Supplementary Information Appendix

Supplementary Materials and Methods

The Ig variable sequences of the new anti-PD1 antibodies that were used in the study:

The sequences encode the Ig variable regions of the new anti-PD1 antibodies that were used in the experiments in Figure 3-4, Figure S2 and S3. In the expression constructs, the human V_H3-33 leader sequence and the human V_K3-11 leader sequence were appended to the 5' end of the variable region sequences of the HCs and LCs, respectively; human V_H3-33 and V_K3-11 segments are germline components of the 17D8 antibody heavy and light chains, respectively. The IgH variable regions were expressed in association with the human IgG4 constant region, and the IgL variable regions were expressed in association with the human kappa constant region. The complete cDNAs for the HCs and LCs were cloned into the pcDNA expression vector and expressed in 293F or Expi293 cells.

M1-1HC

CAGGTGCAGTTGGTGGAGTCTGGGGGGAGACGTGGTCCAGCCTGGGGGGGTCCCTGAGACT CTCCTGTGCAGCGTCTGGAGTCGCCTTCAGGGACTATGGCATGCACTGGGTCCGCCAGG CACCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATGGTATGATGGAAGTAAGAAATATT ATGGAGACTCCGTGAAGGGCCGATTCACCGTCTCCAGAGACAATTCCAAGAACATGTTGTA TCTGGAAATGAACGGCCTGAAAGCCGAGGACACGGCAATGTATTATTGTGCGAGGAACGA TGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCAG

M1-2HC

M1-3HC

M1-4HC

M1-5HC

CCCAGGCAAGGGGCTGGAATGGTTGGCAGTTATATGGTACGACGGAGGTAGAAAACACTA TGCGGACTCCGTGAAGGGCCGGTTCACCATCTCCAGAGACAATTCCAGGAACATGCTCTT TCTGCAAATGAATGGACTGAGAGTCGACGACACGGCTATGTATTACTGTACGAGAAGCCAC TCTACCGATGATTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCAG

M1-6HC

M1-7HC

CAATTACAACTGGTGGAGTCTGGGGGGAGACGTGGTCCAGCCTGGGGGGTCCCTGAGACT CTCCTGTGCGGCGTCTGGGGTCGTCTTCAGTGACTTTGGCCTGGAATGGGTCCGCCAGGC TCCAGGCAAGGGGCTGGAGTGGCTGGCAGTTATCTGGTATGATGGAAGTCGGAAACATTA TGCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGGGACAATTCCAAGAACATGCTCTAT CTGCAAATGAACAGTCTGAGAGTCGAGGACACGGCTATGTACTACTGTGCGCGATGTCAC TCTAAAGAGGACTACTGGGGCCAGGGAACCCTTGTCACCGTCTCCTCAG

M2-1HC

M2-2HC

CAGGTGCAGTTGGTGGAGTCTGGGGGGAGACGTGGTCCAGCCTGGGGGGTCCCTGAGACT CTCCTGTTCAGCGTCTGGGCTCGTAATCAGTGACTATGGCATGAACTGGGTCCGCCAGGC TCCAGGCAAGGGGCTGGAGTGGGTCGGACTTATATGGTATGATGGAAGTAAAAAATATTAC TCAGACTTCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACATATTGTATC TACAAATGAACAACCTGAGAGCCGAGGACACGGCTATGTATTACTGTGCGAGATTTCTAAT AGGTGCGACGAGGAGGGGCAATGCTATGGACTATTGGGGTCAAGGAACCTCAGTCACCGT CTCATCAG

M2-3HC

M2-4HC

GCAGACTCCGTGAAGGGCCGCTTCACCATCTCCAGAGACAATTCCAAGAATATGTTGTATC TTCAAATGAATAGCCTGAGAGCCGAGGACACGGCTATGTATTACTGTGCGAGACTCTCTAT AGGTACGACCCATTACTTTGATACGGACGACTACTGGGGTCAAGGAACCTCAGTCACCGT CTCCTCAG

