQRNNWPLTFGGGTKVEIKRTVAA  
 PSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQD  
 SKDSTYLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC

SEQ ID NO: 31 (DNA of HC of Antibody A S228P IgG4)

15 CAGGTGCAGCTGGTGCAGTCTGGGGCTGAGGTGAAGAAGCCTGGGGCCTCAG  
 TGAAGGTTTCTGCAAGGCATCTGGATACACCTTCACCAGCTACTATATGCAC  
 TGGGTGCGACAGGCCCTGGACAAGGGCTTGAGTGGATGGGAATAATCAACC  
 CTAGTGGTGGTAGCACAAGCTACGCACAGAAGTTCCAGGGCAGAGTCACCAT  
 GACCAGGGACACGTCCACGAGCACAGTCTACATGGAGCTGAGCAGCCTGAGA  
 20 TCTGAGGACACGGCGGTGTACTACTGCGCCAAAGAGGGAGTGGCCGACGGAT  
 ATGGATTGGTAGACGTATGGGGTCAGGGTACAATGGTCACCATCTCCTCAGC  
 CAGCACCAAGGGACCCTCCGTGTTCCCTCTGGCTCCTTGCAGCAGGTCCACCA  
 GCGAATCCACCGCTGCCCTGGGCTGTCTGGTGAAAGACTACTTTCCCGAGCCT  
 GTGACCGTGAGCTGGAACCTCCGGCGCTCTGACCAGCGGCGTGACACATTTC  
 25 CTGCCGTGCTGCAGAGCTCCGGCCTGTACTCCCTGTCCTCCGTGGTGACAGTC  
 CCCAGCAGCAGCCTGGGAACCAAGACCTACACCTGCAACGTCGACCACAAGC  
 CTTCCAACACCAAGGTGGACAAGAGGGTGGAGTCCAAATATGGCCCCCCTG  
 CCCTCCTTGTCCCGCTCCTGAGTTCCCTGGGCGGCCCTTCCGTGTTCCCTGTTCCC  
 TCCCAAGCCCAAGGACACCCTGATGATCTCCCGGACCCCCGAGGTGACCTGT  
 30 GTGGTGGTGGACGTGTCCCAGGAGGACCCTGAGGTGCAATTCAACTGGTACG  
 TGGACGGCGTCGAGGTGCACAACGCCAAGACCAAGCCCCGGGAGGAGCAGT  
 TCAACAGCACCTACCGGGTCGTGTCCGTGCTGACCGTGCTGCACCAGGATTGG  
 CTAACGGCAAGGAGTACAAGTGCAAAGTGCCAATAAGGGCCTGCCCTCCT  
 CCATCGAGAAGACCATCTCCAAGGCCAAGGGACAACCCCGTGAGCCCCAGGT  
 35 GTACACCCTGCCTCCTTCCCAGGAGGAGATGACCAAGAATCAGGTGTCCCTC  
 ACCTGCCTGGTGAAGGGCTTCTACCCTTCCGACATCGCCGTGGAATGGGAGTC  
 CAACGGCCAGCCCGAGAACAACTACAAGACAACCCCCCTGTCCTGGACAGC  
 GACGGCTCCTTCTTTCTGTACAGCAGGCTGACCGTGGACAAGAGCCGGTGGC  
 AGGAGGGCAACGTGTTTAGCTGTAGCGTCATGCACGAGGCCCTGCACAACCA  
 40 CTACACCCAGAAATCCCTGTCCCTGTCCCTGGGCAAGTGATGA

SEQ ID NO: 32 (DNA of HC of Antibody B S228P IgG4)

