

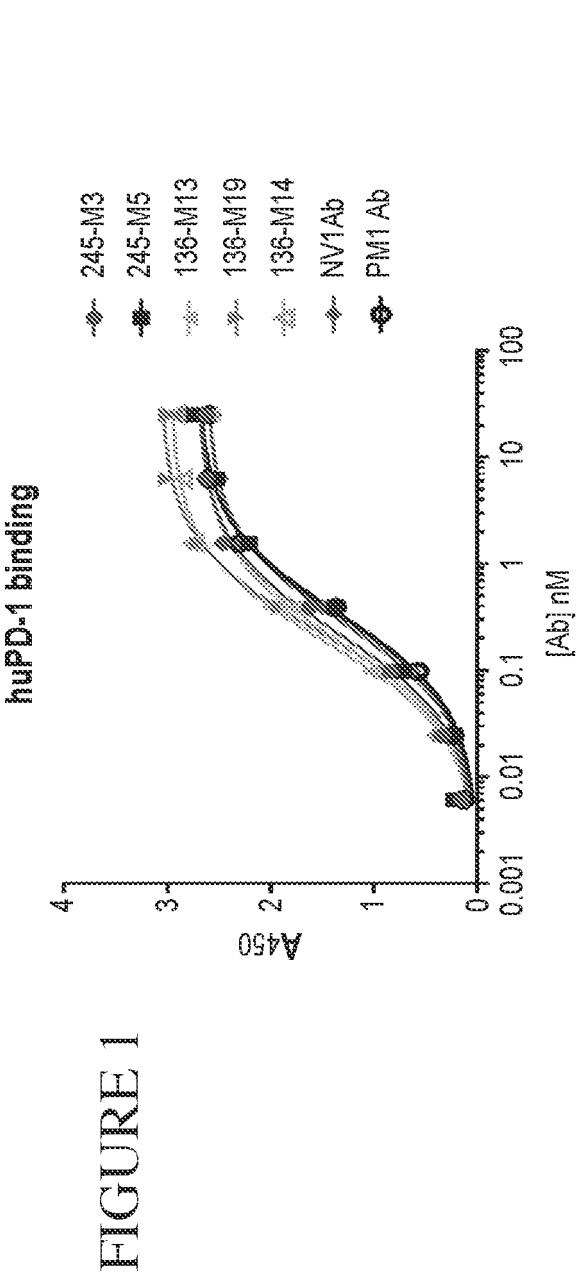


FIGURE 2

