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(54) ISOLATED ANTIBODY WHICH BINDS TO AN EPITOPE WITHIN THE EXTRACELLULAR DOMAIN OF PD - 1

(54) נוגדן מבודד הנקשר לאפיטופ באיזור החוץ תאי של DP–1

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נוגדן מבודד הנקשר לאפיטופ באיזור החוץ תאי של PD -1

An isolated antibody which binds to an epitope within the extracellular domain of PD-1

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DESCRIPTION OF THE INVENTION

Field of the Invention

[01] The technical field relates to modulation of immune responses regulated by the Programmed Death 1 (PD-1) receptor.

Background of the Invention

[02] An adaptive immune response involves activation, selection, and clonal proliferation of two major classes of lymphocytes termed T cells and B cells. After encountering an antigen, T cells proliferate and differentiate into antigen-specific effector cells, while B cells proliferate and differentiate into antibody-secreting cells.

[03] T cell activation is a multi-step process requiring several signaling events between the T cell and an antigen-presenting cell (APC). For T cell activation to occur, two types of signals must be delivered to a resting T cell. The first type is mediated by the antigen-specific T cell receptor (TcR), and confers specificity to the immune response. The second, costimulatory, type regulates the magnitude of the response and is delivered through accessory receptors on the T cell.

[04] A primary costimulatory signal is delivered through the activating CD28 receptor upon engagement of its ligands B7-1 or B7-2. In contrast, engagement of the inhibitory CTLA-4 receptor by the same B7-1 or B7-2 ligands results in attenuation of T cell response. Thus, CTLA-4 signals antagonize costimulation mediated by CD28. At high antigen concentrations, CD28 costimulation overrides the CTLA-4 inhibitory effect. Temporal regulation of the CD28 and CTLA-4 expression maintains a balance between activating and inhibitory signals and ensures the development of an effective immune response, while safeguarding against the development of autoimmunity.

[005] Molecular homologues of CD28 and CTLA-4 and their B-7 like ligands have been recently identified. ICOS is a CD28-like costimulatory receptor. PD-1 (Programmed Death 1) is an inhibitory receptor and a counterpart of CTLA-4. This disclosure relates to modulation of immune responses mediated by the PD-1 receptor.

[006] PD-1 is a 50-55 kDa type I transmembrane receptor that was originally identified in a T cell line undergoing activation-induced apoptosis. PD-1 is expressed on T cells, B cells, and macrophages. The ligands for PD-1 are the B7 family members PD-L1 (B7-H1) and PD-L2 (B7-DC).

[007] PD-1 is a member of the immunoglobulin (Ig) superfamily that contains a single Ig V–like domain in its extracellular region. The PD-1 cytoplasmic domain contains two tyrosines, with the most membrane-proximal tyrosine (VAYEEL in mouse PD-1) located within an ITIM (immuno-receptor tyrosine-based inhibitory motif). The presence of an ITIM on PD-1 indicates that this molecule functions to attenuate antigen receptor signaling by recruitment of cytoplasmic phosphatases. Human and murine PD-1 proteins share about 60% amino acid identity with conservation of four potential N-glycosylation sites, and residues that define the Ig-V domain. The ITIM in the cytoplasmic region and the ITIM-like motif surrounding the carboxy-terminal tyrosine (TEYATI in human and mouse) are also conserved between human and murine orthologues.

[008] PD-1 is expressed on activated T cells, B cells, and monocytes. Experimental data implicates the interactions of PD-1 with its ligands in downregulation of central and peripheral immune responses. In particular, proliferation in wild-type T cells but not in PD-1-deficient T cells is inhibited in the presence of PD-L1. Additionally, PD-1-deficient mice exhibit an autoimmune phenotype. PD-1 deficiency in the C57BL/6 mice results in chronic progressive lupus-like glomerulonephritis and arthritis. In Balb/c mice, PD-1 deficiency leads to severe cardiomyopathy due to the presence of heart-tissue-specific self-reacting antibodies.

[009] In general, a need exists to provide safe and effective therapeutic methods for immune disorders such as, for example, autoimmune diseases, inflammatory disorders, allergies, transplant rejection, cancer, immune deficiency, and other immune system-related disorders. Modulation of the immune responses involved in these disorders can be accomplished by manipulation of the PD-1 pathway.

SUMMARY OF THE INVENTION

The present invention provides for an isolated antibody including a VH domain including SEQ ID NO:6, SEQ ID NO:10, SEQ ID NO:14 or SEQ ID NO:47, wherein the antibody specifically binds an epitope within the extracellular domain of human or mouse PD-1.

The present invention also provides for an isolated antibody including a VL domain including SEQ ID NO:8, SEQ ID NO:12, SEQ ID NO:16 or SEQ ID NO:49, wherein the antibody specifically binds an epitope within the extracellular domain of human or mouse PD-1.

The present invention further provides for an isolated antibody including a VH domain or an antigen-binding fragment thereof that includes 3 CDRs and a VL domain or an antigen-binding fragment thereof that includes 3 CDRs, wherein the antibody includes at least 3 VH domain CDRs having sequences of: SEQ ID NO:23, SEQ ID NO:24 and SEQ ID NO:25; or SEQ ID NO:29, SEQ ID NO:30 and SEQ ID NO:31; or SEQ ID NO:35, SEQ ID NO:36 and SEQ ID NO:37; or SEQ ID NO:50, SEQ ID NO:51 and SEQ ID NO:52; or at least 3 VL domain CDRs having sequences of: SEQ ID NO:26, SEQ ID NO:27 and SEQ ID NO:28; or SEQ ID NO:32, SEQ ID NO:33 and SEQ ID NO:34; or SEQ ID NO:38, SEQ ID NO:39 and SEQ ID NO:40; or SEQ ID NO:53, SEQ ID NO:54, and SEQ ID NO:55; and wherein the antibody specifically binds an epitope within the extracellular domain of human or mouse PD-1.

The present invention additionally provides for an antibody produced by steps including providing a nucleic acid encoding 3 VL domain CDRs

having sequences of: SEQ ID NO:26, SEQ ID NO:27 and SEQ ID NO:28; or SEQ ID NO:32, SEQ ID NO:33 and SEQ ID NO:34; or SEQ ID NO:38, SEQ ID NO:39 and SEQ ID NO:40; or SEQ ID NO:53, SEQ ID NO:54, and SEQ ID NO:55, combining a repertoire of nucleic acids encoding 3 VH domain CDRs with the nucleic acid encoding the 3 VL domain CDRs, so as to provide a product repertoire of nucleic acids encoding the 3 VL domain CDRs and the repertoire of 3 VH domain CDRs, expressing the nucleic acids of the product repertoire, selecting an antigen-binding fragment comprising a variable domain that specifically binds to PD-1 and that is expressed from the nucleic acids of the product repertoire, and producing an antibody comprising the antigen-binding fragment.

NOTICE UNDER REGISTRAR'S CIRCULAR NO. 23(P) OF 5 APRIL 1992:

Inasmuch as the invention is defined in the appended claims, it will be apparent that the portions of the present specification which fall outside the scope of the claims do not relate directly to the invention. This Notice is not meant to disclaim any legitimate rights to which the Patentee is legally entitled, especially any rights in accordance with Section 49 of the Israel Patents Law.

ADDITIONAL ASPECTS OF THE APPLICATION

[010] The present disclosure provides antibodies that can act as agonists and/or antagonists of PD-1, thereby modulating immune responses regulated by PD-1. The disclosure further provides anti-PD-1 antibodies that comprise novel antigen-binding fragments. Anti-PD-1 antibodies of the invention are capable of (a) specifically binding to PD-1, including human PD-1; (b) blocking PD-1 interactions with its natural ligand(s); or (c) performing both functions. Furthermore, the antibodies may possess immunomodulatory properties, i.e., they may be effective in modulating the PD-1-associated downregulation of immune responses. Depending on the method of use and the desired effect, the antibodies may be used to either enhance or inhibit immune responses.

[011] Nonlimiting illustrative embodiments of the antibodies are referred to as PD1-17, PD1-28, PD1-33, PD1-35, and PD1-F2. Other embodiments comprise a V_H and/or V_L domain of the Fv fragment of PD1-17, PD1-28, PD1-33, PD1-35, or PD1-F2. Further embodiments comprise one or more complementarity determining regions (CDRs) of any of these V_H and V_L domains. Other embodiments comprise an H3 fragment of the V_H domain of PD1-17, PD1-28, PD1-33, PD1-35, or PD1-F2.

[012] The disclosure also provides compositions comprising PD-1 antibodies, and their use in methods of modulating immune response, including methods of treating humans or animals. In particular embodiments, anti-PD-1 antibodies are used to treat or prevent immune disorders by virtue of increasing or reducing the T cell response mediated by TcR/CD28.

Disorders susceptible to treatment with compositions of the invention include but are not limited to rheumatoid arthritis, multiple sclerosis, inflammatory bowel disease, Crohn's disease, systemic lupus erythematosis, type I diabetes, transplant rejection, graft-versus-host disease, hyperproliferative immune disorders, cancer, and infectious diseases.

[013] Additionally, anti-PD-1 antibodies may be used diagnostically to detect PD-1 or its fragments in a biological sample. The amount of PD-1 detected may be correlated with the expression level of PD-1, which, in turn, is correlated with the activation status of immune cells (e.g., activated T cells, B cells, and monocytes) in the subject.

[014] The disclosure also provides isolated nucleic acids, which comprise a sequence encoding a V_H or V_L domain from the Fv fragment of PD1-17, PD1-28, PD1-33, PD1-35, or PD1-F2. Also provided are isolated nucleic acids, which comprise a sequence encoding one or more CDRs from any of the presently disclosed V_H and V_L domains. The disclosure also provides vectors and host cells comprising such nucleic acids.

[015] The disclosure further provides a method of producing new V_H and V_L domains and/or functional antibodies comprising all or a portion of such domains derived from the V_H or V_L domains of PD1-17, PD1-28, PD1-33, PD1-35, or PD1-F2.

[016] Additional aspects of the disclosure will be set forth in part in the description which follows, and in part will be obvious from the description, or may be learned by practicing the invention. The invention is set forth and particularly pointed out in the appended claims, and the present disclosure should not be construed as limiting the scope of the claims in any way. The following detailed description includes exemplary representations of various embodiments of the invention, which are not restrictive of the invention, as claimed. The accompanying figures constitute a part of this specification and, together with the description, serve only to illustrate various embodiments and not limit the invention. Citation of references is not an admission that these references are prior art to the invention.

BRIEF DESCRIPTION OF THE FIGURES

[017] Figures 1A and 1B show reactivity of scFv antibodies with human PD-1 as determined by phage ELISA.

[018] Figures 2A-2C show reactivity of IgG-converted antibodies with human or mouse PD-1 as determined by ELISA.

[019] Figure 3 shows results of an ELISA demonstrating that selected PD-1 antibodies inhibit binding of PD-L1 to PD-1.

[020] Figure 4 shows results of an ELISA demonstrating that immunomodulatory PD-1 antibodies bind to distinct sites on PD-1 as determined by cross-blocking ELISA assays.

[021] Figure 5 shows results of T-cell proliferation assays demonstrating that co-engagement by TcR and anti-PD-1 antibody PD1-17 or PD-L1.Fc reduces proliferation. Co-engagement by TcR and anti-PD-1 J110 has no effect on proliferation.

[022] Figure 6 demonstrates enhanced proliferation of primary T cells by PD1-17 in a soluble form.

DETAILED DESCRIPTION

Definitions

[023] The term "antibody," as used in this disclosure, refers to an immunoglobulin or a fragment or a derivative thereof, and encompasses any polypeptide comprising an antigen-binding site, regardless whether it is produced in vitro or in vivo. The term includes, but is not limited to, polyclonal, monoclonal, monospecific, polyspecific, non-specific, humanized, single-chain, chimeric, synthetic, recombinant, hybrid, mutated, and grafted antibodies. Unless otherwise modified by the term "intact," as in "intact antibodies," for the purposes of this disclosure, the term "antibody" also includes antibody fragments such as Fab, $F(ab')_2$, Fv, scFv, Fd, dAb, and other antibody fragments that retain antigen-binding function, i.e., the ability to

bind PD-1 specifically. Typically, such fragments would comprise an antigen-binding domain.

[024] The terms "antigen-binding domain," "antigen-binding fragment," and "binding fragment" refer to a part of an antibody molecule that comprises amino acids responsible for the specific binding between the antibody and the antigen. In instances, where an antigen is large, the antigen-binding domain may only bind to a part of the antigen. A portion of the antigen molecule that is responsible for specific interactions with the antigen-binding domain is referred to as "epitope" or "antigenic determinant."

[025] An antigen-binding domain typically comprises an antibody light chain variable region (V_L) and an antibody heavy chain variable region (V_H), however, it does not necessarily have to comprise both. For example, a so-called Fd antibody fragment consists only of a V_H domain, but still retains some antigen-binding function of the intact antibody.

[026] The term "repertoire" refers to a genetically diverse collection of nucleotides derived wholly or partially from sequences that encode expressed immunoglobulins. The sequences are generated by in vivo rearrangement of, e.g., V, D, and J segments for H chains and, e.g., V and J segment for L chains. Alternatively, the sequences may be generated from a cell line by in vitro stimulation, in response to which the rearrangement occurs. Alternatively, part or all of the sequences may be obtained by combining, e.g., unrearranged V segments with D and J segments, by nucleotide synthesis, randomised mutagenesis, and other methods, e.g., as disclosed in U.S. Patent No. 5,565,332.

[027] The terms "specific interaction" and "specific binding" refer to two molecules forming a complex that is relatively stable under physiologic conditions. Specific binding is characterized by a high affinity and a low to moderate capacity as distinguished from nonspecific binding which usually has a low affinity with a moderate to high capacity. Typically, binding is considered specific when the affinity constant K_A is higher than $10^6\ M^{-1}$, or more preferably higher than $10^8\ M^{-1}$. If necessary, non-specific binding can be

reduced without substantially affecting specific binding by varying the binding conditions. The appropriate binding conditions such as concentration of antibodies, ionic strength of the solution, temperature, time allowed for binding, concentration of a blocking agent (e.g., serum albumin, milk casein), etc., may be optimized by a skilled artisan using routine techniques. Illustrative conditions are set forth in Examples 1, 2, 4, 6, and 7.

[028] The phrase "substantially as set out" means that the relevant CDR, V_H, or V_L domain of the invention will be either identical to or have only insubstantial differences in the specified regions (e.g., a CDR), the sequence of which is set out. Insubstantial differences include minor amino acid changes, such as substitutions of 1 or 2 out of any 5 amino acids in the sequence of a specified region.

[029] The term "PD-1 activity" refers to one or more immunoregulatory activities associated with PD-1. For example, PD-1 is a negative regulator of the TcR/CD28-mediated immune response. Procedures for assessing the PD-1 activity in vivo and in vitro are described in Examples 8, 9, and 10.

[030] The terms "modulate," "immunomodulatory," and their cognates refer to a reduction or an increase in the activity of PD-1 associated with downregulation of T cell responses due to its interaction with an anti-PD-1 antibody, wherein the reduction or increase is relative to the activity of PD-1 in the absence of the same antibody. A reduction or an increase in activity is preferably at least about 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, or more. When PD-1 activity is reduced, the terms "modulatory" and "modulate" are interchangeable with the terms "inhibitory" and "inhibit." When PD-1 activity is increased, the terms "modulatory" and "modulate" are interchangeable with the terms "activating" and "activate." The activity of PD-1 can be determined quantitatively using T cell proliferation assays as described in Examples 8 and 9.

[031] The terms "treatment" and "therapeutic method" refer to both therapeutic treatment and prophylactic/preventative measures. Those in need of treatment may include individuals already having a particular medical

disorder as well as those who may ultimately acquire the disorder (i.e., those needing preventative measures).

[032] The term "effective amount" refers to a dosage or amount that is sufficient to reduce the activity of PD-1 to result in amelioration of symptoms in a patient or to achieve a desired biological outcome, e.g., increased cytolytic activity of T cells, induction of immune tolerance, reduction or increase of the PD-1 activity associated with the negative regulation of T-cell mediated immune response, etc.

[033] The term "isolated" refers to a molecule that is substantially free of its natural environment. For instance, an isolated protein is substantially free of cellular material or other proteins from the cell or tissue source from which it is derived. The term "isolated" also refers to preparations where the isolated protein is sufficiently pure to be administered as a pharmaceutical composition, or at least 70-80% (w/w) pure, more preferably, at least 80-90% (w/w) pure, even more preferably, 90-95% pure; and, most preferably, at least 95%, 96%, 97%, 98%, 99%, or 100% (w/w) pure.

Anti-PD-1 Antibodies

[034] The disclosure provides anti-PD-1 antibodies that comprise novel antigen-binding fragments.