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CAGGTGCAGCTGGTGCAGTCTGGGGCTGAGGTGAAGAAGCCTGGGGCCTCAG  
TGAAGGTTTCCTGCAAGGCATCTGGATACACCTTCGAGGGTTACTATATGCAC  
TGGGTGCGACAGGCCCCCTGGACAAGGGCTTGAGTGGATGGGAATAATCAACC  
CTGAGGGTGGTGAAGCTACGCACAGAAGTTCCAGGGCAGAGTCACCAT  
5 GACCAGGGACACGTCCATCAGCACAGCCTACATGGAGCTGAGCAGGCTGAGA  
TCTGACGACACGGCGGTGTACTACTGCGCCAAAGAGGGAGTGGCCGACGGAT  
ATGGATTGGTAGACGTATGGGGTCAGGGTACAATGGTCACCGTCTCCTCAGC  
CAGCACCAAGGGACCCCTCCGTGTTCCCTCTGGCTCCTTGACGACAGGTCCACCA  
GCGAATCCACCGCTGCCCTGGGCTGTCTGGTGAAAGACTACTTTCCCGAGCCT  
10 GTGACCGTGAGCTGGAACCTCCGGCGCTCTGACCAGCGGCGTGCACACATTTTC  
CTGCCGTGCTGCAGAGCTCCGGCCTGTACTCCCTGTCCTCCGTGGTGACAGTC  
CCCAGCAGCAGCCTGGGAACCAAGACCTACACCTGCAACGTCGACCACAAGC  
CTTCCAACACCAAGGTGGACAAGAGGGTGGAGTCCAAATATGGCCCCCCTG  
CCCTCCTTGTCCCGCTCCTGAGTTCCTGGGCGGCCCTTCCGTGTTCTGTTCCC  
15 TCCCAAGCCCAAGGACACCCTGATGATCTCCCGGACCCCGAGGTGACCTGT  
GTGGTGGTGGACGTGTCCCAGGAGGACCCTGAGGTGCAATTCAACTGGTACG  
TGGACGGCGTCGAGGTGCACAACGCCAAGACCAAGCCCCGGGAGGAGCAGT  
TCAACAGCACCTACCGGGTCGTGTCCGTGCTGACCGTGCTGCACCAGGATTGG  
CTCAACGGCAAGGAGTACAAGTGCAAAGTGTCCAATAAGGGCCTGCCCTCCT  
20 CCATCGAGAAGACCATCTCCAAGGCCAAGGGACAACCCCGTGAGCCCCAGGT  
GTACACCCTGCCTCCTTCCCAGGAGGAGATGACCAAGAATCAGGTGTCCCTC  
ACCTGCCTGGTGAAGGGCTTCTACCCTTCCGACATCGCCGTGGAATGGGAGTC  
CAACGGCCAGCCCGAGAACAATAACAAGACAACCCCCCTGTCTGGACAGC  
GACGGCTCCTTCTTTCTGTACAGCAGGCTGACCGTGGACAAGAGCCGGTGGC  
25 AGGAGGGCAACGTGTTTAGCTGTAGCGTCATGCACGAGGCCCTGCACAACCA  
CTACACCCAGAAATCCCTGTCCCTGTCCCTGGGCAAGTGATGA

SEQ ID NO: 33 (DNA of HC of Antibody C S228P IgG4)

CAGGTGCAGCTGGTGCAGTCTGGGGCTGAGGTGAAGAAGCCTGGGGCCTCAG  
30 TGAAGGTTTCCTGCAAGGCATCTGGATACACCTTCACCGCTCAGTATATGCAC  
TGGGTGCGACAGGCCCCCTGGACAAGGGCTTGAGTGGATGGGAATAATCAACC  
CTAGTGGTGGTGAAGCTACGCACAGAAGTTCCAGGGCAGAGTCACCAT  
GACCAGGGACACGTCCACGAGCACAGTCTACATGGAGCTGAGCAGCCTGAGA  
TCTGAGGACACGGCGGTGTACTACTGCGCCAAAGAGGGAGTGGCCGACGGAT  
35 ATGGATTGGTAGACGTATGGGGTCAGGGTACAATGGTCACCGTCTCCTCAGC  
CAGCACCAAGGGACCCCTCCGTGTTCCCTCTGGCTCCTTGACGACAGGTCCACCA  
GCGAATCCACCGCTGCCCTGGGCTGTCTGGTGAAAGACTACTTTCCCGAGCCT  
GTGACCGTGAGCTGGAACCTCCGGCGCTCTGACCAGCGGCGTGCACACATTTTC  
CTGCCGTGCTGCAGAGCTCCGGCCTGTACTCCCTGTCCTCCGTGGTGACAGTC  
40 CCCAGCAGCAGCCTGGGAACCAAGACCTACACCTGCAACGTCGACCACAAGC  
CTTCCAACACCAAGGTGGACAAGAGGGTGGAGTCCAAATATGGCCCCCCTG  
CCCTCCTTGTCCCGCTCCTGAGTTCCTGGGCGGCCCTTCCGTGTTCTGTTCCC  
TCCCAAGCCCAAGGACACCCTGATGATCTCCCGGACCCCGAGGTGACCTGT  
GTGGTGGTGGACGTGTCCCAGGAGGACCCTGAGGTGCAATTCAACTGGTACG  
45 TGGACGGCGTCGAGGTGCACAACGCCAAGACCAAGCCCCGGGAGGAGCAGT