M2-5HC

M1-1LC

M1-2LC

GAAATTGTGTTGACACAGTTTCCGGCCACCCTGTCTCTGTCTCCCGGGGAAAGAGCCACC CTCTCCTGCAGGACCAGTCAGAATATTGACAGCGACTTAGCCTGGTTCCAACAGAAACCTG GCCAGGCTCCCAGGCTCATCATCTATGATGCATCCAACAGGGCCACTGGCATCCCAGCCA GGTTCAGTGGCGGTGGGTCTGGGACAGACTTCACTCTCACCATCACCAGCCTAGAGCCTG AAGATTTTGCAGTTTATTACTGTCAGCAGCGTACCACCTGGCCTCTCACTTTCGGCGGAGG GACCAAGGTGGAGATCAAAC

M1-3LC

GAAATTGTGTTGACACAGTCTCCAGCCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACCC TCTCCTGCCGGACCAGTCAGAGTGTTAGCAGCGACTTAGCCTGGTTCCAACAGAAACCTG GCCAGGCTCCCAGGCTCTTCATATTTGATGCATCCAAAAGGGTCAATGGCATCCCAGCCA GGTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCAGCAGCCTGGAACCTG AAGATTTTGCAGTTTATTATTGTCAGCAACGTACCGACTGGCCTCTCACTTTCGGCGGAGG GTCCAGGGTGGAGATCAAAC

M1-4LC

GAAATTGTGTTGACACAGTCTCCAGCCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACCC TCTCCTGTAGGGCCAGTCAGAGTATTAGCAACTACTTAGCCTGGTTCCAACAGAAATCTGG CCAGGCTCCCAGGCTCATCATCCATGATGCATTTAAACGGGCCGCTGGCATCCCAACCAG GTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCAGCAGTCTAGAGCCTGA AGATTTTGCAGTTTATTATTGTCAGCAGCGTGACAACTGGCCTCTCAATTTCGGCGGAGGG ACTAAGGTGGAGATCAAAC

M1-5LC

GAAATTGTGTTGACACAGTCGCCAGCCACTCTGTCTGTGTCTCCAGGGGAAAGAGCCACC CTCTCCTGTAGGGCCAGTCAGAGTATTAGCAGCGACTTAGCCTGGTTCCAACAGAAACCTG GCCAGGCTCCCAGGCTCATCATCCATGGTGCATCCAAAAGGGCCACTGGCATCCCAGCCA GGTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCAGCAGCCTAGAGCCTG AAGATTTTGCGGTTTATTACTGTCAGCAGCGTGACAGCTGGCCTCTCAATTTCGGCGGAGG GACCAAGGTGGAGATCAAAC

M1-6LC

M1-7LC

GAAATTGAAGTGACACAGTCTCCGGCCACCCTGTCCTTGTCTCCAGGGGAAAGAGCCACC CTCTCCTGTAGGGCCAGTCAGAGTATTGACACCGACTTAGCCTGGTTCCAGCAGAGACCT GGCCAGACTCCCAGACTCATCATCTATGATGCATCCAAAAGGGCCACTGGCATCCCAGCC AGGTTCAGTGGCGGTGGGTCTGGGACAGACTTCACTCTCACCATCAGCAGCCTAGAGCCT GAAGATTTTGCAGTTTACTACTGTCAGCAGCGTACCACCTGGCCTCTCACTTTCGGCGGAG GGACCAAGGTGGAGATCAAAC

M2-1LC

GAAGTTGTGTTGACACAGTCTCCAGCCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC CTCTCCTGCAGGGCCAGTCAGAGTATTGACAGCGACTTAGCCTGGTCCCAACAGAAAACT GGCCAGCCTCCCAGACTCATCATCTATGATGCATCCAACAGGGCCACTGGCATCCCAGCC AGGTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCAGTAGTCTAGAGCCT GAAGATTTTGCAGTTTATTATTGTCAGCAACGTAGCGACTGGCCTCTCACTTTCGGCGGAG GGACCAAGGTGGAGATCAGAC

M2-2LC

GAAGTTGTGTTGACACAGTCTCCAGCCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC CTCTCCTGCAGGGCCAGTCAGAGTATTGACAGCGACTTAGCCTGGTCCCAACAGAAACCT GGCCAGCCTCCCAGACTCATCATCTATGATGCATCCAACAGGGGCCACTGGCATCCCAGCC AGGTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCAGCAGCCTAGAGCCT GAAGATTTTGCAGTTTATTATTGTCAGCAACGTAGCGACTGGCCTCTCACTTTCGGCGGAG GGACCAAGGTGGAGATCAGAC