[035] In general, antibodies can be made, for example, using traditional hybridoma techniques (Kohler and Milstein (1975) Nature, 256: 495-499), recombinant DNA methods (U.S. Patent No. 4,816,567), or phage display performed with antibody libraries (Clackson et al. (1991) Nature, 352: 624-628; Marks et al. (1991) J. Mol. Biol., 222: 581-597). For other antibody production techniques, see also Antibodies: A Laboratory Manual, eds. Harlow et al., Cold Spring Harbor Laboratory, 1988. The invention is not limited to any particular source, species of origin, method of production.

[036] Intact antibodies, also known as immunoglobulins, are typically tetrameric glycosylated proteins composed of two light (L) chains of approximately 25 kDa each and two heavy (H) chains of approximately 50 kDa each. Two types of light chain, designated as the λ chain and the κ

chain, are found in antibodies. Depending on the amino acid sequence of the constant domain of heavy chains, immunoglobulins can be assigned to five major classes: A, D, E, G, and M, and several of these may be further divided into subclasses (isotypes), e.g., IgG_1, IgG_2, IgG_3, IgG_4, IgA_1, and IgA_2.

[037] The subunit structures and three-dimensional configurations of different classes of immunoglobulins are well known in the art. For a review of antibody structure, see Harlow et al., supra. Briefly, each light chain is composed of an N-terminal variable domain (V_L) and a constant domain (C_L). Each heavy chain is composed of an N-terminal variable domain (V_H), three or four constant domains (C_H), and a hinge region. The C_H domain most proximal to V_H is designated as C_H1. The V_H and V_L domains consist of four regions of relatively conserved sequence called framework regions (FR1, FR2, FR3, and FR4), which form a scaffold for three regions of hypervariable sequence called complementarity determining regions (CDRs). The CDRs contain most of the residues responsible for specific interactions with the antigen. The three CDRs are referred to as CDR1, CDR2, and CDR3. CDR constituents on the heavy chain are referred to as H1, H2, and H3, while CDR constituents on the light chain are referred to as L1, L2, and L3, accordingly. CDR3 and, particularly H3, are the greatest source of molecular diversity within the antigen-binding domain. H3, for example, can be as short as two amino acid residues or greater than 26.

[038] The Fab fragment (Fragment antigen-binding) consists of the V_H-C_H1 and V_L-C_L domains covalently linked by a disulfide bond between the constant regions. To overcome the tendency of non-covalently linked V_H and V_L domains in the Fv to dissociate when co-expressed in a host cell, a so-called single chain (sc) Fv fragment (scFv) can be constructed. In a scFv, a flexible and adequately long polypeptide links either the C-terminus of the V_H to the N-terminus of the V_L or the C-terminus of the V_L to the N-terminus of the V_H. Most commonly, a 15-residue $(Gly_4Ser)_3$ peptide is used as a linker but other linkers are also known in the art.

[039] Antibody diversity is a result of combinatorial assembly of multiple germline genes encoding variable regions and a variety of somatic events. The somatic events include recombination of variable gene segments with diversity (D) and joining (J) gene segments to make a complete V_H region and the recombination of variable and joining gene segments to make a complete V_L region. The recombination process itself is imprecise, resulting in the loss or addition of amino acids at the V(D)J junctions. These mechanisms of diversity occur in the developing B cell prior to antigen exposure. After antigenic stimulation, the expressed antibody genes in B cells undergo somatic mutation.

[040] Based on the estimated number of germline gene segments, the random recombination of these segments, and random V_H-V_L pairing, up to 1.6×10^7 different antibodies could be produced (Fundamental Immunology, 3rd ed., ed. Paul, Raven Press, New York, NY, 1993). When other processes which contribute to antibody diversity (such as somatic mutation) are taken into account, it is thought that upwards of 1×10^{10} different antibodies could be potentially generated (Immunoglobulin Genes, 2nd ed., eds. Jonio et al., Academic Press, San Diego, CA, 1995). Because of the many processes involved in antibody diversity, it is highly unlikely that independently generated antibodies will have identical or even substantially similar amino acid sequences in the CDRs.

[041] The disclosure provides novel CDRs derived from human immunoglobulin gene libraries. The structure for carrying a CDR will generally be an antibody heavy or light chain or a portion thereof, in which the CDR is located at a location corresponding to the CDR of naturally occurring V_H and V_L. The structures and locations of immunoglobulin variable domains may be determined, for example, as described in Kabat et al., Sequences of Proteins of Immunological Interest, No. 91-3242, National Institutes of Health Publications, Bethesda, MD, 1991.

[042] DNA and amino acid sequences of anti-PD-1 antibodies, their scFv fragment, V_H and V_L domains, and CDRs are set forth in the Sequence

Listing and are enumerated as listed in Table 1. Particular nonlimiting illustrative embodiments of the antibodies are referred to as PD1-17, PD1-28, PD1-33, PD1-35, and PD1-F2. The positions for each CDR within the V_H and V_L domains of the illustrative embodiments are listed in Tables 2 and 3.

Table 1: DNA and Amino Acid (AA) Sequences of V_H and V_L Domains and CDRs

Sequence	PD1-17	PD1-28	PD1-33	PD1-35	PD1-F2
V_H DNA	SEQ ID NO:1	SEQ ID NO:5	SEQ ID NO:9	SEQ ID NO:13	SEQ ID NO:46
V_H AA	SEQ ID NO:2	SEQ ID NO:6	SEQ ID NO:10	SEQ ID NO:14	SEQ ID NO:47
V_L DNA	SEQ ID NO:3	SEQ ID NO:7	SEQ ID NO:11	SEQ ID NO:15	SEQ ID NO:48
V_L AA	SEQ ID NO:4	SEQ ID NO:8	SEQ ID NO:12	SEQ ID NO:16	SEQ ID NO:49
H1 AA	SEQ ID NO:17	SEQ ID NO:23	SEQ ID NO:29	SEQ ID NO:35	SEQ ID NO:50
H2 AA	SEQ ID NO:18	SEQ ID NO:24	SEQ ID NO:30	SEQ ID NO:36	SEQ ID NO:51
H3 AA	SEQ ID NO:19	SEQ ID NO:25	SEQ ID NO:31	SEQ ID NO:37	SEQ ID NO:52
L1 AA	SEQ ID NO:20	SEQ ID NO:26	SEQ ID NO:32	SEQ ID NO:38	SEQ ID NO:53
L2 AA	SEQ ID NO:21	SEQ ID NO:27	SEQ ID NO:33	SEQ ID NO:39	SEQ ID NO:54
L3 AA	SEQ ID NO:22	SEQ ID NO:28	SEQ ID NO:34	SEQ ID NO:40	SEQ ID NO:55

Table 2: Positions of Heavy Chain CDRs

CDR	PD1-17 SEQ ID NO:2	PD1-28 SEQ ID NO:6	PD1-33 SEQ ID NO:10	PD1-35 SEQ ID NO:14	PD1-F2 SEQ ID NO:47
H1	31-42	31-35	31-35	31-37	34-42
H2	57-72	50-66	50-66	52-67	57-73
H3	105-117	99-108	99-108	100-116	106-114

Table 3: Positions of Light Chain CDRs

CDR	PD1-17 SEQ ID NO:4	PD1-28 SEQ ID NO:8	PD1-33 SEQ ID NO:12	PD1-35 SEQ ID NO:16	PD1-F2 SEQ ID NO:49
L1	23-35	23-33	23-36	23-35	28-35
L2	51-57	49-55	52-58	51-57	54-61
L3	92-100	88-98	91-102	90-100	94-101

[043] Anti-PD-1 antibodies may optionally comprise antibody constant regions or parts thereof. For example, a V_L domain may have attached, at its C terminus, antibody light chain constant domains including human Cκ or Cλ chains. Similarly, a specific antigen-binding domain based on a V_H domain may have attached all or part of an immunoglobulin heavy chain derived from any antibody isotope, e.g., IgG, IgA, IgE, and IgM and any of the isotope sub-classes, which include but are not limited to, IgG_1 and IgG_4. In the exemplary embodiments, PD1-17, PD1-28, PD1-33, and PD1-35, antibodies comprise C-terminal fragments of heavy and light chains of human $IgG_{1\lambda}$, while PD1-F2 comprises C-terminal fragments of heavy and light chains of human $IgG_{1\kappa}$. The DNA and amino acid sequences for the C-terminal fragment of are well known in the art (see, e.g, Kabat et al., Sequences of Proteins of Immunological Interest, No. 91-3242, National Institutes of Health Publications, Bethesda, MD, 1991). Nonlimiting exemplary sequences are set forth in Table 4.

Table 4

C-Terminal Region	DNA	Amino acid
IgG1 heavy chain	SEQ ID NO:44	SEQ ID NO:45
λ light chain	SEQ ID NO:42	SEQ ID NO:43
κ light chain	SEQ ID NO:57	SEQ ID NO:58

[044] Certain embodiments comprise a V_H and/or V_L domain of an Fv fragment from PD1-17, PD1-28, PD1-33, PD1-35, and PD1-F2. Further embodiments comprise at least one CDR of any of these V_H and V_L domains. Antibodies, comprising at least one of the CDR sequences set out in SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, SEQ ID NO:8, SEQ ID NO:10, SEQ ID NO:12, SEQ ID NO:14, SEQ ID NOs:16-40, SEQ ID NO:47, or SEQ ID NO:49 are encompassed within the scope of this invention. An embodiment, for example, comprises an H3 fragment of the V_H domain of antibodies chosen from at least one of PD1-17, PD1-28, PD1-33, PD1-35, and PD1-F2.

[045] In certain embodiments, the V_H and/or V_L domains may be germlined, i.e., the framework regions (FRs) of these domains are mutated using conventional molecular biology techniques to match those produced by the germline cells. In other embodiments, the framework sequences remain diverged from the consensus germline sequences.

[046] In certain embodiments, the antibodies specifically bind an epitope within the extracellular domain of human PD-1. The predicted extracellular domain consists of a sequence from about amino acid 21 to about amino acid 170 of SEQ ID NO:41 (Swissport Accession No. Q15116). In certain other embodiments, the antibodies specifically bind an epitope within the extracellular domain of mouse PD-1, with an affinity of more than 10^7 M^{-1}, and preferably more than 10^8 M^{-1}. The amino acid sequence of mouse PD-1 is set out in SEQ ID NO:56 (Accession No. NM_008798) and is as a whole about 60% identical to its human counterpart. In further embodiments, antibodies of the invention bind to the PD-L-binding domain of PD-1.

[047] It is contemplated that antibodies of the invention may also bind with other proteins, including, for example, recombinant proteins comprising all or a portion of the PD-1 extracellular domain.

[048] One of ordinary skill in the art will recognize that the antibodies of this invention may be used to detect, measure, and inhibit proteins that differ somewhat from PD-1. The antibodies are expected to retain the

specificity of binding so long as the target protein comprises a sequence which is at least about 60%, 70%, 80%, 90%, 95%, or more identical to any sequence of at least 100, 80, 60, 40, or 20 of contiguous amino acids in the sequence set forth SEQ ID NO:41. The percent identity is determined by standard alignment algorithms such as, for example, Basic Local Alignment Tool (BLAST) described in Altshul et al. (1990) J. Mol. Biol., 215: 403-410, the algorithm of Needleman et al. (1970) J. Mol. Biol., 48: 444-453, or the algorithm of Meyers et al. (1988) Comput. Appl. Biosci., 4: 11-17.

[049] In addition to the sequence homology analyses, epitope mapping (see, e.g., Epitope Mapping Protocols, ed. Morris, Humana Press, 1996) and secondary and tertiary structure analyses can be carried out to identify specific 3D structures assumed by the disclosed antibodies and their complexes with antigens. Such methods include, but are not limited to, X-ray crystallography (Engstom (1974) Biochem. Exp. Biol., 11:7-13) and computer modeling of virtual representations of the presently disclosed antibodies (Fletterick et al. (1986) Computer Graphics and Molecular Modeling, in Current Communications in Molecular Biology, Cold Spring Harbor Laboratory, Cold Spring Harbor, NY).

Derivatives

[050] This disclosure also provides a method for obtaining an antibody specific for PD-1. CDRs in such antibodies are not limited to the specific sequences of V_H and V_L identified in Table 1 and may include variants of these sequences that retain the ability to specifically bind PD-1. Such variants may be derived from the sequences listed in Table 1 by a skilled artisan using techniques well known in the art. For example, amino acid substitutions, deletions, or additions, can be made in the FRs and/or in the CDRs. While changes in the FRs are usually designed to improve stability and immunogenicity of the antibody, changes in the CDRs are typically designed to increase affinity of the antibody for its target. Variants of FRs also include naturally occurring immunoglobulin allotypes. Such affinity-increasing changes may be determined empirically by routine

techniques that involve altering the CDR and testing the affinity antibody for its target. For example, conservative amino acid substitutions can made within any one of the disclosed CDRs. Various alterations can be made according to the methods described in Antibody Engineering, 2nd ed., Oxford University Press, ed. Borrebaeck, 1995. These include but are not limited to nucleotide sequences that are altered by the substitution of different codons that encode a functionally equivalent amino acid residue within the sequence, thus producing a "silent" change. For example, the nonpolar amino acids include alanine, leucine, isoleucine, valine, proline, phenylalanine, tryptophan, and methionine. The polar neutral amino acids include glycine, serine, threonine, cysteine, tyrosine, asparagine, and glutamine. The positively charged (basic) amino acids include arginine, lysine, and histidine. The negatively charged (acidic) amino acids include aspartic acid and glutamic acid. Substitutes for an amino acid within the sequence may be selected from other members of the class to which the amino acid belongs (see Table 5). Furthermore, any native residue in the polypeptide may also be substituted with alanine (see, e.g., MacLennan et al. (1998) Acta Physiol. Scand. Suppl. 643:55-67; Sasaki et al. (1998) Adv. Biophys. 35:1-24).

[051] Derivatives and analogs of antibodies of the invention can be produced by various techniques well known in the art, including recombinant and synthetic methods (Maniatis (1990) Molecular Cloning, A Laboratory Manual, 2nd ed., Cold Spring Harbor Laboratory, Cold Spring Harbor, NY, and Bodansky et al. (1995) The Practice of Peptide Synthesis, 2nd ed., Spring Verlag, Berlin, Germany).

Table 5

Original Residues	Exemplary Substitutions	Typical Substitutions
Ala (A)	Val, Leu, Ile	Val
Arg (R)	Lys, Gln, Asn	Lys
Asn (N)	Gln	Gln
Asp (D)	Glu	Glu
Cys (C)	Ser, Ala	Ser
Gln (Q)	Asn	Asn
Gly (G)	Pro, Ala	Ala
His (H)	Asn, Gln, Lys, Arg	Arg
Ile (I)	Leu, Val, Met, Ala, Phe, Norleucine	Leu
Leu (L)	Norleucine, Ile, Val, Met, Ala, Phe	Ile
Lys (K)	Arg, 1,4-Diamino-butyric Acid, Gln, Asn	Arg
Met (M)	Leu, Phe, Ile	Leu
Phe (F)	Leu, Val, Ile, Ala, Tyr	Leu
Pro (P)	Ala	Gly
Ser (S)	Thr, Ala, Cys	Thr
Thr (T)	Ser	Ser
Trp (W)	Tyr, Phe	Tyr
Tyr (Y)	Trp, Phe, Thr, Ser	Phe
Val (V)	Ile, Met, Leu, Phe, Ala, Norleucine	Leu

[052] In one embodiment, a method for making a V_H domain which is an amino acid sequence variant of a V_H domain of the invention comprises a step of adding, deleting, substituting, or inserting one or more amino acids in the amino acid sequence of the presently disclosed V_H domain, optionally combining the V_H domain thus provided with one or more V_L domains, and testing the V_H domain or V_H/V_L combination or combinations for a specific binding to PD-1 or and, optionally, testing the ability of such antigen-binding domain to modulate PD-1 activity. The V_L domain may have an amino acid sequence that is identical or is substantially as set out according to Table 1.

[053] An analogous method can be employed in which one or more sequence variants of a V_L domain disclosed herein are combined with one or more V_H domains.

[054] A further aspect of the disclosure provides a method of preparing antigen-binding fragment that specifically binds with PD-1. The method comprises:

(a) providing a starting repertoire of nucleic acids encoding a V_H domain that either includes a CDR3 to be replaced or lacks a CDR3 encoding region;

(b) combining the repertoire with a donor nucleic acid encoding an amino acid sequence substantially as set out herein for a V_H CDR3 (i.e., H3) such that the donor nucleic acid is inserted into the CDR3 region in the repertoire, so as to provide a product repertoire of nucleic acids encoding a V_H domain;

(c) expressing the nucleic acids of the product repertoire;

(d) selecting a binding fragment specific for PD-1; and

(e) recovering the specific binding fragment or nucleic acid encoding it.