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TCAACAGCACCTACCGGGTCGTGTCCGTGCTGACCGTGCTGCACCAGGATTGG  
 CTCAACGGCAAGGAGTACAAGTGCAAAGTGTCCAATAAGGGCCTGCCCTCCT  
 CCATCGAGAAGACCATCTCCAAGGCCAAGGGACAACCCCGTGAGCCCCAGGT  
 GTACACCCTGCCTCCTTCCCAGGAGGAGATGACCAAGAATCAGGTGTCCCTC  
 5 ACCTGCCTGGTGAAGGGCTTCTACCCTTCCGACATCGCCGTGGAATGGGAGTC  
 CAACGGCCAGCCCGAGAACAACACTACAAGACAACCCCCCCTGTCCTGGACAGC  
 GACGGCTCCTTCTTTCTGTACAGCAGGCTGACCGTGGACAAGAGCCGGTGGC  
 AGGAGGGCAACGTGTTTAGCTGTAGCGTCATGCACGAGGCCCTGCACAACCA  
 CTACACCCAGAAATCCCTGTCCCTGTCCCTGGGCAAGTGATGA

10

SEQ ID NO: 34 (DNA of HC of Antibody D S228P IgG4)

CAGGTGCAGCTGGTGCAGTCTGGGGCTGAGGTGAAGAAGCCTGGGGCCTCAG  
 TGAAGGTTTCCTGCAAGGCATCTGGATACACCTTCGAGAAGTACTATATGCAC  
 TGGGTGCGACAGGCCCTGGACAAGGGCTTGAGTGGATGGGAATAATCAACC  
 15 CTGATGGTGGTAGCACAGGGTACGCACAGAAGTTCCAGGGCAGAGTCACCAT  
 GACCAGGGACACGTCCACGAGCACAGTCTACATGGAGCTGAGCAGCCTGAGA  
 TCTGAGGACACGGCGGTGTACTACTGCGCCAAAGAGGGAGTGGCCGACGGAT  
 ATGGATTGGTAGACGTATGGGGTCAGGGTACAATGGTCACCGTCTCCTCAGC  
 CAGCACCAAGGGACCCTCCGTGTTCCCTCTGGCTCCTTGCAGCAGGTCCACCA  
 20 GCGAATCCACCGCTGCCCTGGGCTGTCTGGTGAAAGACTACTTTCCCGAGCCT  
 GTGACCGTGAGCTGGAACCTCCGGCGCTCTGACCAGCGGCGTGCACACATTTC  
 CTGCCGTGCTGCAGAGCTCCGGCCTGTACTCCCTGTCTCCGTGGTGACAGTC  
 CCCAGCAGCAGCCTGGGAACCAAGACCTACACCTGCAACGTCGACCACAAGC  
 CTTCCAACACCAAGGTGGACAAGAGGGTGGAGTCCAAATATGGCCCCCCTG  
 25 CCCTCCTTGTCCCGCTCCTGAGTTCTTGGGCGGCCCTTCCGTGTTCTGTTCCC  
 TCCCAAGCCCAAGGACACCCTGATGATCTCCCGGACCCCCGAGGTGACCTGT  
 GTGGTGGTGGACGTGTCCCAGGAGGACCCTGAGGTGCAATTCAACTGGTACG  
 TGGACGGCGTCGAGGTGCACAACGCCAAGACCAAGCCCCGGGAGGAGCAGT  
 TCAACAGCACCTACCGGGTCGTGTCCGTGCTGACCGTGCTGCACCAGGATTGG  
 30 CTCAACGGCAAGGAGTACAAGTGCAAAGTGTCCAATAAGGGCCTGCCCTCCT  
 CCATCGAGAAGACCATCTCCAAGGCCAAGGGACAACCCCGTGAGCCCCAGGT  
 GTACACCCTGCCTCCTTCCCAGGAGGAGATGACCAAGAATCAGGTGTCCCTC  
 ACCTGCCTGGTGAAGGGCTTCTACCCTTCCGACATCGCCGTGGAATGGGAGTC  
 CAACGGCCAGCCCGAGAACAACACTACAAGACAACCCCCCCTGTCCTGGACAGC  
 35 GACGGCTCCTTCTTTCTGTACAGCAGGCTGACCGTGGACAAGAGCCGGTGGC  
 AGGAGGGCAACGTGTTTAGCTGTAGCGTCATGCACGAGGCCCTGCACAACCA  
 CTACACCCAGAAATCCCTGTCCCTGTCCCTGGGCAAGTGATGA