M2-3LC

GAAATTGTGTTGACACAGTCTCCAGTCATCCTGTCTTTGTCTCCAGGGGAAAGAGCCACCC TCTCCTGCAGGGCCAGTCAGAGTATTAGCAGCGACTTGGCCTGGTTCCAACAGACACCTG GCCAGGCTCCCAGGCTCATCATCTATGATGCATCCAACAGGGCCACTGGCATCCCAGCCA GGTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCAGCAGCCTAGAGCCTG AAGATTTTGCAGTTTATTACTGTCAGCAGCGGAGCAACTGGCCTCTCACTTTCGGCGGAGG GACCAAGGTGGAGATCAAAC

M2-4LC

GAAATTGTGTTGACACAGTCTCCAGCCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACCC TCTCCTGCAGGGCCAGTCAGAGTGTTAGCAGCTACTTAGCCTGGTACCAACAGAAGGTTG GCCAGGCTCCCAGGCTCATCATCTTTGATGCATCCAACAGGGCCACTGGCATCCCAGCCA GGTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCACCAGCCTAGATCCTG AAGATTTTGCAGTTTATTACTGTCAGCAACGTAGCGCCTGGCCTCTCACTTTCGGCGGAGG GACCAAGGTGGAGATCAGAC M2-5LC

GAAATTGTGCTGACACAGTCTCCAGCCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC CTCTCCTGCAGGGCCAGTCAGAGCATTAGCAGCGACTTAACCTGGTTCCAACAGAAACCT GGCCAGGCTCCCAGGCTCATCATCTATGATGCATCCAACAGGGGCCACTGGCATCCCAGCC AGGTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCAGTAGCCTCGAGCCT GAAGATTTTGTAGTTTATTACTGTCTGCAACGTAGCGACTGGCCTCTCACTTTCGGCGGAG GGACCAAGGTGGAGATCAAAC