[055] Again, an analogous method may be employed in which a V_L CDR3 (i.e., L3) of the invention is combined with a repertoire of nucleic acids encoding a V_L domain, which either include a CDR3 to be replaced or lack a CDR3 encoding region. The donor nucleic acid may be selected from nucleic acids encoding an amino acid sequence substantially as set out in SEQ ID NO:17-40 or SEQ ID NO:50-55.

[056] A sequence encoding a CDR of the invention (e.g., CDR3) may be introduced into a repertoire of variable domains lacking the respective CDR (e.g., CDR3), using recombinant DNA technology, for example, using methodology described by Marks et al. (Bio/Technology (1992) 10: 779-783). In particular, consensus primers directed at or adjacent to the 5' end of the variable domain area can be used in conjunction with consensus primers to the third framework region of human V_H genes to provide a repertoire of V_H

variable domains lacking a CDR3. The repertoire may be combined with a CDR3 of a particular antibody. Using analogous techniques, the CDR3-derived sequences may be shuffled with repertoires of V_H or V_L domains lacking a CDR3, and the shuffled complete V_H or V_L domains combined with a cognate V_L or V_H domain to make the PD-1-specific antibodies of the invention. The repertoire may then be displayed in a suitable host system such as the phage display system such as described in WO92/01047 so that suitable antigen-binding fragments can be selected.

[057] Analogous shuffling or combinatorial techniques are also disclosed by Stemmer (Nature (1994) 370: 389-391), who describes the technique in relation to a β-lactamase gene but observes that the approach may be used for the generation of antibodies.

[058] In further embodiments, one may generate novel V_H or V_L regions carrying one or more sequences derived from the sequences disclosed herein using random mutagenesis of one or more selected V_H and/or V_L genes. One such technique, error-prone PCR, is described by Gram et al. (Proc. Nat. Acad. Sci. U.S.A. (1992) 89: 3576-3580).

[059] Another method that may be used is to direct mutagenesis to CDRs of V_H or V_L genes. Such techniques are disclosed by Barbas et al. (Proc. Nat. Acad. Sci. U.S.A. (1994) 91: 3809-3813) and Schier et al. (J. Mol. Biol. (1996) 263: 551-567).

[060] Similarly, one or more, or all three CDRs may be grafted into a repertoire of V_H or V_L domains, which are then screened for an antigen-binding fragment specific for PD-1.

[061] A portion of an immunoglobulin variable domain will comprise at least one of the CDRs substantially as set out herein and, optionally, intervening framework regions from the scF_V fragments as set out herein. The portion may include at least about 50% of either or both of FR1 and FR4, the 50% being the C-terminal 50% of FR1 and the N-terminal 50% of FR4. Additional residues at the N-terminal or C-terminal end of the substantial part of the variable domain may be those not normally associated with naturally

occurring variable domain regions. For example, construction of antibodies by recombinant DNA techniques may result in the introduction of N- or C-terminal residues encoded by linkers introduced to facilitate cloning or other manipulation steps. Other manipulation steps include the introduction of linkers to join variable domains to further protein sequences including immunoglobulin heavy chain constant regions, other variable domains (for example, in the production of diabodies), or proteinaceous labels as discussed in further detail below.

[062] Although the embodiments illustrated in the Examples comprise a "matching" pair of V_H and V_L domains, a skilled artisan will recognize that alternative embodiments may comprise antigen-binding fragments containing only a single CDR from either V_L or V_H domain. Either one of the single chain specific binding domains can be used to screen for complementary domains capable of forming a two-domain specific antigen-binding fragment capable of, for example, binding to PD-1. The screening may be accomplished by phage display screening methods using the so-called hierarchical dual combinatorial approach disclosed in WO92/01047, in which an individual colony containing either an H or L chain clone is used to infect a complete library of clones encoding the other chain (L or H) and the resulting two-chain specific binding domain is selected in accordance with phage display techniques as described.

[063] Anti-PD1 antibodies described herein can be linked to another functional molecule, e.g., another peptide or protein (albumin, another antibody, etc.), toxin, radioisotope, cytotoxic or cytostatic agents. For example, the antibodies can be linked by chemical cross-linking or by recombinant methods. The antibodies may also be linked to one of a variety of nonproteinaceous polymers, e.g., polyethylene glycol, polypropylene glycol, or polyoxyalkylenes, in the manner set forth in U.S. Patent Nos. 4,640,835; 4,496,689; 4,301,144; 4,670,417; 4,791,192; or 4,179,337. The antibodies can be chemically modified by covalent conjugation to a polymer, for example, to increase their circulating half-life. Exemplary polymers and methods to

attach them are also shown in U.S. Patent Nos. 4,766,106; 4,179,337; 4,495,285; and 4,609,546.

[064] The disclosed antibodies may also be altered to have a glycosylation pattern that differs from the native pattern. For example, one or more carbohydrate moieties can be deleted and/or one or more glycosylation sites added to the original antibody. Addition of glycosylation sites to the presently disclosed antibodies may be accomplished by altering the amino acid sequence to contain glycosylation site consensus sequences known in the art. Another means of increasing the number of carbohydrate moieties on the antibodies is by chemical or enzymatic coupling of glycosides to the amino acid residues of the antibody. Such methods are described in WO 87/05330 and in Aplin et al. (1981) CRC Crit. Rev. Biochem., 22: 259-306. Removal of any carbohydrate moieties from the antibodies may be accomplished chemically or enzymatically, for example, as described by Hakimuddin et al. (1987) Arch. Biochem. Biophys., 259: 52; and Edge et al. (1981) Anal. Biochem., 118: 131 and by Thotakura et al. (1987) Meth. Enzymol., 138: 350. The antibodies may also be tagged with a detectable, or functional, label. Detectable labels include radiolabels such as ^{131}I or ^{99}Tc, which may also be attached to antibodies using conventional chemistry. Detectable labels also include enzyme labels such as horseradish peroxidase or alkaline phosphatase. Detectable labels further include chemical moieties such as biotin, which may be detected via binding to a specific cognate detectable moiety, e.g., labeled avidin.

[065] Antibodies, in which CDR sequences differ only insubstantially from those set out in SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, SEQ ID NO:8, SEQ ID NO:10, SEQ ID NO:12, SEQ ID NO:14, SEQ ID NOs:16-40, SEQ ID NO:47, or SEQ ID NO:49 are encompassed within the scope of this invention. Typically, an amino acid is substituted by a related amino acid having similar charge, hydrophobic, or stereochemical characteristics. Such substitutions would be within the ordinary skills of an artisan. Unlike in CDRs, more substantial changes can be made in FRs without adversely affecting the

binding properties of an antibody. Changes to FRs include, but are not limited to, humanizing a non-human derived or engineering certain framework residues that are important for antigen contact or for stabilizing the binding site, e.g., changing the class or subclass of the constant region, changing specific amino acid residues which might alter the effector function such as Fc receptor binding, e.g., as described in U.S. Patent Nos. 5,624,821 and 5,648,260 and Lund et al. (1991) J. Immun. 147: 2657-2662 and Morgan et al. (1995) Immunology 86: 319-324, or changing the species from which the constant region is derived.

[066] One of skill in the art will appreciate that the modifications described above are not all-exhaustive, and that many other modifications would obvious to a skilled artisan in light of the teachings of the present disclosure.

Nucleic Acids, Cloning and Expression Systems

[067] The present disclosure further provides isolated nucleic acids encoding the disclosed antibodies. The nucleic acids may comprise DNA or RNA and may be wholly or partially synthetic or recombinant. Reference to a nucleotide sequence as set out herein encompasses a DNA molecule with the specified sequence, and encompasses a RNA molecule with the specified sequence in which U is substituted for T, unless context requires otherwise.

[068] The nucleic acids provided herein comprise a coding sequence for a CDR, a V_H domain, and/or a V_L domain disclosed herein.

[069] The present disclosure also provides constructs in the form of plasmids, vectors, phagemids, transcription or expression cassettes which comprise at least one nucleic acid encoding a CDR, a V_H domain, and/or a V_L domain disclosed here.

[070] The disclosure further provides a host cell which comprises one or more constructs as above.

[071] Also provided are nucleic acids encoding any CDR (H1, H2, H3, L1, L2, or L3), V_H or V_L domain, as well as methods of making of the encoded products. The method comprises expressing the encoded product from the

encoding nucleic acid. Expression may be achieved by culturing under appropriate conditions recombinant host cells containing the nucleic acid. Following production by expression a V_H or V_L domain, or specific binding member may be isolated and/or purified using any suitable technique, then used as appropriate.

[072] Antigen-binding fragments, V_H and/or V_L domains, and encoding nucleic acid molecules and vectors may be isolated and/or purified from their natural environment, in substantially pure or homogeneous form, or, in the case of nucleic acid, free or substantially free of nucleic acid or genes of origin other than the sequence encoding a polypeptide with the required function.

[073] Systems for cloning and expression of a polypeptide in a variety of different host cells are well known in the art. For cells suitable for producing antibodies, see Gene Expression Systems, Academic Press, eds. Fernandez et al., 1999. Briefly, suitable host cells include bacteria, plant cells, mammalian cells, and yeast and baculovirus systems. Mammalian cell lines available in the art for expression of a heterologous polypeptide include Chinese hamster ovary cells, HeLa cells, baby hamster kidney cells, NS0 mouse myeloma cells, and many others. A common bacterial host is *E. coli*. Any protein expression system compatible with the invention may be used to produce the disclosed antibodies. Suitable expression systems include transgenic animals described in Gene Expression Systems, Academic Press, eds. Fernandez et al., 1999.

[074] Suitable vectors can be chosen or constructed, so that they contain appropriate regulatory sequences, including promoter sequences, terminator sequences, polyadenylation sequences, enhancer sequences, marker genes and other sequences as appropriate. Vectors may be plasmids or viral, e.g., phage, or phagemid, as appropriate. For further details see, for example, Sambrook et al., Molecular Cloning: A Laboratory Manual, 2nd ed., Cold Spring Harbor Laboratory Press, 1989. Many known techniques and protocols for manipulation of nucleic acid, for example, in preparation of nucleic acid constructs, mutagenesis, sequencing, introduction of DNA into

cells and gene expression, and analysis of proteins, are described in detail in Current Protocols in Molecular Biology, 2nd Edition, eds. Ausubel et al., John Wiley & Sons, 1992.

[075] A further aspect of the disclosure provides a host cell comprising a nucleic acid as disclosed here. A still further aspect provides a method comprising introducing such nucleic acid into a host cell. The introduction may employ any available technique. For eukaryotic cells, suitable techniques may include calcium phosphate transfection, DEAE-Dextran, electroporation, liposome-mediated transfection and transduction using retrovirus or other virus, e.g., vaccinia or, for insect cells, baculovirus. For bacterial cells, suitable techniques may include calcium chloride transformation, electroporation and transfection using bacteriophage. The introduction of the nucleic acid into the cells may be followed by causing or allowing expression from the nucleic acid, e.g., by culturing host cells under conditions for expression of the gene.

Methods of Use

[076] The disclosed anti-PD-1 antibodies are capable of modulating the PD-1-associated downregulation of the immune responses. In particular embodiments, the immune response is TcR/CD28-mediated. The disclosed antibodies can act as either agonists or antagonists of PD-1, depending on the method of their use. The antibodies can be used to prevent, diagnose, or treat medical disorders in mammals, especially, in humans. Antibodies of the invention can also be used for isolating PD-1 or PD-1-expressing cells. Furthermore, the antibodies can be used to treat a subject at risk of or susceptible to a disorder or having a disorder associated with aberrant PD-1 expression or function.

[077] Antibodies of the invention can be used in methods for induction of tolerance to a specific antigen (e.g., a therapeutic protein). In one embodiment, tolerance is induced against a specific antigen by co-administration of antigen and an anti-PD-1 antibody of the invention. For example, patients that received Factor VIII frequently generate antibodies to

this protein; co-administration of an anti-PD-1 antibody of the invention in combination with recombinant Factor VIII is expected to result in the downregulation of immune responses to this clotting factor.

[078] Antibodies of the invention can be used in circumstances where a reduction in the level of immune response may be desirable, for example, in certain types of allergy or allergic reactions (e.g., by inhibition of IgE production), autoimmune diseases (e.g., rheumatoid arthritis, type I diabetes mellitus, multiple sclerosis, inflammatory bowel disease, Crohn's disease, and systemic lupus erythematosis), tissue, skin and organ transplant rejection, and graft-versus-host disease (GVHD).

[079] When diminished immune response is desirable, the anti-PD-1 antibodies of the invention may be used as agonists to PD-1 in order to enhance the PD-1-associated attenuation of the immune response. In these embodiments, co-presentation and physical proximity between positive (i.e., mediated by an antigen receptor, e.g., TcR or BcR) and negative (i.e., PD-1) signals are required. The preferred distance is less than or comparable to the size of a naturally occurring antigen-presenting cell, i.e., less than about 100 μm; more preferably, less than about 50 μm; and most preferably, less than about 20 μm.

[080] In some embodiments, the positive (activating) and the negative (inhibiting) signals are provided by a ligand or antibodies immobilized on solid support matrix, or a carrier. In various embodiments, the solid support matrix may be composed of polymer such as activated agarose, dextran, cellulose, polyvinylidene fluoride (PVDF). Alternatively, the solid support matrix may be based on silica or plastic polymers, e.g., as nylon, dacron, polystyrene, polyacrylates, polyvinyls, teflons, etc.

[081] The matrix can be implanted into the spleen of a patient. Alternatively, the matrix may be used for the ex vivo incubation of T cells obtained from a patient, which are then separated and implanted back into the patient. The matrix may also be made from a biodegradable material such polyglycolic acid, polyhydroxyalkanoate, collagen, or gelatin so that they can

be injected into the patient's peritoneal cavity, and dissolve after some time following the injection. The carrier can be shaped to mimic a cell (e.g., bead or microsphere).

[082] In some embodiments, the positive signal is delivered by a T-cell-activating anti-CD3 antibody, which binds TcR. Activating anti-CD3 antibodies are known in the art (see, for example, U.S. Patent Nos. 6,405,696 and 5,316,763). The ratio between the activating TcR signal and negative PD-1 signal is determined experimentally using conventional procedures known in the art or as described in Examples 8, 9, and 10.

[083] Under certain circumstances, it may be desirable to elicit or enhance a patient's immune response in order to treat an immune disorder or cancer. The disorders being treated or prevented by the disclosed methods include but are not limited to infections with microbes (e.g. bacteria), viruses (e.g., systemic viral infections such as influenza, viral skin diseases such as herpes or shingles), or parasites; and cancer (e.g., melanoma and prostate cancers).

[084] Stimulation of T cell activation with anti-PD-1 antibodies enhances T-T cell responses. In such cases, antibodies act as antagonists of PD-1. Thus, in some embodiments, the antibodies can be used to inhibit or reduce the downregulatory activity associated with PD-1, i.e., the activity associated with downregulation of TcR/CD28-mediated immune response. In these embodiments, the antibodies are not coupled to a positive signal such as the TcR-mediated stimulation, e.g., the antibodies are in their soluble, support-unbound, form. As demonstrated in the Examples, a blockade of PD-1/PD-L interaction with antagonizing anti-PD-1 antibodies leads to enhanced T cell proliferative responses, consistent with a downregulatory role for the PD-1 pathway in T-T interactions. In various embodiments, the antibodies inhibit binding of PD-L to PD-1 with an IC_{50} of less than 10 nM, and more preferably less then 5 nM, and most preferably less than 1 nM. Inhibition of PD-L binding can be measured as described in Example 6 or using techniques known in the art.

[085] The antibodies or antibody compositions of the present invention are administered in therapeutically effective amounts. Generally, a therapeutically effective amount may vary with the subject's age, condition, and sex, as well as the severity of the medical condition of the subject. A therapeutically effective amount of antibody ranges from about 0.001 to about 30 mg/kg body weight, preferably from about 0.01 to about 25 mg/kg body weight, from about 0.1 to about 20 mg/kg body weight, or from about 1 to about 10 mg/kg. The dosage may be adjusted, as necessary, to suit observed effects of the treatment. The appropriate dose is chosen based on clinical indications by a treating physician.

[086] The antibodies may given as a bolus dose, to maximize the circulating levels of antibodies for the greatest length of time after the dose. Continuous infusion may also be used after the bolus dose.