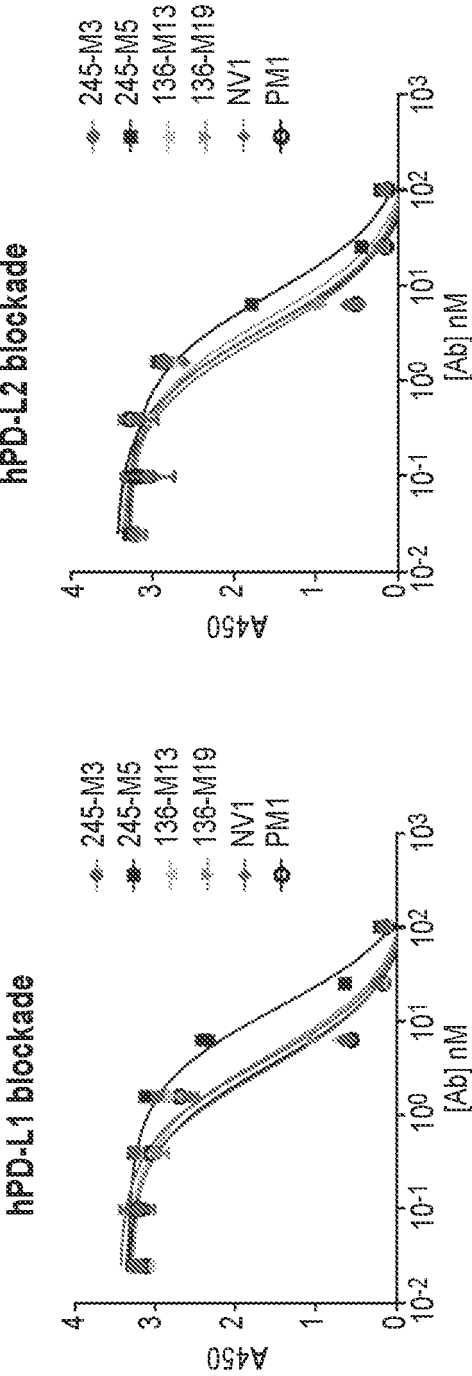
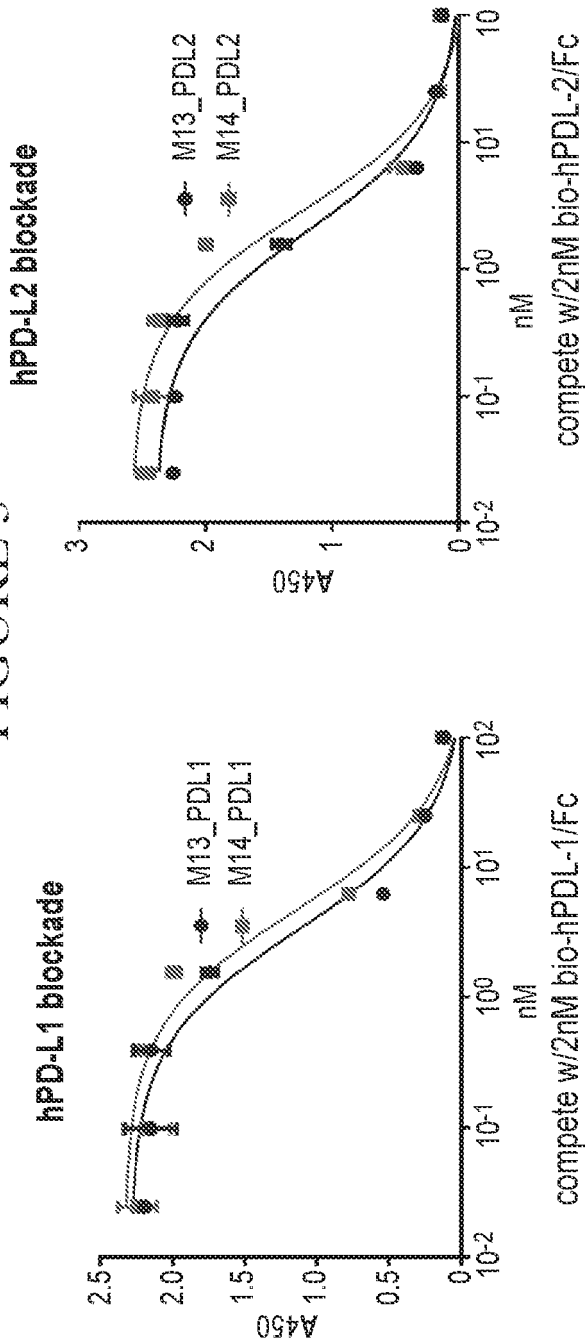