SEQ ID NO: 35 (DNA of HC of Antibody E S228P IgG4)

40 CAGGTGCAGCTGGTGCAGTCTGGGGCTGAGGTGAAGAAGCCTGGGGCCTCAG  
 TGAAGGTTTCCTGCAAGGCATCTGGATACACCTTCACCAGCAATTATATGCAC  
 TGGGTGCGACAGGCCCTGGACAAGGGCTTGAGTGGATGGGAATAATCAACC  
 CTAGTGAGGGTAGCACAGGTTACGCACAGAAGTTCCAGGGCAGAGTCACCAT  
 GACCAGGGACACGTCCACGAGCACAGTCTACATGGAGCTGAGCAGCCTGAGA

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TCTGAGGACACGGCGGTGTACTACTGCGCCAAAGAGGGAGTGGCCGACGGAT  
ATGGATTGGTAGACGTATGGGGTCAGGGTACAATGGTCACCGTCTCCTCAGC  
CAGCACCAAGGGACCCTCCGTGTTCCCTCTGGCTCCTTGCAGCAGGTCCACCA  
GCGAATCCACCGCTGCCCTGGGCTGTCTGGTGAAAGACTACTTTCCCGAGCCT  
5 GTGACCGTGAGCTGGAACCTCCGGCGCTCTGACCAGCGGCGTGCACACATTTC  
CTGCCGTGCTGCAGAGCTCCGGCCTGTACTCCCTGTCTCCTCCGTGGTGACAGTC  
CCCAGCAGCAGCCTGGGAACCAAGACCTACACCTGCAACGTCGACCACAAGC  
CTTCCAACACCAAGGTGGACAAGAGGGTGGAGTCCAAATATGGCCCCCCTG  
CCCTCCTTGTCCCGCTCCTGAGTTCCTGGGCGGCCCTTCCGTGTTCTGTTCCC  
10 TCCCAAGCCCAAGGACACCCTGATGATCTCCCGGACCCCCGAGGTGACCTGT  
GTGGTGGTGGACGTGTCCCAGGAGGACCCTGAGGTGCAATTCAACTGGTACG  
TGGACGGCGTCGAGGTGCACAACGCCAAGACCAAGCCCCGGGAGGAGCAGT  
TCAACAGCACCTACCGGGTTCGTGTCCGTGCTGACCGTGCTGCACCAGGATTGG  
CTCAACGGCAAGGAGTACAAGTGCAAAGTGTCCAATAAGGGCCTGCCCTCCT  
15 CCATCGAGAAGACCATCTCCAAGGCCAAGGGACAACCCCGTGAGCCCCAGGT  
GTACACCCTGCCTCCTTCCCAGGAGGAGATGACCAAGAATCAGGTGTCCCTC  
ACCTGCCTGGTGAAGGGCTTCTACCCTTCCGACATCGCCGTGGAATGGGAGTC  
CAACGGCCAGCCCGAGAACAATAAGACAACCCCCCTGTCCTGGACAGC  
GACGGCTCCTTCTTTCTGTACAGCAGGCTGACCGTGGACAAGAGCCGGTGGC  
20 AGGAGGGCAACGTGTTTAGCTGTAGCGTCATGCACGAGGCCCTGCACAACCA  
CTACACCCAGAAATCCCTGTCCCTGTCCCTGGGCAAGTGATGA