[087] Immune cells (e.g., activated T cells, B cells, or monocytes) can also be isolated from a patient and incubated ex vivo with antibodies of the invention. In some embodiments, immune responses can be inhibited by removing immune cells from a subject, contacting the immune cells in vitro with an anti-PD-1 antibody of the invention concomitantly with activation of the immune cells (e.g., by antibodies to the TcR and/or BcR antigen receptor). In such embodiments, the anti-PD-1 antibody should be used in a multivalent form such that PD-1 molecules on the surface of an immune cell become "crosslinked" upon binding to such antibodies. For example, the anti-PD-1 antibodies can be bound to solid support, such as beads, or crosslinked via a secondary antibody. The immune cells may be then isolated using methods known in the art and reimplanted into the patient.

[088] In another aspect, the antibodies of the invention can be used as a targeting agent for delivery of another therapeutic or a cytotoxic agent (e.g., a toxin) to a cell expressing PD-1. The method includes administering an anti-PD-1 antibody coupled to a therapeutic or a cytotoxic agent or under conditions that allow binding of the antibody to PD-1.

[089] The antibodies of the invention may also be used to detect the presence of PD-1 in biological samples. The amount of PD-1 detected may be correlated with the expression level of PD-1, which, in turn, is correlated with the activation status of immune cells (e.g., activated T cells, B cells, and monocytes) in the subject.

[090] Detection methods that employ antibodies are well known in the art and include, for example, ELISA, radioimmunoassay, immunoblot, Western blot, immunofluorescence, immunoprecipitation. The antibodies may be provided in a diagnostic kit that incorporates one or more of these techniques to detect PD-1. Such a kit may contain other components, packaging, instructions, or other material to aid the detection of the protein.

[091] Where the antibodies are intended for diagnostic purposes, it may be desirable to modify them, for example, with a ligand group (such as biotin) or a detectable marker group (such as a fluorescent group, a radioisotope or an enzyme). If desired, the antibodies of the invention may be labeled using conventional techniques. Suitable detectable labels include, for example, fluorophores, chromophores, radioactive atoms, electron-dense reagents, enzymes, and ligands having specific binding partners. Enzymes are typically detected by their activity. For example, horseradish peroxidase can be detected by its ability to convert tetramethylbenzidine (TMB) to a blue pigment, quantifiable with a spectrophotometer. For detection, suitable binding partners include, but are not limited to, biotin and avidin or streptavidin, IgG and protein A, and the numerous receptor-ligand couples known in the art. Other permutations and possibilities will be readily apparent to those of ordinary skill in the art, and are considered as equivalents within the scope of the instant invention.

[092] Antibodies of the invention can be used in screening methods to identify inhibitors of the PD-1 pathway effective as therapeutics. In such a screening assay, a first binding mixture is formed by combining PD-1 and an antibody of the invention; and the amount of binding in the first binding mixture (M_0) is measured. A second binding mixture is also formed by

combining PD-1, the antibody, and the compound or agent to be screened, and the amount of binding in the second binding mixture (M_1) is measured. A compound to be tested may be another anti-PD-1 antibody, as illustrated in the Examples. The amounts of binding in the first and second binding mixtures are then compared, for example, by calculating the M_1/M_0 ratio. The compound or agent is considered to be capable of modulating a PD-1-associated downregulation of immune responses if a decrease in binding in the second binding mixture as compared to the first binding mixture is observed. The formulation and optimization of binding mixtures is within the level of skill in the art, such binding mixtures may also contain buffers and salts necessary to enhance or to optimize binding, and additional control assays may be included in the screening assay of the invention. Compounds found to reduce the PD-1-antibody binding by at least about 10% (i.e., $M_1/M_0<0.9$), preferably greater than about 30% may thus be identified and then, if desired, secondarily screened for the capacity to ameliorate a disorder in other assays or animal models as described below. The strength of the binding between PD-1 and an antibody can be measured using, for example, an enzyme-linked immunoadsorption assay (ELISA), radio-immunoassay (RIA), surface plasmon resonance-based technology (e.g., Biacore), all of which are techniques well known in the art.

[093] The compound may then be tested in vitro as described in the Examples or in an animal model (see, generally, Immunologic Defects in Laboratory Animals, eds. Gershwin et al., Plenum Press, 1981), for example, such as the following: the SWR X NZB (SNF1) transgenic mouse model (Uner et al. (1998) J. Autoimmune. 11(3): 233-240), the KRN transgenic mouse (K/BxN) model (Ji et al. (1999) Immunol. Rev. 169: 139); NZB X NZW (B/W) mice, a model for SLE (Riemekasten et al. (2001) Arthritis Rheum., 44(10): 2435-2445); experimental autoimmune encephalitis (EAE) in mouse, a model for multiple sclerosis (Tuohy et al. (1988) J. Immunol. 141: 1126-1130, Sobel et al. (1984) J. Immunol. 132: 2393-2401, and Traugott, Cell Immunol. (1989)

119: 114-129); the NOD mouse model of diabetes (Baxter et al. (1991) Autoimmunity, 9(1): 61-67), etc.).

[094] Preliminary doses as, for example, determined according to animal tests, and the scaling of dosages for human administration is performed according to art-accepted practices. Toxicity and therapeutic efficacy can be determined by standard pharmaceutical procedures in cell cultures or experimental animals. The data obtained from the cell culture assays or animal studies can be used in formulating a range of dosage for use in humans. Therapeutically effective dosages achieved in one animal model can be converted for use in another animal, including humans, using conversion factors known in the art (see, e.g., Freireich et al. (1966) Cancer Chemother. Reports, 50(4): 219-244).

Pharmaceutical Compositions and Methods of Administration

[095] The disclosure provides compositions comprising anti-PD-1 antibodies. Such compositions may be suitable for pharmaceutical use and administration to patients. The compositions typically comprise one or more antibodies of the present invention and a pharmaceutically acceptable excipient. The phrase "pharmaceutically acceptable excipient" includes any and all solvents, dispersion media, coatings, antibacterial agents and antifungal agents, isotonic agents, and absorption delaying agents, and the like, that are compatible with pharmaceutical administration. The use of such media and agents for pharmaceutically active substances is well known in the art. The compositions may also contain other active compounds providing supplemental, additional, or enhanced therapeutic functions. The pharmaceutical compositions may also be included in a container, pack, or dispenser together with instructions for administration.

[096] A pharmaceutical composition of the invention is formulated to be compatible with its intended route of administration. Methods to accomplish the administration are known to those of ordinary skill in the art. The administration may, for example, be intravenous, intraperitoneal, intramuscular, intracavity, subcutaneous or transdermal. It may also be

possible to obtain compositions which may be topically or orally administered, or which may be capable of transmission across mucous membranes.

[097] Solutions or suspensions used for intradermal or subcutaneous application typically include one or more of the following components: a sterile diluent such as water for injection, saline solution, fixed oils, polyethylene glycols, glycerin, propylene glycol, or other synthetic solvents; antibacterial agents such as benzyl alcohol or methyl parabens; antioxidants such as ascorbic acid or sodium bisulfite; chelating agents such as ethylenediaminetetraacetic acid; buffers such as acetates, citrates or phosphates; and agents for the adjustment of tonicity such as sodium chloride or dextrose. The pH can be adjusted with acids or bases, such as hydrochloric acid or sodium hydroxide. Such preparations may be enclosed in ampoules, disposable syringes or multiple dose vials made of glass or plastic.

[098] Pharmaceutical compositions suitable for injection include sterile aqueous solutions or dispersions and sterile powders for the extemporaneous preparation of sterile injectable solutions or dispersion. For intravenous administration, suitable carriers include physiological saline, bacteriostatic water, Cremophor EL (BASF, Parsippany, NJ) or phosphate buffered saline (PBS). In all cases, the composition must be sterile and should be fluid to the extent that easy syringability exists. It should be stable under the conditions of manufacture and storage and must be preserved against the contaminating action of microorganisms such as bacteria and fungi. Prevention of the action of microorganisms can be achieved by various antibacterial and antifungal agents, for example, parabens, chlorobutanol, phenol, ascorbic acid, thimerosal, and the like. In many cases, it will be preferable to include isotonic agents, for example, sugars, polyalcohols such as mannitol, sorbitol, and sodium chloride in the composition. The carrier can be a solvent or dispersion medium containing, for example, water, ethanol, polyol (for example, glycerol, propylene glycol, and liquid polyetheylene glycol, and the like), and suitable mixtures thereof. The proper fluidity can be maintained, for

example, by the use of a coating such as lecithin, by the maintenance of the required particle size in the case of dispersion and/or by the use of surfactants. Prolonged absorption of the injectable compositions can be brought about by including in the composition an agent which delays absorption, for example, aluminum monostearate, and gelatin.

[099] Oral compositions generally include an inert diluent or an edible carrier. They can be enclosed in gelatin capsules or compressed into tablets. For oral administration, the antibodies can be combined with excipients and used in the form of tablets, troches, or capsules. Pharmaceutically compatible binding agents, and/or adjuvant materials can be included as part of the composition. The tablets, pills, capsules, troches, and the like can contain any of the following ingredients, or compounds of a similar nature; a binder such as microcrystalline cellulose, gum tragacanth or gelatin; an excipient such as starch or lactose, a disintegrating agent such as alginic acid, Primogel, or corn starch; a lubricant such as magnesium stearate or Sterotes; a glidant such as colloidal silicon dioxide; a sweetening agent such as sucrose or saccharin; or a flavoring agent such as peppermint, methyl salicylate, or orange flavoring.

[0100] Systemic administration can also be by transmucosal or transdermal means. For transmucosal or transdermal administration, penetrants appropriate to the barrier to be permeated are used in the formulation. Such penetrants are generally known in the art, and include, for example, detergents, bile salts, and fusidic acid derivatives. Transmucosal administration may be accomplished, for example, through the use of lozenges, nasal sprays, inhalers, or suppositories. For example, in case of antibodies that comprise the Fc portion, compositions may be capable of transmission across mucous membranes in intestine, mouth, or lungs (e.g., via the FcRn receptor-mediated pathway as described in U.S. Patent No. 6,030,613). For transdermal administration, the active compounds may be formulated into ointments, salves, gels, or creams as generally known in the art. For administration by inhalation, the antibodies may be delivered in the

form of an aerosol spray from pressured container or dispenser, which contains a suitable propellant, e.g., a gas such as carbon dioxide, or a nebulizer.

[0101] In certain embodiments, the presently disclosed antibodies are prepared with carriers that will protect the compound against rapid elimination from the body, such as a controlled release formulation, including implants and microencapsulated delivery systems. Biodegradable, biocompatible polymers can be used, such as ethylene vinyl acetate, polyanhydrides, polyglycolic acid, collagen, polyorthoesters, and polylactic acid. Methods for preparation of such formulations will be apparent to those skilled in the art. Liposomal suspensions containing the presently disclosed antibodies can also be used as pharmaceutically acceptable carriers. These can be prepared according to methods known to those skilled in the art, for example, as described in U.S. Patent No. 4,522,811.

[0102] It may be advantageous to formulate oral or parenteral compositions in a dosage unit form for ease of administration and uniformity of dosage. The term "dosage unit form" as used herein refers to physically discrete units suited as unitary dosages for the subject to be treated; each unit containing a predetermined quantity of active compound calculated to produce the desired therapeutic effect in association with the required pharmaceutical carrier.

[0103] Toxicity and therapeutic efficacy of the composition of the invention can be determined by standard pharmaceutical procedures in cell cultures or experimental animals, e.g., for determining the LD_{50} (the dose lethal to 50% of the population) and the ED_{50} (the dose therapeutically effective in 50% of the population). The dose ratio between toxic and therapeutic effects is the therapeutic index and it can be expressed as the ratio LD_{50}/ED_{50}. Compositions that exhibit large therapeutic indices are preferred.

[0104] For any composition used in the present invention, the therapeutically effective dose can be estimated initially from cell culture

assays. Examples of suitable bioassays include DNA replication assays, cytokine release assays, transcription-based assays, PD-1/PD-L1 binding assays, creatine kinase assays, assays based on the differentiation of pre-adipocytes, assays based on glucose uptake in adipocytes, immunological assays other assays as, for example, described in the Examples. The data obtained from the cell culture assays and animal studies can be used in formulating a range of dosage for use in humans. A dose may be formulated in animal models to achieve a circulating plasma concentration range that includes the IC_{50} (i.e., the concentration of the antibody which achieves a half-maximal inhibition of symptoms). Circulating levels in plasma may be measured, for example, by high performance liquid chromatography. The effects of any particular dosage can be monitored by a suitable bioassay. The dosage lies preferably within a range of circulating concentrations with little or no toxicity. The dosage may vary depending upon the dosage form employed and the route of administration utilized.

[0105] The following Examples do not in any way limit the scope of the invention. One of ordinary skill in the art will recognize the numerous modifications and variations that may be performed without altering the spirit or scope of the present invention. Such modifications and variations are encompassed within the scope of the invention. The entire contents of all references, patents, and published patent applications cited throughout this application are herein incorporated by reference.

EXAMPLES

Example 1: Selection of PD-1 Binding ScFv's

[0106] An scFv phagemid library, which is an expanded version of the $1.38x10^{10}$ library described by Vaughan et al. (Nature Biotech. (1996) 14: 309-314) was used to select antibodies specific for human PD-1. Soluble PD-1 fusion protein (at 20 µg/ml in phosphate buffered saline (PBS)) or control fusion protein (at 50 µg/ml in PBS) was coated onto wells of a microtiter plate overnight at 4°C. Wells were washed in PBS and blocked for

1 hour at 37°C in MPBS (3% milk powder in PBS). Purified phage (10^{12} transducing units (tu)) was blocked for 1 hour in a final volume of 100 µl of 3% MPBS. Blocked phage was added to blocked control fusion protein wells and incubated for 1 hour. The blocked and deselected phage were then transferred to the blocked wells coated with the PD-1 fusion protein and were incubated for an additional hour. Wells were washed 5 times with PBST (PBS containing 0.1% v/v Tween 20), then 5 times with PBS. Bound phage particles were eluted and used to infect 10 ml exponentially growing *E. coli* TG1. Infected cells were grown in 2TY broth for 1 hour at 37°C, then spread onto 2TYAG plates and incubated overnight at 30°C. Colonies were scraped off the plates into 10 ml 2TY broth and 15% glycerol added for storage at –70°C.

[0107] Glycerol stock cultures from the first round of panning selection were superinfected with helper phage and rescued to give scFv antibody-expressing phage particles for the second round of panning. A total of two rounds of panning were carried out in this way for isolation of PD1-17, except in the second round of panning 20 µg/ml of control protein were used for deselection. Clones PD1-28, PD1-33, and PD1-35 were selected following three rounds of selection. Deselection in the second and third rounds was carried out using 10 µg/ml control fusion protein.

[0108] Antibodies to murine PD-1 were selected by soluble selection using biotinylated murine PD-1 fusion protein at a final concentration of 100 nM. An scFv phagemid library, as described above, was used. Purified scFv phage (10^{12} tu) in 1 ml 3% MPBS were blocked for 30 minutes, then biotinylated antigen was added and incubated at room temperature for 1 hour. Phage/antigen was added to 250 µl of Dynal M280 Streptavidin magnetic beads that had been blocked for 1 hour at 37°C in 1 ml of 3% MPBS and incubated for a further 15 minutes at room temperature. Beads were captured using a magnetic rack and washed 4 times in 1 ml of 3% MPBS/ 0.1% (v/v) Tween 20 followed by 3 washes in PBS. After the last PBS wash, beads were resuspended in 100 µl PBS and used to infect 5 ml exponentially growing *E.*

coli TG-1 cells. Infected cells were incubated for 1 hour at 37°C (30 minutes stationary, 30 minutes shaking at 250 rpm), then spread on 2TYAG plates and incubated overnight at 30°C. Output colonies were scraped off the plates and phage rescued as described above. A second round of soluble selection was carried out as described above.

Example 2: Specificity of Antibodies for PD-1 by a Phage ELISA

[0109] To determine the specificity of antibodies for PD-1, a phage ELISA was performed against PD-1 fusion protein and control proteins. Individual *E. coli* colonies from selection outputs were picked into 96 well plates containing 100 µl of 2TYAG medium per well. M13K07 helper phage was added to a multiplicity of infection (moi) of 10 to the exponentially growing culture and the plates incubated an additional 1 hour at 37°C. Plates were centrifuged in a benchtop centrifuge at 2000 rpm for 10 minutes. The supernatant was removed and cell pellets were resuspended in 100 µl 2TYAK and incubated at 30°C overnight with shaking. The next day, plates were centrifuged at 2000 rpm for 10 minutes and phage-containing supernatant from each well was transferred to a fresh 96 well plate. Phage samples were blocked in a final concentration of 3% MPBS prior to ELISA.