FIGURE 3



huPD-1 Binding

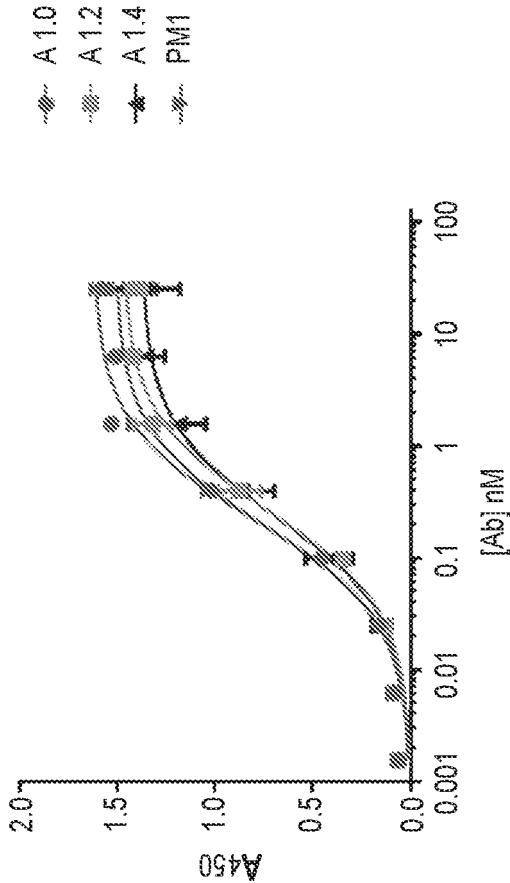


FIGURE 4

FIGURE 5

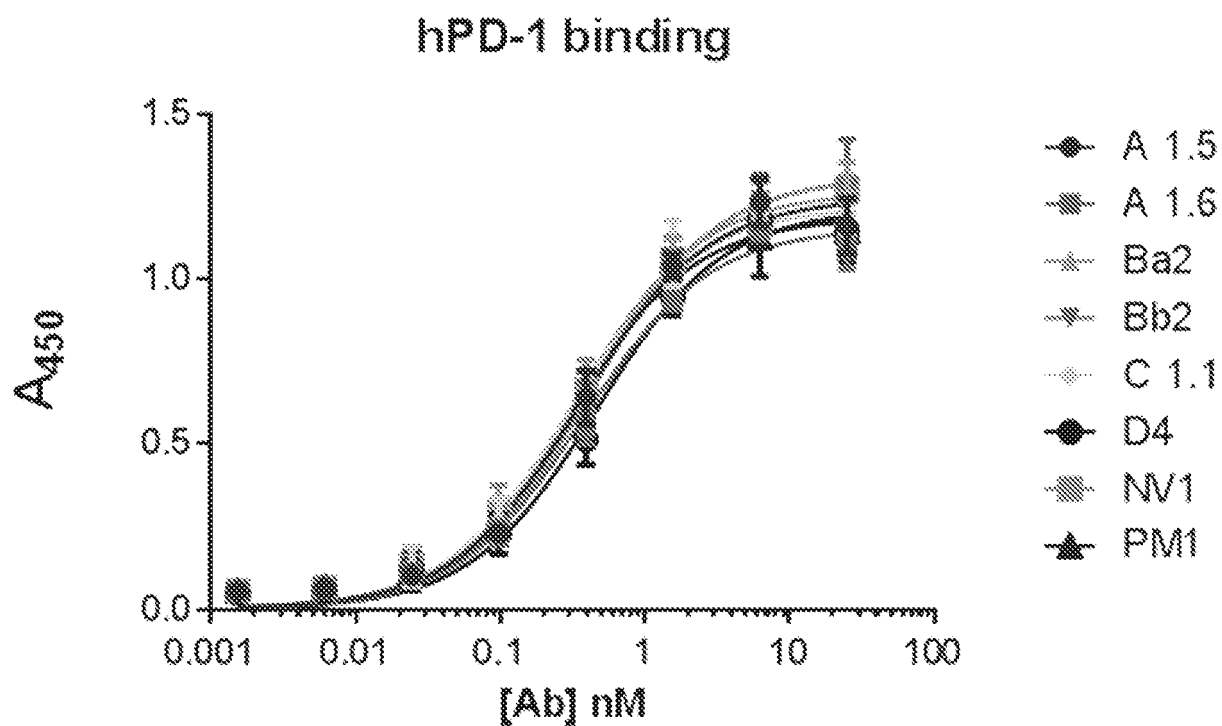
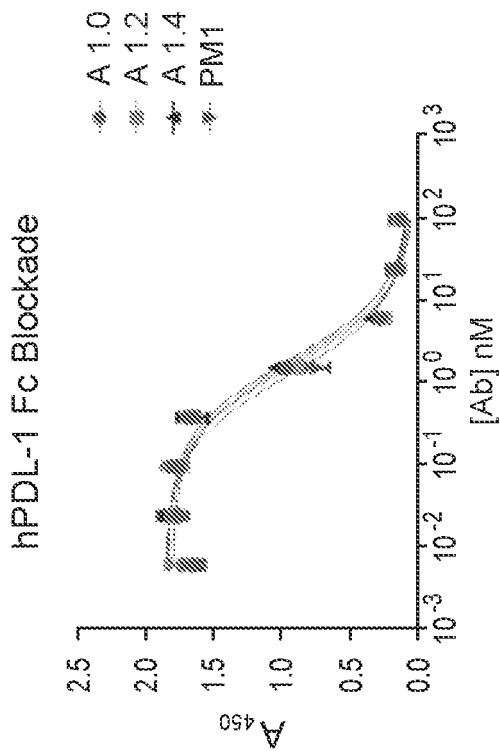
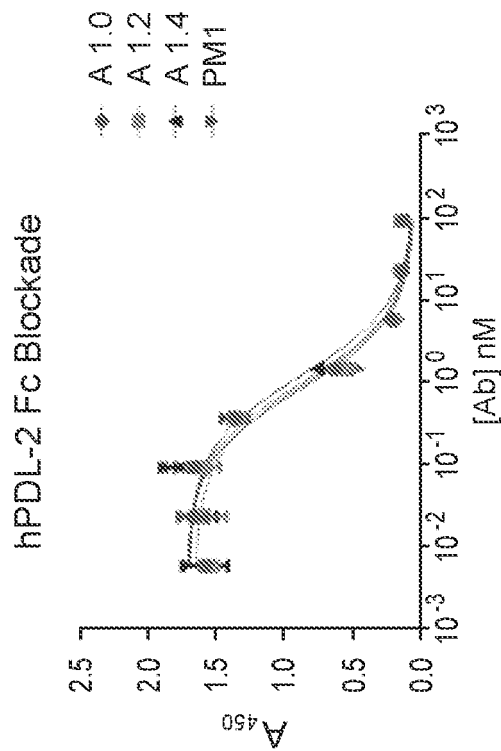