SEQ ID NO: 36 (DNA of HC of Antibody F S228P IgG4)

CAGGTGCAGCTGGTGCAGTCTGGGGCTGAGGTGAAGAAGCCTGGGGCCTCAG  
25 TGAAGGTTTCTGCAAGGCATCTGGATACACCTTCAGTGCGTACTATATGCAC  
TGGGTGCGACAGGCCCTGGACAAGGGCTTGAGTGGATGGGAATAATCAACC  
CTGATGGTGGTAGCACAGGGTACGCACAGAAGTTCCAGGGCAGAGTCACCAT  
GACCAGGGACACGTCCACGAGCACAGTCTACATGGAGCTGAGCAGCCTGAGA  
TCTGAGGACACGGCGGTGTACTACTGCGCCAAAGAGGGAGTGGCCGACGGAT  
30 ATGGATTGGTAGACGTATGGGGTCAGGGTACAATGGTCACCGTCTCCTCAGC  
CAGCACCAAGGGACCCTCCGTGTTCCCTCTGGCTCCTTGCAGCAGGTCCACCA  
GCGAATCCACCGCTGCCCTGGGCTGTCTGGTGAAAGACTACTTTCCCGAGCCT  
GTGACCGTGAGCTGGAACCTCCGGCGCTCTGACCAGCGGCGTGCACACATTTC  
CTGCCGTGCTGCAGAGCTCCGGCCTGTACTCCCTGTCTCCTCCGTGGTGACAGTC  
35 CCCAGCAGCAGCCTGGGAACCAAGACCTACACCTGCAACGTCGACCACAAGC  
CTTCCAACACCAAGGTGGACAAGAGGGTGGAGTCCAAATATGGCCCCCCTG  
CCCTCCTTGTCCCGCTCCTGAGTTCCTGGGCGGCCCTTCCGTGTTCTGTTCCC  
TCCCAAGCCCAAGGACACCCTGATGATCTCCCGGACCCCCGAGGTGACCTGT  
GTGGTGGTGGACGTGTCCCAGGAGGACCCTGAGGTGCAATTCAACTGGTACG  
40 TGGACGGCGTCGAGGTGCACAACGCCAAGACCAAGCCCCGGGAGGAGCAGT  
TCAACAGCACCTACCGGGTTCGTGTCCGTGCTGACCGTGCTGCACCAGGATTGG  
CTCAACGGCAAGGAGTACAAGTGCAAAGTGTCCAATAAGGGCCTGCCCTCCT  
CCATCGAGAAGACCATCTCCAAGGCCAAGGGACAACCCCGTGAGCCCCAGGT  
GTACACCCTGCCTCCTTCCCAGGAGGAGATGACCAAGAATCAGGTGTCCCTC  
45 ACCTGCCTGGTGAAGGGCTTCTACCCTTCCGACATCGCCGTGGAATGGGAGTC

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CAACGGCCAGCCCGAGAACAACCTACAAGACAACCCCCCTGTCCTGGACAGC  
GACGGCTCCTTCTTTCTGTACAGCAGGCTGACCGTGGACAAGAGCCGGTGGC  
AGGAGGGCAACGTGTTTAGCTGTAGCGTCATGCACGAGGCCCTGCACAACCA  
CTACACCCAGAAATCCCTGTCCCTGTCCCTGGGCAAGTGATGA

5

SEQ ID NO: 37 (DNA of LC of Antibody C and E)