[0110] Human or mouse PD-1 fusion protein and control fusion and non-fusion proteins were coated overnight at 4°C onto 96-well microtiter plates at 0.5-2.5 µg/ml in PBS. After coating, the solutions were removed from the wells, and the plates blocked for 1 hour in 3% MPBS. Plates were rinsed with PBS and then 50 µl of pre-blocked phage were added to each well. The plates were incubated for 1 hour and then washed 3 times with PBST followed by 3 washes with PBS. To each well, 50 µl of a 1:5000 dilution of anti-M13-HRP conjugate (Pharmacia, Peapack, NJ) was added, and the plates incubated for 40-60 minutes. Each plate was washed three times with PBST then 3 times with PBS. Fifty µl of TMB substrate was added to each well, and the samples were incubated until color development. The reaction was stopped by the addition of 25 µl of 0.5 M H_2SO_4. The signal

generated was measured by reading the absorbance at 450 nm using a microtiter plate reader. Clones showing specific binding to PD-1 fusion protein but not to control fusion proteins were thus identified and confirmed.

[0111] Specificity data for the PD1-17 scFv is shown in Figure 1A. Reactivity of PD1-28, PD1-33, and PD1-35 scFv's with human PD-1 is shown in Figure 1B (an IgG_1 control did not bind PD-1).

Example 3: Identification of Antibody Clones

[0112] PD-1-binding scFv *E. coli* clones were streaked out onto 2TYAG plates and incubated overnight at 30°C. Colonies from these plates were sequenced using pCANTAB6 vector sequence oligos to amplify the V_H and V_L regions from the scFv clone. Unique PD-1 binding clones were assayed for neutralization of PD-L1 binding to PD-1 as described in Example 4. Sequence differences between scFv and IgG formats are due to changes introduced by PCR primers during the conversion from scFv to IgG.

Example 4: Biochemical Binding Inhibition Assay and Screen

[0113] ScFv production was induced by addition of 1 mM IPTG to exponentially growing cultures and incubation overnight at 30°C. Crude scFv-containing periplasmic extracts were obtained by subjecting the bacterial pellets from the overnight induction to osmotic shock. Pellets were resuspended in 20% (w/v) sucrose, 50mM Tris-HCl, pH 7.5, 1 mM EDTA and cooled on ice for 30 minutes. Cellular debris was removed by centrifugation, and the scFv was purified by chromatography and buffer-exchanged into PBS. Purified scFv's (PD1-17, PD1-28, PD1-33, and PD1-35) were tested for the ability to inhibit the binding of biotinylated human PD-L1 fusion protein to human PD-1 fusion protein immobilized on plastic in a 96 well microtiter plate assay. Binding of biotinylated PD-L1 fusion protein was detected with AMDEX-alkaline phosphatase, and the signal generated was measured by reading the absorbance at 405 nm using a microtiter plate reader. Data was expressed as a percentage of the total binding and a titration of scFv

concentrations was tested to establish clone potency as calculated IC_{50} values. Clone potency data for the scFv and IgG antibodies is shown in Table 5.

[0114] PD1-F2 scFv was produced and purified as described above. Cells expressing murine PD-1 were added at 10^5 cells/well in a final volume of 100 µl to a poly-D-lysine-coated 96 well microtiter plate. Cells were centrifuged and washed twice in PBS, then blocked with 300 µl 1% BSA in PBS for 1 hour at room temperature. Blocked cells were washed three times in PBST, prior to addition of 25 µl/well of assay buffer (0.05% BSA, 0.05% Tween 20 in Dulbecco's PBS) or sample, followed by 25 µl of biotinylated murine PD-L1 fusion protein at 300 ng/ml. Binding of biotinylated PD-L1 fusion protein was detected with Amdex alkaline phosphatase and signals read as described above. Potencies of PD1-F2 scFv and IgG are shown in Table 6.

Table 6: Potency of Anti-PD-1 ScFv and IgG Antibodies

Clone	ScFv IC_{50} (nM)	IgG IC_{50} (nM)
PD1-17	726	2.5
PD1-28	560	1.4
PD1-33	74	1.8
PD1-35	85	2.3
PD1-F2	28	1.0

Example 5: Conversion of ScFv to IgG

[0115] Heavy and light chain V regions from scFv clones were amplified by PCR using clone-specific primers. PCR products were digested with appropriate restriction enzymes and subcloned into vectors containing human IgG_1 heavy chain constant domain (Takahashi et al. (1982) Cell 29, 671) or vectors containing human lambda or kappa light chain constant domains (Hieter et al. (1982) Nature 294, 536). Based on the germlines of the

V_H and V_L segments, it was determined whether kappa or lambda light chain constant domains were used for conversion (Table 7).

Table 7: Germlines of V_H and V_L Regions of PD-1 Antibody Clones

Clone	V_H germline	V_L germline
PD1-17	DP-70	DPL-8
PD1-28	DP-14	DPL-23
PD1-33	DP-7	DPL-11
PD1-35	DP-65	DPL-2
PD1-F2	DP-47	L12 (κ)

[0116] The insertion of V region domains into plasmids was verified by sequencing of plasmid DNA from individual *E. coli* colonies. Plasmids were prepared from *E. coli* cultures by standard techniques and heavy and light chain constructs cotransfected into eukaryotic cells using standard techniques. Secreted IgG was purified using Protein A Sepharose (Pharmacia) and buffer-exchanged into PBS.

[0117] The binding affinity of the anti-mouse PD1 antibody PD1-F2 was determined with a Surface Plasmon Resonance (SPR) system (BIAcore 3000) (Biacore, Piscataway, NJ) using murine PD-1 fusion immobilized on a CM5 sensor chip. The concentration of PD1-F2 in the flow cell ranged from 7.81 to 125 nM, while the concentration of the anti-mouse PD1 antibody J43 (eBioscience, San Diego, CA) ranged from 25 nM to 500 nM. The equilibrium constant K_D for PD1-F2 is $6.7x10^{-9}$ M (K_A=$1.5x10^8$ M^{-1}), whereas K_D for J43 is $3.8x10^{-7}$ M (K_A=$2.6x10^6$ M^{-1}).

[0118] The ability of anti-PD-1 IgG's to bind human or murine PD-1 was determined as follows. ELISA plates were incubated with 2.5 μg/ml human PD-1/IgG chimera overnight. Plates were washed with PBS/1 % BSA and incubated with serial dilutions of a test antibody for 2 hours at room temperature (RT). After washing, saturating concentrations of HRP-conjugated goat anti-human antibody or HRP-conjugated rabbit

anti-murine antibody were added, and the samples were incubated for 1 hour at RT. Unbound goat and rabbit antibodies were washed using PBS/1 % BSA. The assay was developed using TBM. Results were expressed as OD 405 absorbency values and are presented in Figures 2A-2C. Murine anti-human PD-1 antibody J110 is commercially available (eBioscience, San Diego, CA) and was included for comparison.

Example 6: Selected PD-1 Antibodies Inhibit Binding of PD-L1 to PD-1

[0119] Inhibition assays were performed to assess the ability of the antibodies to block binding of PD-L1 to PD-1. ELISA was performed as described in Example 2 with modifications. After incubation with a primary, anti-PD-1 antibody for 2 hours at RT, a fixed concentration (1 µg/ml) of biotin-conjugated PD-L1-Ig was added, and the samples were further incubated for 1 hour at RT. After washing, saturating concentrations of avidin-HRP were added, and incubated for 1 hour at RT. Unbound avidin-HRP was washed using PBS/1% BSA. The assay was developed using TMB.

[0120] Results were compared to those obtained with J110 as shown in Figure 3. Anti-human PD-1 antibodies J110 and PD1-30 did not inhibit the binding of PD-L1 to PD-1. Anti-human antibodies PD1-17, PD1-28, PD1-33, and PD1-35 and anti-mouse antibody PD1-F2 block PD-1/PD-L1 interaction.

Example 7: PD-1 Antibodies Recognize Distinct Sites on PD-1

[0121] Inhibition assays were performed to map sites recognize by the various human anti-human PD-1 antibodies. ELISA was performed as described in Example 6 with minor modifications. After incubation with primary antibody for 2 hours at RT, a fixed concentration (0.25 µg/ml) of biotin-conjugated anti-PD-1 antibody J110 was added, and the samples were further incubated for 1 hour at RT. After washing, saturating concentrations of avidin-HRP were added, and incubated for 1 hour at RT. Unbound

avidin-HRP was washed using PBS/1% BSA. The assay was developed using TMB.

[0122] As shown in Figure 4, binding of anti-human PD-1 antibodies (J110, J116, PD1-17, PD1-28, PD1-33, and PD1-35) defines at least two distinct sites on PD-1. Cross-blocking results show that J110 and J116, bind to identical or overlapping sites while PD1-17, 28, 33, and 35 bind to another distinct site. Binding of J116 or J110 to PD-1 blocks the binding of J110. In contrast, binding of PD1-17, PD1-28, PD1-33, and PD1-35 do not block binding of J110. This suggests that the tested anti-PD-1 antibodies bind to at least two distinct epitopes: one recognized by J110 and J116, and the other one recognized by PD1-17, PD1-28, PD1-33, and PD1-35.

Example 8: PD-1 Engagement Results in Decreased T Cell Responses

[0123] CD4+ T cells (5×10^4 cells/well) were stimulated with tosyl-beads (Dynal, Great Neck, NY) coated with anti-hCD3 +/- PD-L1-Fc or anti-PD-1 (PD1-17 or J110). Concentration of fusion protein or antibody titer was as indicated in the X-axis of Figure 5. After 72 hours, proliferation was determined by ^{3}H-thymidine incorporation. Incorporated radioactivity was determined using a LKB 1205 plate reader.

[0124] As shown in Figure 5, PD-1 engagement by anti-PD-1 antibody PD1-17 or PD-L1.Fc caused a decrease in T cell proliferation. Thus, PD1-17 can mimic PD-1 ligands and delivered an inhibitory signal. As discussed below (Example 9), this inhibitory signal results in decreased T cell proliferation and IL-2 production. Antibodies PD1- 28, PD1-33, and PD1-35 have the same effect as PD1-17. The effect is dose-dependent, as activation of cells in the presence of increasing concentrations of PD1-17 or PD-L1.Fc results in decreased T cell proliferation. The control anti-PD-1 antibodies, J110 (Figure 5) or J116 (data not shown), do not inhibit T cell responses and increasing the concentration of J110 has minimal effect on T cell proliferation. For comparison, values are represented as percentage of the anti-CD3 response. "100%" represents CPMs obtained when cells were activated with

anti-CD3/murine IgG-coated microspheres. Altogether these results indicate that some but not all antibodies that recognize PD-1 can act as agonists of the PD-1 pathway.

[0125] Further experiments were performed to address whether PD-1 downregulation of T cell responses required coordinate TcR/PD-1 engagement on a single (CIS) or a separate (TRANS) cell surfaces. Two sets of microspheres were prepared: one set contained anti-CD3 and PD-L1.Fc (CIS), the other set contained anti-CD3 or PD-L1.Fc (TRANS). Inhibition through PD-1 was only observed under conditions where both PD-1 and TcR were engaged by ligands on the same surface (CIS). At all bead:cell ratios tested, no inhibition was observed in conditions where TCR and PD-1 signals were delivered on separate surfaces (TRANS).

[0126] To rule out steric hindrance in the TRANS experiments, similar assays were set up using anti-CD3 antibody and B7.2.Fc. In these assays, B7 costimulation of T cell responses was observed in both CIS and TRANS conditions. Altogether, these findings demonstrate that PD-1 proximity to TCR is required for the receptor modulatory function on T cell activation. Therefore, to modulate a T cell response, both activating and inhibitory signals must emanate from the same surface whether the surface is that of a cell or a bead.

Example 9: Blockage of PD-1 Engagement by Antibody Results in Enhanced Proliferation

[0127] For assessing effect of soluble anti-PD-1 antibody on proliferation, CD4+ T cells were pre-activated for 48 hours with anti-CD3/anti-CD28-coated beads, harvested, and restimulated with the indicated concentration of PHA plus 10 ng/ml IL-2 in the presence of PD1-17, J110, or control IgG. Each of the antibodies was added at various concentrations at initiation of the culture. Proliferation was measured at 72 hr.

[0128] The results demonstrate that PD1- 17 (Figure 6) and PD1-35 (data not shown) enhanced proliferation of primary T cells. The control

antibody J110 did not enhance in vitro T cell responses. Selected anti-PD1 antibodies, as exemplified by PD1-17 and PD-35, inhibit the interaction of PD-1 with its natural ligands and thereby block delivery of a negative signal. The blockade of the negative signal also results in enhanced proliferation and IL-2 production.

Example 10: Treatment of Disorders

[0129] Modulation of immune response regulated by PD-1 is useful in instances where an immunosuppressive effect or augmentation of immune response is desired. This example describes the use of PD-1 antibodies as PD-1 agonists or antagonists to treat a subject at disease onset or having an established immune disorder or cancer, respectively.

[0130] Subjects at risk for or afflicted with cancer may be in need of immune response augmentation would benefit from treatment with a PD-1 antagonist, such as an anti-PD-1 antibody of the present invention in a soluble form. Most commonly, antibodies are administered in an outpatient setting by weekly administration at about 0.1-10 mg/kg dose by slow intravenous (IV) infusion. The appropriate therapeutically effective dose of an antagonist is selected by a treating clinician and would range approximately from 1 µg/kg to 20 mg/kg, from 1 µg/kg to 10 mg/kg, from 1 µg/kg to 1 mg/kg, from 10 µg/kg to 1 mg/kg, from 10 µg/kg to 100 µg/kg, from 100 µg to 1 mg/kg, and from 500 µg/kg to 5 mg/kg.

[0131] The antibodies are also used to prevent and/or to reduce severity and/or symptoms of diseases or conditions that involve an aberrant or undesirable immune response, such as in autoimmune disorders exemplified below.

[0132] Multiple sclerosis (MS) is a central nervous system disease that is characterized by inflammation and loss of myelin sheaths. In the experimental autoimmune encephalitis (EAE) mouse model for multiple sclerosis (Tuohy et al. (J. Immunol. (1988) 141: 1126-1130), Sobel et al. (J. Immunol. (1984) 132: 2393-2401), and Traugott (Cell Immunol. (1989) 119:

114-129), treatment of mice with a PD-1 agonist prior (and continuously) to EAE induction is expected to prevent or delay the onset of MS.

[0133] Arthritis is a disease characterized by inflammation in the joints. In the collagen induced arthritis (CIA) mouse model for rheumatoid arthritis (Courtenay et al. (Nature (1980) 283: 666-628) and Williams et al. (Immunol. (1995) 84: 433-439)), treatment with a PD-1 agonist is expected to prevent or treat rheumatoid arthritis (RA) or other arthritic diseases.

[0134] Systemic Lupus Erythematosis (SLE) is an autoimmune disease characterized by the presence of autoantibodies. The antibodies and compositions of this invention can be used as PD-1 agonists to inhibit activities of autoreactive T cells and B cells, and prevent or treat SLE or related diseases in NZB X NZW mice (a mouse model for SLE) (Immunologic Defects in Laboratory Animals, Gershwin et al. eds., Plenum Press, 1981) or in humans.

[0135] It is anticipated that PD-1 antibodies of the invention would be administered as PD-1 agonists in ex vivo therapy with a frequency of one per month or less. Treatment duration could range between one month and several years.

[0136] To test the clinical efficacy of antibodies in humans, individuals with melanoma, prostate cancer, RA, SLE, MS, type I diabetes, are identified and randomized to a treatment group. Treatment groups include a placebo group and one to three groups treated with a PD-1 agonist (different doses). Individuals are followed prospectively for one to three years. It is anticipated that individuals receiving treatment would exhibit an improvement.

[0137] The specification is most thoroughly understood in light of the teachings of the references cited within the specification, all of which are hereby incorporated by reference in their entirety. The embodiments within the specification provide an illustration of embodiments of the invention and should not be construed to limit the scope of the invention. The skilled artisan recognizes that many other embodiments are encompassed by the claimed invention and that it is intended that the specification and examples be

considered as exemplary only, with a true scope and spirit of the invention being indicated by the following claims.