FIGURE 6

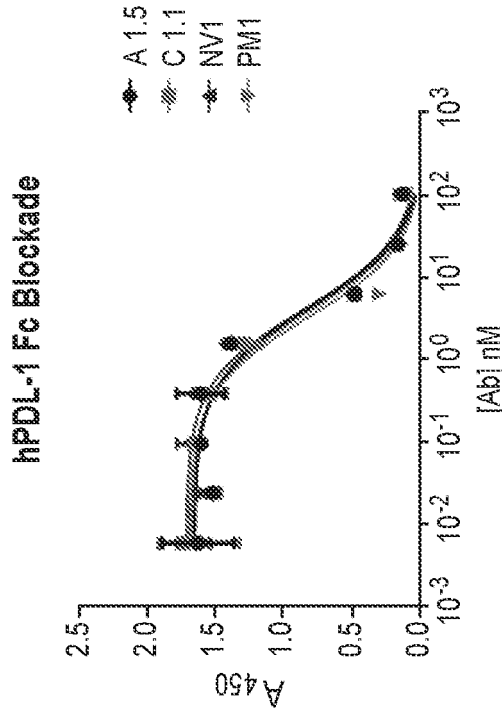


2nM of hPDL-1 Fc/hPDL-2Fc as blockade

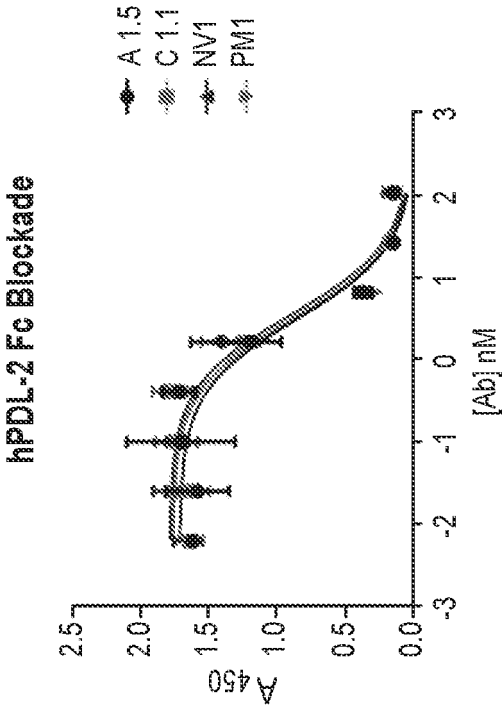
	A 1.0	A 1.2	A 1.4	PM1
log(inhibitor) vs. response (three parameters)				
Best-fit values				
Bottom	0.02473	0.03887	0.06139	0.07407
Top	1.824	1.869	1.862	1.876
LogIC50	0.2671	0.1900	0.2838	0.08887
IC50	1.850	1.549	1.922	1.227
Span	1.800	1.830	1.800	1.802