| Α | | | | |
|------------------|--|------------------------------|--|---|
| 1708HC | | CDR H1 | GCATGCACTGGGTCCGCCAGGCTCCAGGCAAGGGGCTGGAGTGGGTG | |
| M1-1HC | | | | G176 |
| M1-2HC M1-3HC | Δ ΤG Τ G G ΔΔΔ | G Δ Δ G T | G.АС т | GGC. 176 |
| M1-4HC | TA | AGGT | | TCCGAC. 176 |
| M1-5HC | | CTCGTG. | | CGGAC. 176 |
| M1-6HC M1-7HC | AT.AA | GTGT | CG.AC | C |
| | | | | |
| | | | | |
| 17D8HC | CTATGCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACATGCTGTATCTGCAAATGAACA | GCCTGAGAGCCGAGGACACG | GCTATGTATTACTGTGCGAGGAACGATGACTACTG | GGGCCAGGGAACCCTGGTCACCGTCTCCTCAG 340 |
| M1-1HC M1-2HC | ТТ | A | AT | |
| M1-3HC | C | A | T | |
| M1-4HC M1-5HC | ТG | .AT | AAAAGTCACTCTCG | |
| M1-6HC | TT.CGT.ACTG | .AT | AACTAA.GCCACTCTCG | |
| M1-7HC | Τ | .тт | C.ATGTCACTCTA.AG | |
| | | | | |
| _ | | | | |
| В | | | | |
| | | CDR L1 | | CDR L2 |
| 17D8LC | GAAATTGTGTTGACACAGTCTCCAGCCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACCCTCTCCTGCAGGGCCA | GTCAGAGTGTTAGCAGCTAG | | ATCTATGATGCATCCAACAG 161 |
| M1-2LC | C | A.AGAG | T | 161 |
| M1-3LC | | G | TT | |
| M1-4LC M1-5LC | | | Т | CG |
| M1-6LC | | | T | TA. 161 |
| M1-7LC | | AUAC.U | AA | |
| | | | | |
| | | | CDR I 3 | |
| 17D8LC | GGCCACTGGCATCCCAGCCAGGTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCAGCAGCCTAGAG | CCTGAAGATTTTGCAGTTT | | GACCAAGGTGGAGATCAAAC 322 |
| M1-1LC | | Α | | |
| M1-2LC M1-3LC | | | T | |
| M1-4LC | GAT | | T | T |
| M1-5LC M1-6LC | Ст | | | |
| M1-7LC | | | . c | |
| | | | | |
| ~ | | | | |
| 6 | | | | |
| | | CDR H1 | | CDR H2 |
| 17D8HC | | CONTRACTAT | | TATGGTATGATGGAAGTAATAAATACTATGCAGA 185 |
| M2-1HC | | TAAGA | ۹ | |
| M2-2HC M2-3HC | | A.CGTA. | ТАТТС. | .C |
| M2-4HC | | G | | |
| M2-5HC | | | | |
| | | | | |
| | | | CDR H3 | |
| 17D8HC | CTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACATGCTGTATCTGCAAATGAACAGCCTGAGAGCCGAG | GACACGGCTATGTATTACTGT | GCGAGGAACGATGACTAC | IGGGGCCAGGGAACCCTGGTCACCGTCTCCTCAG 340 |
| M2-THC M2-2HC | | | ATTTCTAATAGGTGCGACGAGGAGGAGGGGGCAATGCTATGT | TA |
| M2-3HC | ATTTGGTTTTTT | TC.T | AGGGATACGACGGGGGCCCTG-GTTCACT ACT.TCTATAGGTACGACCCATTACTTTGATACGC | CAGTT.ATA358 T.A |
| M2-5HC | T.ATTC | ттт | TAGGGATACGACAGGGGCCATG-GTTCT | AGT.GTTTC 358 |
| | | | | |
| - | | | | |
| D | | | | |
| | | CDR L1 | | CDR L2 |
| 17D8LC | GAAATTGTGTTGACACAGTCTCCAGCCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACCCTCTCCTGCAGGGCCA | GT CAGAGTGTTAGCAGCTAG | CTTAGCCTGGTACCAACAGAAACCTGGCCAGGCTCCCAGGCTCATG | ATCTAT <mark>GATGCATCC</mark> AACAG 161 |
| M2-1LC M2-2LC | | AGAG A GA G | CACA. | |
| M2-3LC | | | GTC | 161 |
| M2-4LC | ۰۰۰۰۰ ۲ | CA G | GGT | T |
| M2-5LC | | ······· | | 101 |
| | | | | |
| | | | CDR L3 | |
| 17D8LC | GGCCACTGGCATCCCAGGCCAGGTTCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCAGCAGCCTAGAG | CCTGAAGATTTTGCAGTTTA | ATTACTGTCAGCAGCGTAGCAACTGGCCTCTCACTTTCGGCGGAGG | GACCAAGGTGGAGATCAAAC 322 |
| M2-1LC | ттт. | | TAG | G 322 |
| M2-3LC | | | | |
| M2-4LC | | т | AGC | G 322 |
| M2-5LC | | | | |

Fig. S1. DNA Sequence alignments of the Ig variable regions of new anti-PD1 antibodies with the 17D8HC. (*A*) Alignment of M1-1 to M1-7HCs with the 17D8HC. The sequence of the 17D8HC serves as the template for alignment; only SHMs in new anti-PD1 antibodies are shown below, "." Indicates identity, "-" indicates gap in alignment. The CDRs are shaded with color. Sections B-D are organized in the same way as this section. (*B*) Alignment of M1-1 to M1-7LCs with the 17D8LC. (*C*) Alignment of M2-1 to M2-5HCs with the 17D8HC. (*D*) Alignment of M2-1 to M2-5LCs with the 17D8LC.

Fig. S2



PD1 binding

Fig. S2. Analysis of PD1-binding activities of new anti-PD1 antibodies. This figure supplements Fig. 3 by showing the analysis of additional antibodies isolated from Mouse Models 1 and 2. Sections A-C in this figure correspond to A-C in Figure 3, Section D-F in this figure relate to E-G in Fig. 3.





PD-L2 binding

Fig. S3. The effects of new anti-PD1 antibodies on PD1/ligand interaction. This figure supplements Fig. 4 by presenting the characterization of additional antibodies isolated from Mouse Model 1 and 2. Sections A and B in this figure relate to (A) and (B) in Fig. 4.