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<211> 108
<212> PRT
<213> Homo sapiens

<400> 8

Ser Tyr Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
1 5 10 15

Thr Ala Arg Ile Thr Cys Ser Gly Asp Ala Leu Pro Lys Gln Tyr Ala
20 25 30

Tyr Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Met Val Ile Tyr
35 40 45

Lys Asp Thr Glu Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
50 55 60

Ser Ser Gly Thr Lys Val Thr Leu Thr Ile Ser Gly Val Gln Ala Glu
65 70 75 80

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

Arg Val Phe Gly Gly Gly Thr Lys Val Thr Val Leu
100 105

<210> 9
<211> 357
<212> DNA
<213> Homo sapiens

<400> 9
caggtgcagc tggtgcaatc tggggctgag gtgaagaaac ctggggcctc agtgagggtt 60

tcctgcaagg catctggata caccctcacc agttactata ttcactgggt gcgacaggcc 120

cctggacaag ggcttgagtg gatgggaata atcaacccta gaggtgccac cataagctac 180

gcacagaagt tccagggcag agtcaccatg accagggaca cgtccacgag tacagtctac 240

atggaactga gaaacttgaa atctgaggac acggccctgt attactgtgc tactgcaggc 300

atctatggtt ttgactttga ctactggggc agaggaaccc tggtcaccgt ctcctca 357

<210> 10
<211> 119
<212> PRT
<213> Homo sapiens

<400> 10

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15

Ser Val Arg Val Ser Cys Lys Ala Ser Gly Tyr Thr Leu Thr Ser Tyr
20 25 30

Tyr Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
35 40 45

Gly Ile Ile Asn Pro Arg Gly Ala Thr Ile Ser Tyr Ala Gln Lys Phe
50 55 60

Gln Gly Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser Thr Val Tyr
65 70 75 80

Met Glu Leu Arg Asn Leu Lys Ser Glu Asp Thr Ala Leu Tyr Tyr Cys
85 90 95

Ala Thr Ala Gly Ile Tyr Gly Phe Asp Phe Asp Tyr Trp Gly Arg Gly
100 105 110

Thr Leu Val Thr Val Ser Ser
115

<210> 11
<211> 333
<212> DNA
<213> Homo sapiens

<400> 11
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tcctgcactg gaaccagtaa tgacgttggt ggttataatt atgtctcctg gtaccaacat 120

cacccaggca aagcccccaa actcatcatt tatgatgtca ctaaccggcc ctcaggggtt 180

tctgatcgct tctctggctc caagtctggc aacacggcct ccctgaccat ctctgggctc 240

ctggctgagg acgagggtga ttattactgc agctcatata caattgttac caatttcgag 300

gttcttttcg gcggagggac caagctgacc gtc 333

<210> 12
<211> 111
<212> PRT
<213> Homo sapiens

<400> 12

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

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Gly Tyr
20 25 30

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

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

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

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

Thr Asn Phe Glu Val Leu Phe Gly Gly Gly Thr Lys Leu Thr Val
100 105 110

<210> 13
<211> 381
<212> DNA
<213> Homo sapiens

<400> 13
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acctgcactg tctctggtgg ctccatcagc agtggtgctt attactggag ctggatccgc 120

cagcacccag ggaagggcct ggagtggatt gggtacatct attacaatgg gaacacgtac 180

tacaacccgt ccctcaggag tctagttacc atatcagtag acgcgtctaa gaaccagttc 240

tccctgaagc tgagctctgt gactgccgcg gacacggccg tctattactg tgcgagagcg 300

tctgattacg tttgggggggg ttatcgttat atggatgctt ttgatatctg gggccgggga 360

accctggtca ccgtctcctc a 381

<210> 14
<211> 127
<212> PRT
<213> Homo sapiens

<400> 14

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

Thr Leu Ser Leu Thr Cys Thr Val Ser Gly Gly Ser Ile Ser Ser Gly
20 25 30

Ala Tyr Tyr Trp Ser Trp Ile Arg Gln His Pro Gly Lys Gly Leu Glu
35 40 45

Trp Ile Gly Tyr Ile Tyr Tyr Asn Gly Asn Thr Tyr Tyr Asn Pro Ser
50 55 60

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

Ser Leu Lys Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr
85 90 95

Cys Ala Arg Ala Ser Asp Tyr Val Trp Gly Gly Tyr Arg Tyr Met Asp
100 105 110

Ala Phe Asp Ile Trp Gly Arg Gly Thr Leu Ile Thr Val Ser Ser
115 120 125

<210> 15
<211> 336
<212> DNA
<213> Homo sapiens

<400> 15
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tcttgttctg gaagcaactc caacatcgga agtaattctg taaactggta ccagcagctc 120

ccaggaacgg cccccaaact cctcatctat ggtaataatc agcggccctc aggggtccct 180

gaccgattct ctggctccaa gtctggcacc tcagcctccc tggccatcag tgggctccag 240

tctgagaatg aggctgatta ttactgtgca gcatgggatg acagcctgaa tggtccggta 300

ttcggccgag ggaccaaggt caccgtccta ggtgag 336

<210> 16
<211> 112
<212> PRT
<213> Homo sapiens

<400> 16

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

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

Ser Val Asn Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
35 40 45

Ile Tyr Gly Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser
50 55 60

Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln
65 70 75 80

Ser Glu Asn Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu
85 90 95

Asn Gly Pro Val Phe Gly Arg Gly Thr Lys Val Thr Val Leu Gly Glu
100 105 110

<210> 17
<211> 12
<212> PRT

<213> Homo sapiens

<400> 17

Ser Gly Gly Ser Ile Arg Ser Thr Arg Trp Trp Ser
1 5 10

<210> 18
<211> 16
<212> PRT
<213> Homo sapiens

<400> 18

Glu Ile Tyr His Ser Gly Ser Thr Asn Tyr Asn Pro Ser Leu Lys Ser
1 5 10 15

<210> 19
<211> 13
<212> PRT
<213> Homo sapiens

<400> 19

Gln Asp Tyr Gly Asp Ser Gly Asp Trp Tyr Phe Asp Leu
1 5 10

<210> 20
<211> 13
<212> PRT
<213> Homo sapiens

<400> 20

Thr Arg Ser Ser Gly Ser Ile Ala Ser Asn Ser Val Gln
1 5 10

<210> 21
<211> 7
<212> PRT
<213> Homo sapiens

<400> 21

Glu Asp Asn Gln Arg Pro Ser
1 5

<210> 22
<211> 9
<212> PRT
<213> Homo sapiens

<400> 22

Gln Ser Ser Asp Ser Ser Ala Val Val
1 5

<210> 23
<211> 5
<212> PRT
<213> Homo sapiens

<400> 23

Ser Tyr Gly Ile Ser
1 5

<210> 24
<211> 17
<212> PRT

<213> Homo sapiens

<400> 24

Trp Ile Ser Ala Tyr Asn Gly Asn Thr Asn Tyr Ala Gln Lys Leu Gln
1 5 10 15

Gly

<210> 25
<211> 10
<212> PRT
<213> Homo sapiens

<400> 25

Asp Ala Asp Tyr Ser Ser Gly Ser Gly Tyr
1 5 10

<210> 26
<211> 11
<212> PRT
<213> Homo sapiens

<400> 26

Ser Gly Asp Ala Leu Pro Lys Gln Tyr Ala Tyr
1 5 10

<210> 27
<211> 7
<212> PRT
<213> Homo sapiens

<400> 27

Lys Asp Thr Glu Arg Pro Ser
1 5

<210> 28
<211> 11
<212> PRT
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<400> 28

Gln Ser Ala Asp Asn Ser Ile Thr Tyr Arg Val
1 5 10

<210> 29
<211> 5
<212> PRT
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<400> 29

Ser Tyr Tyr Ile His
1 5

<210> 30
<211> 17
<212> PRT
<213> Homo sapiens

<400> 30

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

Gly

<210> 31
<211> 10
<212> PRT
<213> Homo sapiens

<400> 31

Ala Gly Ile Tyr Gly Phe Asp Phe Asp Tyr
1 5 10

<210> 32
<211> 14
<212> PRT
<213> Homo sapiens

<400> 32

Thr Gly Thr Ser Asn Asp Val Gly Gly Tyr Asn Tyr Val Ser
1 5 10

<210> 33
<211> 7
<212> PRT
<213> Homo sapiens

<400> 33

Asp Val Thr Asn Arg Pro Ser
1 5

<210> 34
<211> 12
<212> PRT

<213> Homo sapiens

<400> 34

Ser Ser Tyr Thr Ile Val Thr Asn Phe Glu Val Leu
1 5 10

<210> 35
<211> 7
<212> PRT
<213> Homo sapiens

<400> 35

Ser Gly Ala Tyr Tyr Trp Ser
1 5

<210> 36
<211> 16
<212> PRT
<213> Homo sapiens

<400> 36

Tyr Ile Tyr Tyr Asn Gly Asn Thr Tyr Tyr Asn Pro Ser Leu Arg Ser
1 5 10 15

<210> 37
<211> 17
<212> PRT
<213> Homo sapiens

<400> 37

Ala Ser Asp Tyr Val Trp Gly Gly Tyr Arg Tyr Met Asp Ala Phe Asp
1 5 10 15

Ile

<210> 38
<211> 13
<212> PRT
<213> Homo sapiens

<400> 38

Ser Gly Ser Asn Ser Asn Ile Gly Ser Asn Ser Val Asn
1 5 10

<210> 39
<211> 7
<212> PRT
<213> Homo sapiens

<400> 39

Gly Asn Asn Gln Arg Pro Ser
1 5

<210> 40
<211> 11
<212> PRT
<213> Homo sapiens

<400> 40

Ala Ala Trp Asp Asp Ser Leu Asn Gly Pro Val
1 5 10

<210> 41
<211> 288
<212> PRT
<213> Homo sapiens

<400> 41

Met Gln Ile Pro Gln Ala Pro Trp Pro Val Val Trp Ala Val Leu Gln
1 5 10 15

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

Asn Pro Pro Thr Phe Phe Pro Ala Leu Leu Val Val Thr Glu Gly Asp
35 40 45

Asn Ala Thr Phe Thr Cys Ser Phe Ser Asn Thr Ser Glu Ser Phe Val
50 55 60

Leu Asn Trp Tyr Arg Met Ser Pro Ser Asn Gln Thr Asp Lys Leu Ala
65 70 75 80

Ala Phe Pro Glu Asp Arg Ser Gln Pro Gly Gln Asp Cys Arg Phe Arg
85 90 95

Val Thr Gln Leu Pro Asn Gly Arg Asp Phe His Met Ser Val Val Arg
100 105 110

Ala Arg Arg Asn Asp Ser Gly Thr Tyr Leu Cys Gly Ala Ile Ser Leu
115 120 125

Ala Pro Lys Ala Gln Ile Lys Glu Ser Leu Arg Ala Glu Leu Arg Val
130 135 140

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Thr Glu Arg Arg Ala Glu Val Pro Thr Ala His Pro Ser Pro Ser Pro
145                 150                 155                 160

Arg Pro Ala Gly Gln Phe Gln Thr Leu Val Val Gly Val Val Gly Gly
                165                 170                 175

Leu Leu Gly Ser Leu Val Leu Leu Val Trp Val Leu Ala Val Ile Cys
            180                 185                 190

Ser Arg Ala Ala Arg Gly Thr Ile Gly Ala Arg Arg Thr Gly Gln Pro
        195                 200                 205

Leu Lys Glu Asp Pro Ser Ala Val Pro Val Phe Ser Val Asp Tyr Gly
    210                 215                 220

Glu Leu Asp Phe Gln Trp Arg Glu Lys Thr Pro Glu Pro Pro Val Pro
225                 230                 235                 240

Cys Val Pro Glu Gln Thr Glu Tyr Ala Thr Ile Val Phe Pro Ser Gly
                245                 250                 255

Met Gly Thr Ser Ser Pro Ala Arg Arg Gly Ser Ala Asp Gly Pro Arg
            260                 265                 270

Ser Ala Gln Pro Leu Arg Pro Glu Asp Gly His Cys Ser Trp Pro Leu
        275                 280                 285
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<210> 42
<211> 320
<212> DNA
<213> Homo sapiens

<400> 42
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ccaacaaggc cacactggtg tgtctcataa gtgacttcta cccgggagcc gtgacagtgg 120

cctggaaggc agatagcagc cccgtcaagg cgggagtgga gaccaccaca ccctccaaac 180

aaagcaacaa caagtacgcg gccagcagct atctgagcct gacgcctgag cagtggaagt 240

cccacagaag ctacagctgc caggtcacgc atgaagggag caccgtggag aagacagtgg 300

cccctacaga atgttcatag 320

<210> 43
<211> 106
<212> PRT
<213> Homo sapiens

<400> 43

Gly Gln Pro Lys Ala Ala Pro Ser Val Thr Leu Phe Pro Pro Ser Ser
1 5 10 15

Glu Glu Leu Gln Ala Asn Lys Ala Thr Leu Val Cys Leu Ile Ser Asp
20 25 30

Phe Tyr Pro Gly Ala Val Thr Val Ala Trp Lys Ala Asp Ser Ser Pro
35 40 45

Val Lys Ala Gly Val Glu Thr Thr Thr Pro Ser Lys Gln Ser Asn Asn
50 55 60

Lys Tyr Ala Ala Ser Ser Tyr Leu Ser Leu Thr Pro Glu Gln Trp Lys
65 70 75 80

Ser His Arg Ser Tyr Ser Cys Gln Val Thr His Glu Gly Ser Thr Val
85 90 95

Glu Lys Thr Val Ala Pro Thr Glu Cys Ser
100 105

<210> 44
<211> 960
<212> DNA
<213> Homo sapiens

<400> 44
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gcacagcggc cctgggctgc ctggtcaagg actacttccc cgaaccggtg acggtgtcgt 120

ggaactcagg cgccctgacc agcggcgtgc acaccttccc ggctgtccta cagtcctcag 180

gactctactc cctcagcagc gtggtgaccg tgccctccag cagcttgggc acccagacct 240

acatctgcaa cgtgaatcac aagcccagca acaccaaggt ggacaagaaa gttgagccca 300

aatcttgtga caaaactcac acatgcccac cgtgcccagc acctgaactc ctgggggggac 360

cgtcagtctt cctcttcccc ccaaaaccca aggacaccct catgatctcc cggacccctg 420

aggtcacatg cgtggtggtg gacgtgagcc acgaagaccc tgaggtcaag ttcaactggt 480

acgtggacgg cgtggaggtg cataatgcca agacaaagcc gcgggaggag cagtacaaca 540

gcacgtaccg tgtggtcagc gtcctcaccg tcctgcacca ggactggctg aatggcaagg 600

agtacaagtg caaggtctcc aacaaagccc tcccagcccc catcgagaaa accatctcca 660

aagccaaagg gcagccccga gaaccacagg tgtacaccct gcccccatcc cgggaggaga 720

tgaccaagaa ccaggtcagc ctgacctgcc tggtcaaagg cttctatccc agcgacatcg 780

ccgtggagtg ggagagcaat gggcagccgg agaacaacta caagaccacg cctcccgtgc 840

tggactccga cggctccttc ttcctctata gcaagctcac cgtggacaag agcaggtggc 900

agcaggggaa cgtcttctca tgctccgtga tgcatgaggc tctgcacaac cactacacgc 960

<210> 45
<211> 330
<212> PRT
<213> Homo sapiens

<400> 45

Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys
1 5 10 15

Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr
20 25 30

Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser
35 40 45

Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser
50 55 60

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr
65 70 75 80

Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys
85 90 95

Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys
100 105 110

Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro
115 120 125

Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys
130 135 140

Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp
145 150 155 160

Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu
165 170 175

Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu
180 185 190

His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn
195 200 205

Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly
210 215 220

Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu
225 230 235 240

Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr
245 250 255

Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn
260 265 270

Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe
275 280 285

Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn
290 295 300

Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr
305 310 315 320

Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
325 330

<210> 46
<211> 366
<212> DNA
<213> Homo sapiens

<400> 46
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tccctgagac tctcctgtgc agcgtctgga ttcaccttta gtagctattg gatgagctgg 120

gtccgccagg ctccagggaa ggggctggag tgggtctcag ctattagtgg tagtggtggt 180

agcacatact acgcagactc cgtgaagggc cggttcacca tctccagaga caattccaag 240

aacacgctgt atctgcaaat gaacagccta agagccgagg acacggccgt atattactgt 300

gcgaaagaga actggggatc gtacttcgat ctctgggggc aagggaccac ggtcaccgtc 360

tcctca 366

<210> 47
<211> 125
<212> PRT
<213> Homo sapiens

<400> 47

Gly Ala His Ser Glu Val Gln Leu Val Gln Ser Gly Gly Gly Val Val
1 5 10 15

Gln Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr
20 25 30

Phe Ser Ser Tyr Trp Cys Asp Arg Met Ser Trp Val Arg Gln Ala Pro
35 40 45

Gly Lys Gly Leu Glu Trp Val Ser Ala Ile Ser Gly Ser Gly Gly Ser
50 55 60

Thr Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp
65 70 75 80

Asn Ser Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu
85 90 95

Asp Thr Ala Val Tyr Tyr Cys Ala Lys Glu Asn Trp Gly Ser Tyr Phe
100 105 110

Asp Leu Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser
115 120 125

<210> 48
<211> 332
<212> DNA
<213> Homo sapiens

<400> 48
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gacagagtca ccatcacttg ccgggccagt cagggtatta gtagctggtt ggcctggtat 120

cagcagaaac cagggagagc ccctaaggtc ttgatctata aggcatctac tttagaaagt 180

ggggtcccat caaggttcag cggcagtgga tctgggacag atttcactct caccatcagc 240

agtctgcaac ctgaagattt tgcaacttac tactgtcaac agagttacag taccccgtgg 300

acgttcggcc aggggaccaa gctggaaatc aa 332

<210> 49
<211> 112
<212> PRT
<213> Homo sapiens

<400> 49

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

Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly
20 25 30

Ile Ser Ser Trp Leu Ala Trp Tyr Gln Gln Lys Pro Gly Arg Ala Pro
35 40 45

Lys Val Leu Ile Tyr Lys Ala Ser Thr Leu Glu Ser Gly Val Pro Ser
50 55 60

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

Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr
85 90 95

Ser Thr Pro Trp Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg
100 105 110