	A 1.0	A 1.2	A 1.4	PM1
log(inhibitor) vs. response (three parameters)				
Best-fit values				
Bottom	0.05839	0.06423	0.06416	0.05329
Top	1.742	1.687	1.734	1.750
LogIC50	0.01826	0.002967	0.1109	-0.05539
IC50	1.043	1.007	1.291	0.8803
Span	1.683	1.622	1.670	1.696

FIGURE 7



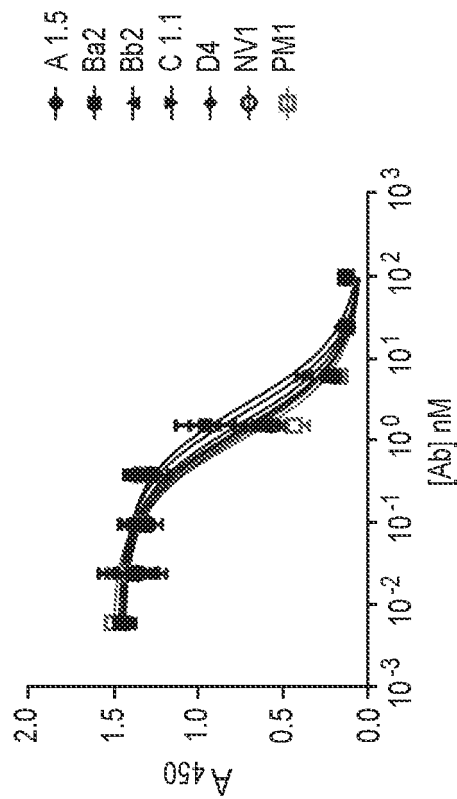
	A 1.5	C 1.1	NV1	PM1
log(inhibitor) vs. response (three parameters)				
Best-fit values				
Bottom	-0.01122	-0.002852	0.01597	-0.01207
Top	1.672	1.691	1.735	1.710
LogIC50	0.6289	0.5712	0.6045	0.5045
IC50	4.255	3.726	4.023	3.196



	A 1.5	C 1.1	NV1	PM1
log(inhibitor) vs. response (three parameters)				
Best-fit values				
Bottom	0.01162	0.002368	0.02642	0.003166
Top	1.733	1.784	1.804	1.779
LogIC50	0.4676	0.4875	0.5231	0.4797
IC50	2.935	3.073	3.335	3.018

FIGURE 8

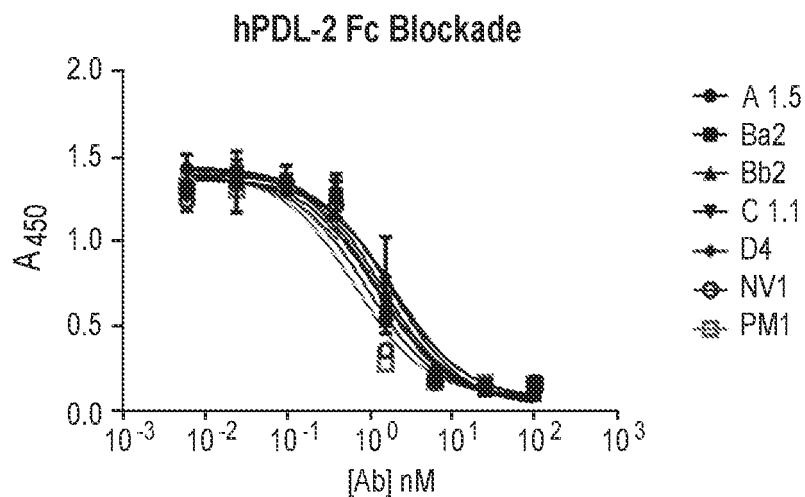
hPDL-1 Fc Blockade



	A 1.5	Ba2	Bb2	C 1.1	D4	NV1	PM1
log(inhibitor) vs. response (three parameters)							
Best-fit values							
Bottom	0.03776	0.03156	0.04762	0.05233	0.04490	0.05720	0.04908
Top	1.494	1.469	1.471	1.472	1.460	1.461	1.529
LogIC50	0.1465	0.3417	0.2221	0.1209	0.4346	0.07608	-0.01427
IC50	1.401	2.196	1.668	1.321	2.720	1.191	0.9677
Span	1.457	1.437	1.423	1.420	1.415	1.404	1.480