GAAATTGTGTTGACACAGTCTCCAGCCACCCTGTCTTTGTCTCCAGGGGAAAG  
AGCCACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGCAGCTACTTAGCCTGGT  
ACCAACAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGATGCATCCAA  
10 AAGGGCCACTGGCATCCCAGCCAGGTTCAAGTGGCAGTGGGTCTGGGACAGAC  
TTCACCTCTACCATCAGCAGCCTAGAGCCTGAAGATTTTGCAGTTTATTACTG  
TGACCAGAGAAACAATTGGCCTCTCACTTTTGGCGGAGGGACCAAGGTTGAG  
ATCAAACGGACCGTGGCTGCCCCTAGCGTGTTTCATCTTCCCTCCCTCCGATGA  
GCAGCTCAAGTCCGGCACAGCCAGCGTGGTGTGCCTGCTGAATAACTTCTACC  
15 CCCGGGAGGCCAAAGTGCAGTGGAAGGTGGACAACGCTCTGCAGTCCGGCAA  
TTCCCAGGAGAGCGTCACCGAGCAGGACAGCAAGGACAGCACCTACTCCCTG  
AGCTCCACCCTGACCCTGAGCAAGGCCGACTACGAGAAGCACAAGGTCTACG  
CCTGCGAGGTCAACCATCAGGGCCTGTCCAGCCCCGTGACCAAGTCCTTCAAC  
CGGGGCGAGTGCTGATGA

20

SEQ ID NO: 38 (mouse PD-1)

MWVRQVPWSFTWAVLQLSWQSGWLLEVPNGPWRS�TFYPAWLTVSEGANATF  
TCSLSNWSEDLMLNWNRLSPSNQTEKQAAFCNGLSQPVQDARFQIIQLPNRHDF  
HMNILDTRRNDSGIYLCGAISLHPKAKIEESPGAELVVTERILETSTRYPSPSPKPE  
25 GRFQGMVIGIMSALVGIPVLLLLAWALAVFCSTSMSEARGAGSKDDTLKEEPSAA  
PVPSVAYEELDFQGREKTPELPTACVHTEYATIVFTEGLGASAMGRRGSADGLQG  
PRPPRHEDGHCSWPL



## WE CLAIM:

1. An antibody that binds human PD-1 (SEQ ID NO: 1), comprising a light chain (LC) and a heavy chain (HC), wherein the light chain comprises light chain complementarity determining regions LCDR1, LCDR2, and LCDR3 consisting of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein the heavy chain comprises heavy chain complementarity determining regions HCDR1, HCDR2, and HCDR3, wherein HCDR1 consists of the amino acid sequence:
- KASGYTFTSYMH (SEQ ID NO: 2),
  - KASGYTFEGYYMH (SEQ ID NO: 3),
  - KASGYTFTAQYMH (SEQ ID NO: 4),
  - KASGYTFEKYYMH (SEQ ID NO: 5),
  - KASGYTFTSNYMH (SEQ ID NO: 6), or
  - KASGYTFSAYMH (SEQ ID NO: 7);
- wherein HCDR2 consists of the amino acid sequence:
- IINPSGGSTSYAQKFQG (SEQ ID NO: 8),
  - IINPEGGETSYAQKFQG (SEQ ID NO: 9),
  - IINPSGGETGYAQKFQG (SEQ ID NO: 10),
  - IINPSEGSTGYAQKFQG (SEQ ID NO: 11), or
  - IINPDGGSTGYAQKFQG (SEQ ID NO: 12);
- and wherein HCDR3 consists of the amino acid sequence AKEGVADGYGLVDV (SEQ ID NO: 13).
2. The antibody of Claim 1, wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFTSYMH (SEQ ID NO: 2),

IINPSGGSTSYAQKFQG (SEQ ID NO: 8), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.

3. The antibody of Claim 1, wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14),  
5 YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFEGYYMH (SEQ ID NO: 3), IINPEGGETSYAQKFQG (SEQ ID NO: 9), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.
- 10 4. The antibody of Claim 1, wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFTAQYMH (SEQ ID NO: 4),  
15 IINPSGGETGYAQKFQG (SEQ ID NO: 10), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.
5. The antibody of Claim 1, wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16),  
20 respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFEKYYMH (SEQ ID NO: 5), IINPDGGSTGYAQKFQG (SEQ ID NO: 12), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.
- 25 6. The antibody of Claim 1, wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFTSNYMH (SEQ ID NO: 6), IINPSEGSTGYAQKFQG (SEQ ID NO: 11), and AKEGVADGYGLVDV  
30 (SEQ ID NO: 13), respectively.