<210> 50
<211> 6
<212> PRT
<213> Homo sapiens

<400> 50

Ser Ser Tyr Trp Met Ser
1 5

<210> 51
<211> 17
<212> PRT
<213> Homo sapiens

<400> 51

Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val Lys
1 5 10 15

Gly

<210> 52
<211> 9
<212> PRT
<213> Homo sapiens

<400> 52

Glu Asn Trp Gly Ser Tyr Phe Asp Leu
1 5

<210> 53
<211> 11
<212> PRT
<213> Homo sapiens

<400> 53

Arg Ala Ser Gln Gly Ile Ser Ser Trp Leu Ala
1 5 10

<210> 54
<211> 7
<212> PRT
<213> Homo sapiens

<400> 54

Lys Ala Ser Thr Leu Glu Ser
1 5

<210> 55
<211> 9
<212> PRT
<213> Homo sapiens

<400> 55

Gln Gln Ser Tyr Ser Thr Pro Trp Thr
1 5

<210> 56
<211> 288
<212> PRT
<213> Murine

<400> 56

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

Leu Ser Trp Gln Ser Gly Trp Leu Leu Glu Val Pro Asn Gly Pro Trp
20 25 30

Arg Ser Leu Thr Phe Tyr Pro Ala Trp Leu Thr Val Ser Glu Gly Ala
35 40 45

Asn Ala Thr Phe Thr Cys Ser Leu Ser Asn Trp Ser Glu Asp Leu Met
50 55 60

Leu Asn Trp Asn Arg Leu Ser Pro Ser Asn Gln Thr Glu Lys Gln Ala
65 70 75 80

Ala Phe Cys Asn Gly Leu Ser Gln Pro Val Gln Asp Ala Arg Phe Gln
85 90 95

```
Ile Ile Gln Leu Pro Asn Arg His Asp Phe His Met Asn Ile Leu Asp
            100                 105                 110

Thr Arg Arg Asn Asp Ser Gly Ile Tyr Leu Cys Gly Ala Ile Ser Leu
        115                 120                 125

His Pro Lys Ala Lys Ile Glu Glu Ser Pro Gly Ala Glu Leu Val Val
    130                 135                 140

Thr Glu Arg Ile Leu Glu Thr Ser Thr Arg Tyr Pro Ser Pro Ser Pro
145                 150                 155                 160

Lys Pro Glu Gly Arg Phe Gln Gly Met Val Ile Gly Ile Met Ser Ala
                165                 170                 175

Leu Val Gly Ile Pro Val Leu Leu Leu Leu Ala Trp Ala Leu Ala Val
            180                 185                 190

Phe Cys Ser Thr Ser Met Ser Glu Ala Arg Gly Ala Gly Ser Lys Asp
        195                 200                 205

Asp Thr Leu Lys Glu Glu Pro Ser Ala Ala Pro Val Pro Ser Val Ala
    210                 215                 220

Tyr Glu Glu Leu Asp Phe Gln Gly Arg Glu Lys Thr Pro Glu Leu Pro
225                 230                 235                 240

Thr Ala Cys Val His Thr Glu Tyr Ala Thr Ile Val Phe Thr Glu Gly
                245                 250                 255
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Leu Gly Ala Ser Ala Met Gly Arg Arg Gly Ser Ala Asp Gly Leu Gln
260 265 270

Gly Pro Arg Pro Pro Arg His Glu Asp Gly His Cys Ser Trp Pro Leu
275 280 285

<210> 57
<211> 321
<212> DNA
<213> Homo sapiens

<400> 57
actgtggctg caccatctgt cttcatcttc ccgccatctg atgagcagtt gaaatctgga 60

actgcctctg ttgtgtgcct gctgaataac ttctatccca gagaggccaa agtacagtgg 120

aaggtggata acgccctcca atcgggtaac tcccaggaga gtgtcacaga gcaggacagc 180

aaggacagca cctacagcct cagcagcacc ctgacgctga gcaaagcaga ctacgagaaa 240

cacaaagtct acgcctgcga agtcacccat cagggcctga gctcgcccgt cacaaagagc 300

ttcaacaggg gagagtgtta g 321

<210> 58
<211> 108
<212> PRT
<213> Homo sapiens

<400> 58

His Met Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
1 5 10 15

Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
20 25 30

Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu
35 40 45

Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp
50 55 60

Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr
65 70 75 80

Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser
85 90 95

Ser Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
100 105

שרד המשפטים

סמך זה הינו העתק שנסרק בשלמותו ביום ובשעה המצוינים .

ריקה ממוחשבת מהימנה מהמסמך המצוי בחיק.

תאם לנוהל הבדיקות במשרד המשפטים.

החתום

משרד המשפטים (חתימה מוסדית).

We claim:

1. An isolated antibody comprising:
a VH domain comprising SEQ ID NO:6, SEQ ID NO:10 , SEQ ID NO:14 or SEQ ID NO:47, wherein the antibody specifically binds an epitope within the extracellular domain of human or mouse PD-1.

2. An isolated antibody comprising:
a VL domain comprising SEQ ID NO:8, SEQ ID NO:12, SEQ ID NO:16 or SEQ ID NO:49, wherein the antibody specifically binds an epitope within the extracellular domain of human or mouse PD-1.

3. An isolated antibody comprising:
a VH domain or an antigen-binding fragment thereof that comprises 3 CDRs; and a VL domain or an antigen-binding fragment thereof that comprises 3 CDRs; wherein the antibody comprises at least 3 VH domain CDRs having sequences of:
SEQ ID NO:23, SEQ ID NO:24 and SEQ ID NO:25;
or SEQ ID NO:29, SEQ ID NO:30 and SEQ ID NO:31;
or SEQ ID NO:35, SEQ ID NO:36 and SEQ ID NO:37;
or SEQ ID NO:50, SEQ ID NO:51 and SEQ ID NO:52;
or at least 3 VL domain CDRs having sequences of:
SEQ ID NO:26, SEQ ID NO:27 and SEQ ID NO:28;
or SEQ ID NO:32, SEQ ID NO:33 and SEQ ID NO:34;
or SEQ ID NO:38, SEQ ID NO:39 and SEQ ID NO:40;
or SEQ ID NO:53, SEQ ID NO:54, and SEQ ID NO:55; and wherein the antibody specifically binds an epitope within the extracellular domain of human or mouse PD-1.