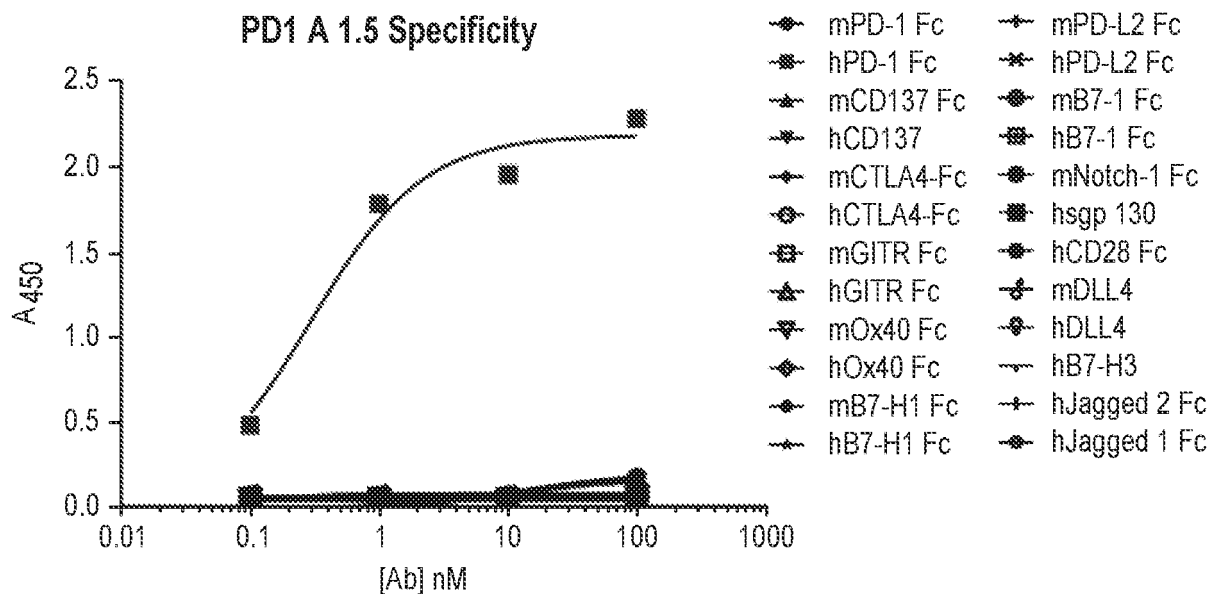
1.4571.4371.4231.4201.4151.4041.480

FIGURE 9



	A 1.5	Ba2	Bb2	C 1.1	D4	NV1	PM1
log(inhibitor) vs. response (three parameters)							
Best-fit values							
Bottom	0.05123	0.04694	0.02765	0.06529	0.06849	0.06895	0.08572
Top	1.445	1.430	1.404	1.385	1.370	1.391	1.404
LogIC50	0.1290	0.08061	0.2635	0.07807	0.2893	-0.03908	-0.1851
IC50	1.346	1.204	1.834	1.197	1.947	0.9140	0.6529
Span	1.344	1.383	1.376	1.320	1.302	1.322	1.318