- 5 7. The antibody of Claim 1, wherein LCDR1, LCDR2, and LCDR3 consist of the amino acid sequences RASQSVSSYLA (SEQ ID NO: 14), YDASKRAT (SEQ ID NO: 15), and DQRNNWPLT (SEQ ID NO: 16), respectively, and wherein HCDR1, HCDR2, and HCDR3 consist of the amino acid sequences KASGYTFSAYYMH (SEQ ID NO: 7), IINPDGGSTGYAQKFQG (SEQ ID NO: 12), and AKEGVADGYGLVDV (SEQ ID NO: 13), respectively.
- 10 8. An antibody, comprising a light chain (LC) and a heavy chain (HC), wherein the light chain comprises a light chain variable region (LCVR) and the heavy chain comprises a heavy chain variable region (HCVR), wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 17, SEQ ID NO: 18, SEQ ID NO: 19, SEQ ID NO: 20, SEQ ID NO: 21, or SEQ ID NO: 22.
- 15 9. The antibody of Claim 8, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 17.
- 20 10. The antibody of Claim 8, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 18.
- 25 11. The antibody of Claim 8, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 19.
12. The antibody of Claim 8, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 20.
13. The antibody of Claim 8, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 21.

14. The antibody of Claim 8, wherein the LCVR has the amino acid sequence given in SEQ ID NO: 23, and the HCVR has the amino acid sequence given in SEQ ID NO: 22.
- 5 15. The antibody of Claim 8, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 24, SEQ ID NO: 25, SEQ ID NO: 26, SEQ ID NO: 27, SEQ ID NO: 28, or SEQ ID NO: 29.
- 10 16. The antibody of Claim 15, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 24.
- 15 17. The antibody of Claim 15, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 25.
18. The antibody of Claim 15, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 26.
- 20 19. The antibody of Claim 15, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 27.
21. The antibody of Claim 15, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 28.
- 25 22. The antibody of Claim 15, wherein the LC has the amino acid sequence given in SEQ ID NO: 30, and the HC has the amino acid sequence given in SEQ ID NO: 29.
- 30 23. The antibody of Claim 15, comprising two light chains and two heavy chains, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 24, SEQ ID NO: 25, SEQ ID NO: 26, SEQ ID NO: 27, SEQ ID NO: 28, or SEQ ID NO: 29.

23. The antibody of Claim 22, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 24.
- 5 24. The antibody of Claim 22, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 25.
25. The antibody of Claim 22, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 26.
- 10 26. The antibody of Claim 22, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 27.
27. The antibody of Claim 22, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 28.
- 15 28. The antibody of Claim 22, wherein each light chain has the amino acid sequence given in SEQ ID NO: 30, and each heavy chain has the amino acid sequence given in SEQ ID NO: 29.
29. The antibody of any one of Claims 22-28, wherein one of the heavy chains forms an inter-chain disulfide bond with one of the light chains, and the other heavy chain forms an inter-chain disulfide bond with the other light chain, and one of the heavy chains forms two inter-chain disulfide bonds with the other heavy chain.
- 20 30. The antibody of any one of Claims 1-29, wherein the antibody is glycosylated.
- 25 31. A pharmaceutical composition, comprising the antibody of any one of Claims 1-30, and an acceptable carrier, diluent, or excipient.
32. A method of treating cancer, comprising administering to a patient in need thereof, an effective amount of the antibody of any one of Claims 1-30.
- 30 33. The method of Claim 32, wherein the cancer is melanoma, lung cancer, head and neck cancer, colorectal cancer, pancreatic cancer, gastric cancer,

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kidney cancer, bladder cancer, prostate cancer, breast cancer, ovarian cancer, or hepatocellular carcinoma.

34. The method of Claim 32 or 33, further comprising administering simultaneously, separately, or sequentially one or more anti-tumor agents.

5 35. The antibody of any one of Claims 1-30, for use in therapy.

36. The antibody of any one of Claims 1-30, for use in the treatment of cancer.

37. The antibody for use of Claim 36, wherein the cancer is melanoma, lung cancer, head and neck cancer, colorectal cancer, pancreatic cancer, gastric cancer, kidney cancer, bladder cancer, prostate cancer, breast cancer, ovarian cancer, or hepatocellular carcinoma.