4. An antibody produced by steps comprising:
(a) providing a nucleic acid encoding 3 VL domain CDRs having sequences of: SEQ ID NO:26, SEQ ID NO:27 and SEQ ID NO:28; or SEQ ID NO:32, SEQ ID NO:33 and SEQ ID NO:34; or SEQ ID NO:38, SEQ ID NO:39 and SEQ ID NO:40; or SEQ ID NO:53, SEQ ID NO:54, and SEQ ID NO:55;
(b) combining a repertoire of nucleic acids encoding 3 VH domain CDRs with the nucleic acid encoding the 3 VL domain CDRs, so as to provide a product repertoire of nucleic acids encoding the 3 VL domain CDRs and the repertoire of 3 VH domain CDRs
(c) expressing the nucleic acids of the product repertoire;
(d) selecting an antigen-binding fragment comprising a variable domain that specifically binds to PD-1 and that is expressed from the nucleic acids of the product repertoire; and
(e) producing an antibody comprising the antigen-binding fragment.

5. The isolated antibody of claim 1, wherein the antibody comprises:
a VH domain comprising SEQ ID NO:6; and
a VL domain comprising SEQ ID NO:8.

6. The isolated antibody of claim 1, wherein the antibody comprises:
a VH domain comprising SEQ ID NO:10; and
a VL domain comprising SEQ ID NO:12.

7. The isolated antibody of claim 1, wherein the antibody comprises:
a VH domain comprising SEQ ID NO:14; and
a VL domain comprising SEQ ID NO:16.

8. The isolated antibody of claim 1, wherein the antibody comprises:
a VH domain comprising SEQ ID NO:47; and
a VL domain comprising SEQ ID NO:49.

9. The isolated antibody of claim 1, wherein the antibody comprises a VL domain comprising SEQ ID NO:8.

10. The isolated antibody of claim 1, wherein the antibody comprises a VL domain comprising SEQ ID NO:12.

11. The isolated antibody of claim 1, wherein the antibody comprises a VL domain comprising SEQ ID NO:16.

12. The isolated antibody of claim 1, wherein the antibody comprises a VL domain comprising SEQ ID NO:49.

13. The isolated antibody of any one of claims 1-3, wherein the antibody is an antigen binding fragment Fab, $F(ab')_2$, Fv, scFv, Fd or dAb.

14. The isolated antibody of any one of claims 1-3, wherein the antibody specifically binds to the extracellular domain of PD-1 with an affinity constant greater than 10^7 M^{-1}.

15. The isolated antibody of any one of claims 1-3, where the antibody inhibits the binding of PD-L1 or PD-L2 to PD-1 with an IC_{50} of less than 10nM.

16. The isolated antibody of any one of claims 1-3, wherein the antibody is a human antibody.

17. The isolated antibody of any one of claims 1-3, wherein the antibody is IgG_1 or IgG_4.

18. The isolated antibody of any one of claims 1-3, wherein the antibody is $IgG_{1\lambda}$ or IgG_{1k}.

19. A pharmaceutical composition comprising the isolated antibody of any one of claims 1-3, 4-12.

20. The isolated antibody of claim 2, wherein the antibody comprises a VL domain comprising SEQ ID NO:8.

21. The isolated antibody of claim 2, wherein the antibody comprises a VL domain comprising SEQ ID NO: 12.

22. The isolated antibody of claim 2, wherein the antibody comprises a VL domain comprising SEQ ID NO:16.

23. The isolated antibody of claim 2, wherein the antibody comprises a VL domain comprising SEQ ID NO:49.

For the Applicant,

Sanford T. Colb & Co
C: 55305

שרד המשפטים

סמך זה הינו העתק שנסרק בשלמותו ביום ובשעה המצוינים .

ריקה ממוחשבת מהימנה מהמסמך המצוי בחיק.

תאם לנוהל הבדיקות במשרד המשפטים.

החתום

משרד המשפטים (חתימה מוסדית).

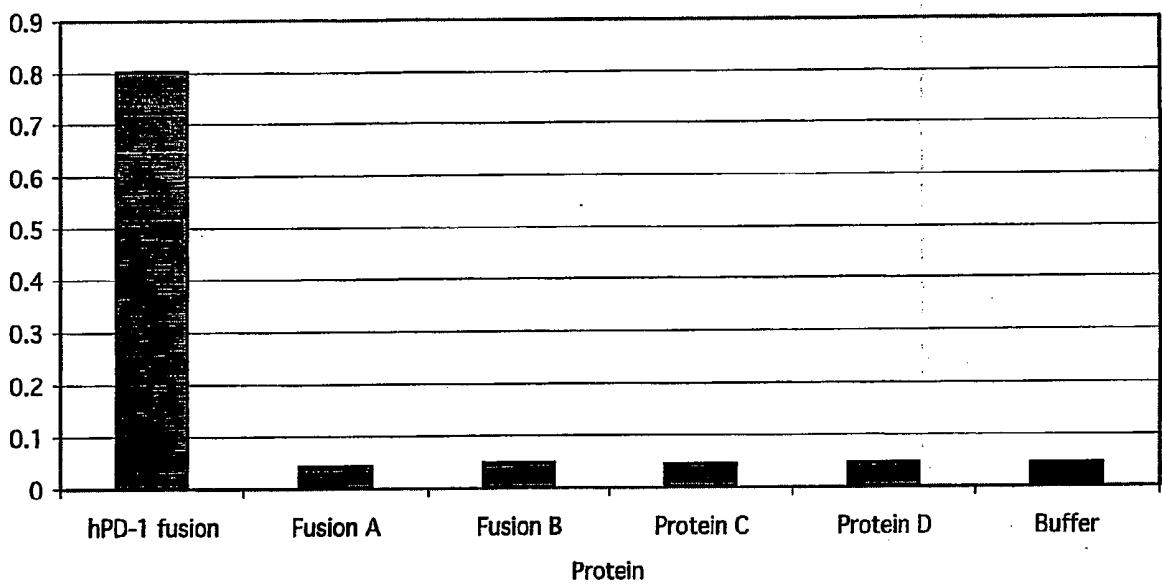
0.9
0.8
0.7
0.6
0.5
0.4
0.3
0.2
0.1
0
hPD-1 fusion
Fusion A
Fusion B
Protein C
Protein D
Buffer
Protein

Fig. 1A

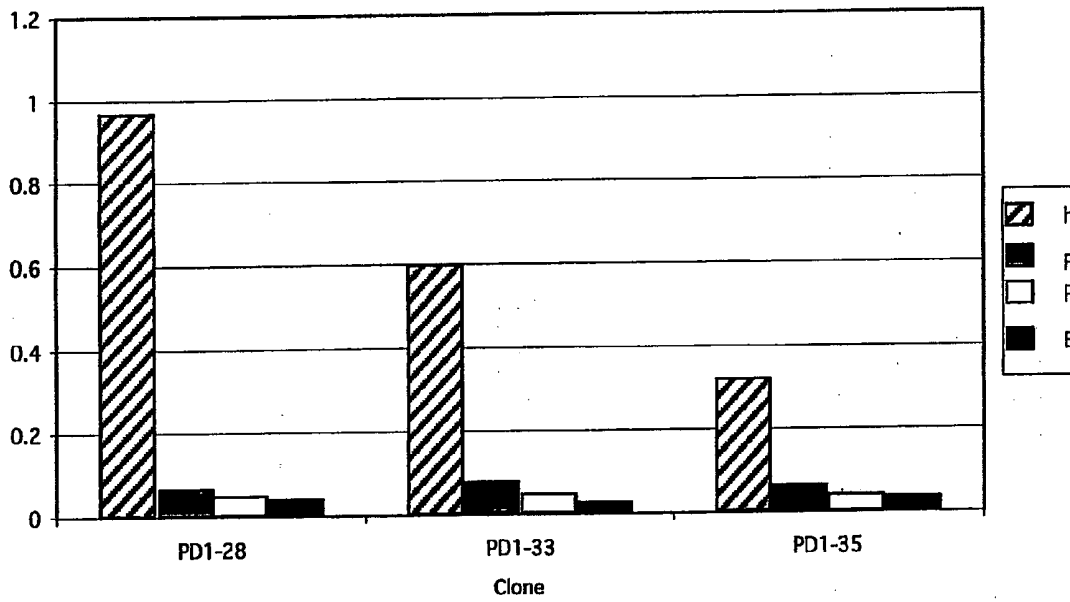

Fig. 1B

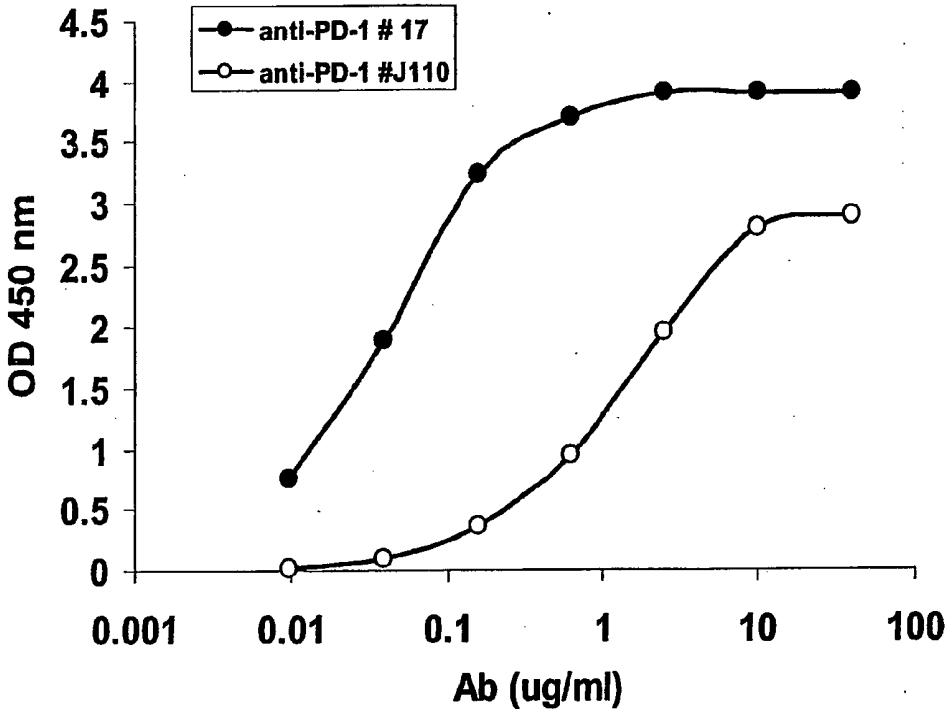
anti-PD-1 # 17
anti-PD-1 #J110
OD 450 nm
4.5
4
3.5
3
2.5
2
1.5
1
0.5
0
0.001
0.01
0.1
1
10
100
Ab (ug/ml)

Fig. 2A

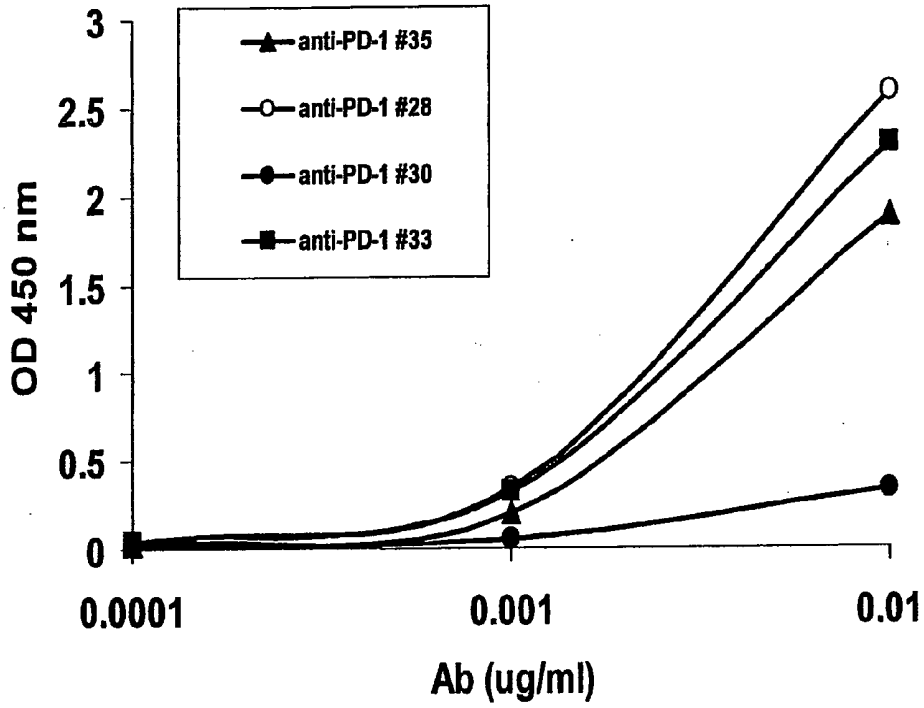
anti-PD-1 #35
anti-PD-1 #28
anti-PD-1 #30
anti-PD-1 #33
OD 450 nm
3
2.5
2
1.5
1
0.5
0
0.0001
0.001
0.01
Ab (ug/ml)

Fig. 2B

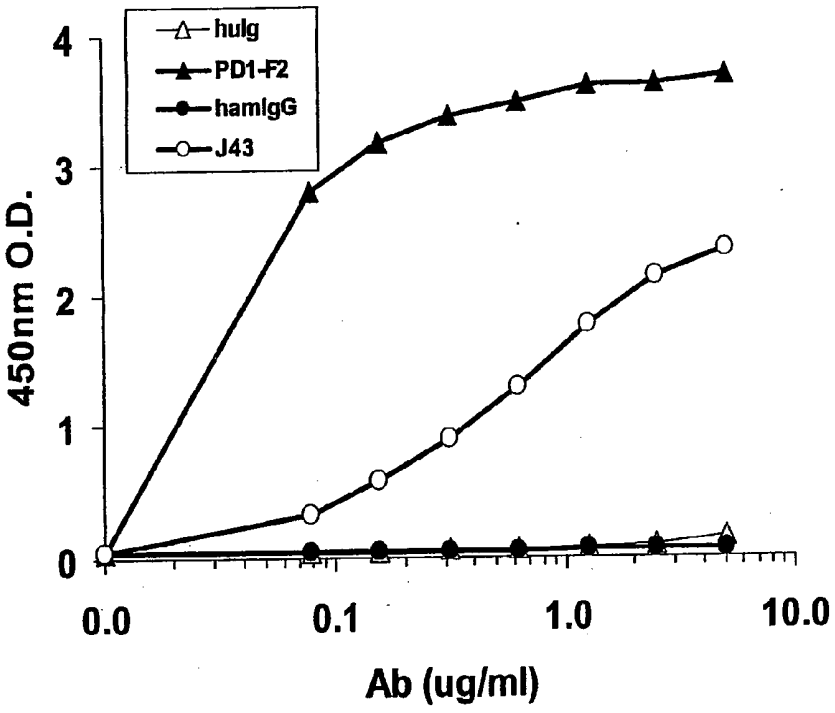

Fig. 2C

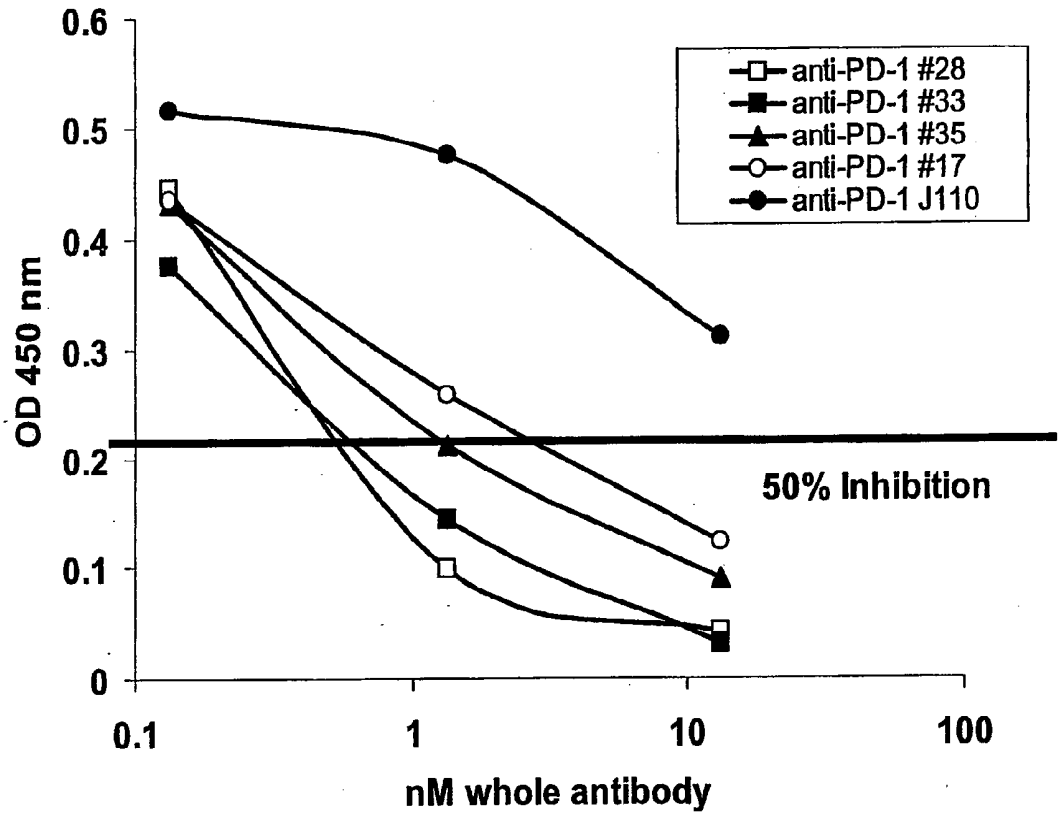
0.6
0.5
0.4
0.3
0.2
0.1
0
OD 450 nm
0.1
1
10
100
nM whole antibody
anti-PD-1 #28
anti-PD-1 #33
anti-PD-1 #35
anti-PD-1 #17
anti-PD-1 J110
50% Inhibition

Fig. 3

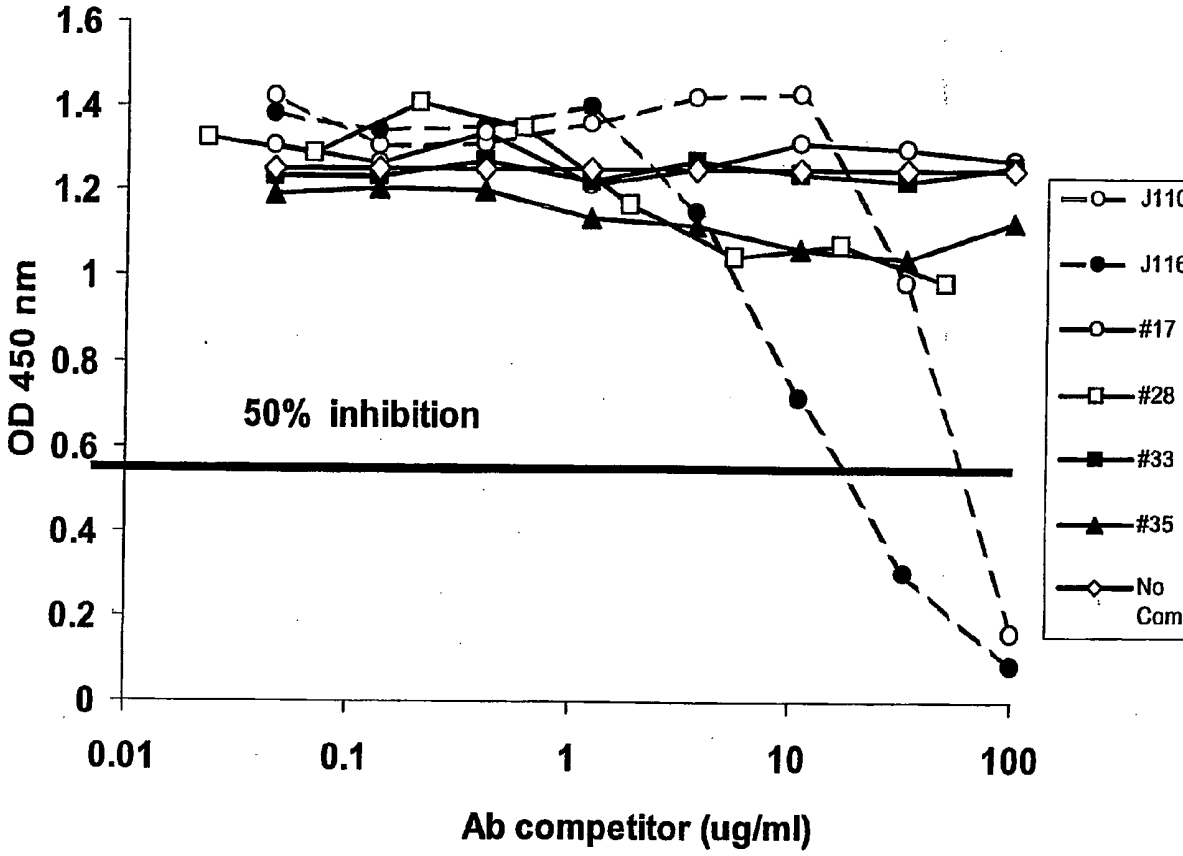
1.6
1.4
1.2
1
0.8
0.6
0.4
0.2
0
OD 450 nm
50% inhibition
0.01
0.1
1
10
100
Ab competitor (ug/ml)
J110
J116
#17
#28
#33
#35
No
Com

Fig. 4

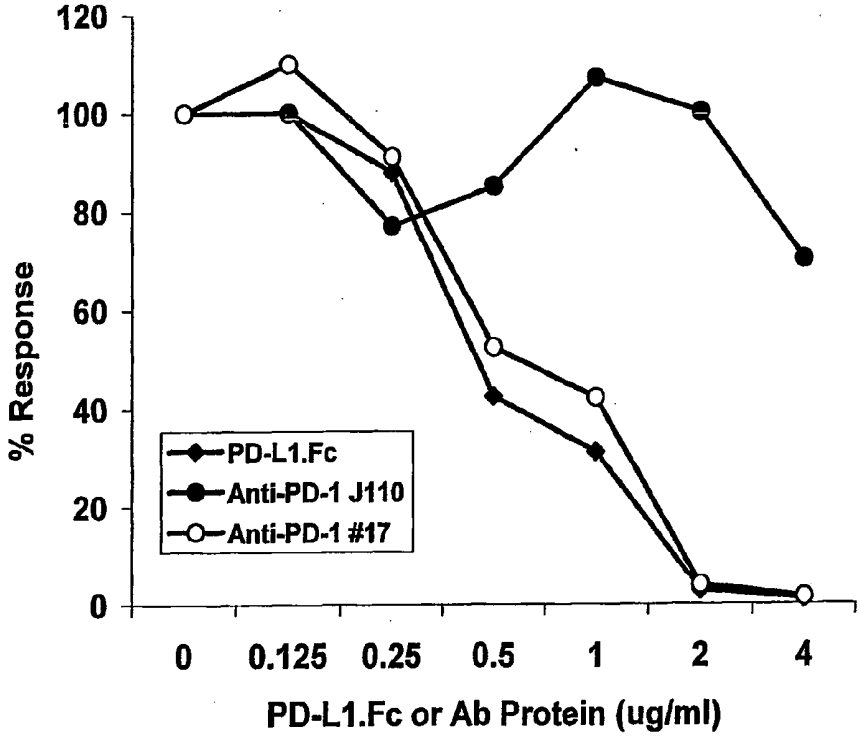
120
100
80
60
40
20
0
% Response
PD-L1.Fc
Anti-PD-1 J110
Anti-PD-1 #17
0
0.125
0.25
0.5
1
2
4
PD-L1.Fc or Ab Protein (ug/ml)

Fig. 5

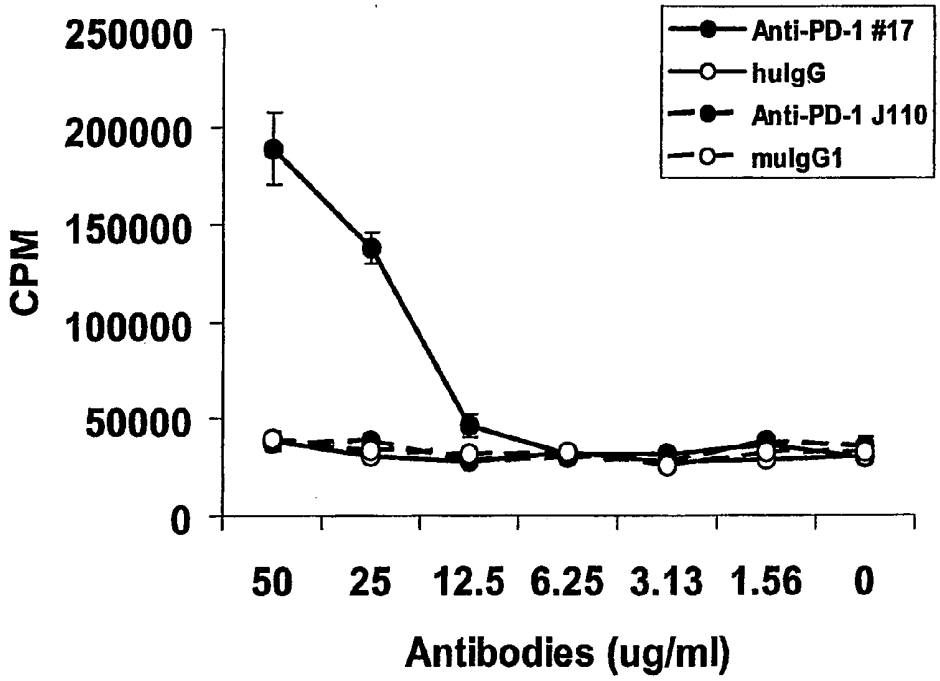
250000
200000
150000
100000
50000
0
CPM
Anti-PD-1 #17
huIgG
Anti-PD-1 J110
muIgG1
50
25
12.5
6.25
3.13
1.56
0
Antibodies (ug/ml)

Fig. 6

שרד המשפטים

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החתום

משרד המשפטים (חתימה מוסדית).