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38. The antibody of any one of Claims 1-30 for use in simultaneous, separate, or sequential combination with one or more anti-tumor agents, in the treatment of cancer.

39. The combination for use of Claim 38, wherein the cancer is melanoma, lung cancer, head and neck cancer, colorectal cancer, pancreatic cancer, gastric cancer, kidney cancer, bladder cancer, prostate cancer, breast cancer, ovarian cancer, or hepatocellular carcinoma.

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## INTERNATIONAL SEARCH REPORT

International application No.

**PCT/CN2016/073169****A. CLASSIFICATION OF SUBJECT MATTER**

C07K 16/28(2006.01)i; C12N 5/10(2006.01)i; A61K 39/395(2006.01)i; A61P 35/00(2006.01)i

According to International Patent Classification (IPC) or to both national classification and IPC

**B. FIELDS SEARCHED**

Minimum documentation searched (classification system followed by classification symbols)

C07K; C12N; A61K; A61P

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

DWPI; SIPOABS; CPRSABS; CNKI; NCBI; ISI Web of knowledge; ELSEVIER and keywords: programmed cell death 1, PD-1, antibody, etc. GenBank; EMBL; Retrieving System for Biological Sequence of Chinese Patent and searched sequences: SEQ ID NOs: 2-29

**C. DOCUMENTS CONSIDERED TO BE RELEVANT**

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	CN 103242448 A (UNIV ZHENGZHOU) 14 August 2013 (2013-08-14) the abstract, claims 1-8	1-31, 35-39
A	CN 104250302 A (SHANGHAI JUNSHI BIOSCIENCES CO LTD) 31 December 2014 (2014-12-31) the abstract, claims 1-15	1-31, 35-39

☐ Further documents are listed in the continuation of Box C.☒ See patent family annex.

\* Special categories of cited documents:

“A” document defining the general state of the art which is not considered to be of particular relevance

“E” earlier application or patent but published on or after the international filing date

“L” document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

“O” document referring to an oral disclosure, use, exhibition or other means

“P” document published prior to the international filing date but later than the priority date claimed

“T” later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

“X” document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

“Y” document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

“&amp;” document member of the same patent family

Date of the actual completion of the international search

**02 November 2016**

Date of mailing of the international search report

**08 November 2016**

Name and mailing address of the ISA/CN

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# INTERNATIONAL SEARCH REPORT

International application No.

**PCT/CN2016/073169**

**Box No. I**      **Nucleotide and/or amino acid sequence(s) (Continuation of item 1.c of the first sheet)**

1. With regard to any nucleotide and/or amino acid sequence disclosed in the international application, the international search was carried out on the basis of a sequence listing filed or furnished:
  - a. (means)  
☐ on paper  
☒ in electronic form
  - b. (time)  
☐ in the international application as filed  
☒ together with the international application in electronic form  
☐ subsequently to this Authority for the purposes of search
2. ☐ In addition, in the case that more than one version or copy of a sequence listing has been filed or furnished, the required statements that the information in the subsequent or additional copies is identical to that in the application as filed or does not go beyond the application as filed, as appropriate, were furnished.
3. Additional comments:
  - [1] The sequence listing contained in the description does not comply with the standard provided for in Annex C of the Administrative Instructions.



# INTERNATIONAL SEARCH REPORT

International application No.

**PCT/CN2016/073169**

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

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

1. ☒ Claims Nos.: **32-34**  
because they relate to subject matter not required to be searched by this Authority, namely:  
[1] Rule 39.1(iv) PCT- Method for treatment of the human or animal body by surgery or therapy.
2. ☐ Claims Nos.:  
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

**INTERNATIONAL SEARCH REPORT**  
**Information on patent family members**

International application No.

**PCT/CN2016/073169**

Patent document cited in search report			Publication date (day/month/year)	Patent family member(s)			Publication date (day/month/year)
CN	103242448	A	14 August 2013	CN	103242448	B	14 January 2015
CN	104250302	A	31 December 2014	EP	3026062	A1	01 June 2016
				PH	12015502819	A1	21 March 2016
				US	2016272708	A1	22 September 2016
				WO	2014206107	A1	31 December 2014
				JP	2016523265	A	08 August 2016