FIGURE 10



K_d of A1.5 = 0.2948

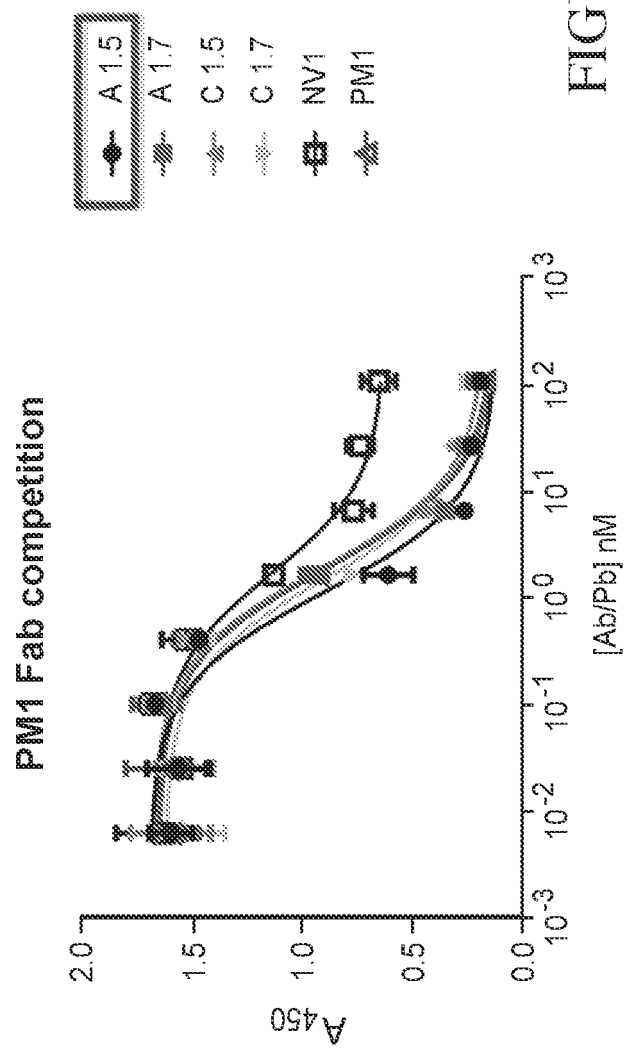


FIGURE 11a

	A 1.5	A 1.7	C 1.5	C 1.7	NV1	PM1
log(inhibitor) vs. response (three parameters)						
Best-fit values						
Bottom	0.1248	0.1450	0.1778	0.1928	0.6340	0.1206
Top	1.694	1.653	1.678	1.626	1.660	1.676
LogIC50	0.03121	0.2458	0.2429	0.1508	0.2042	0.2741
IC50	1.075	1.761	1.749	1.415	1.600	1.880

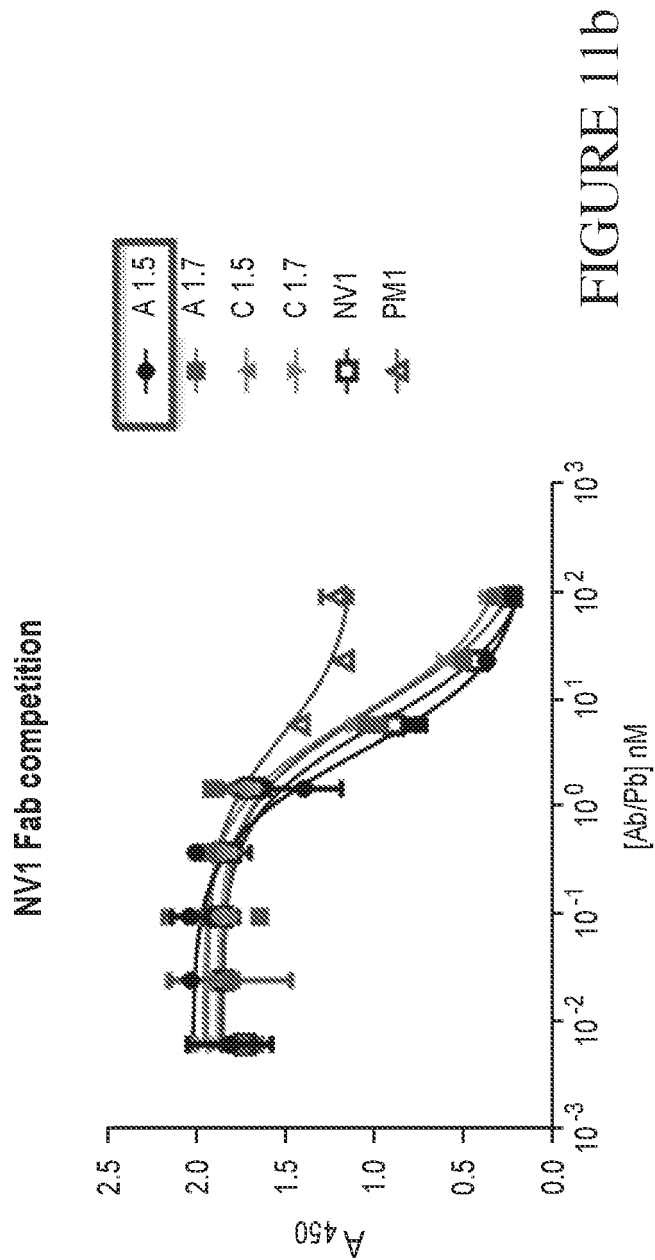
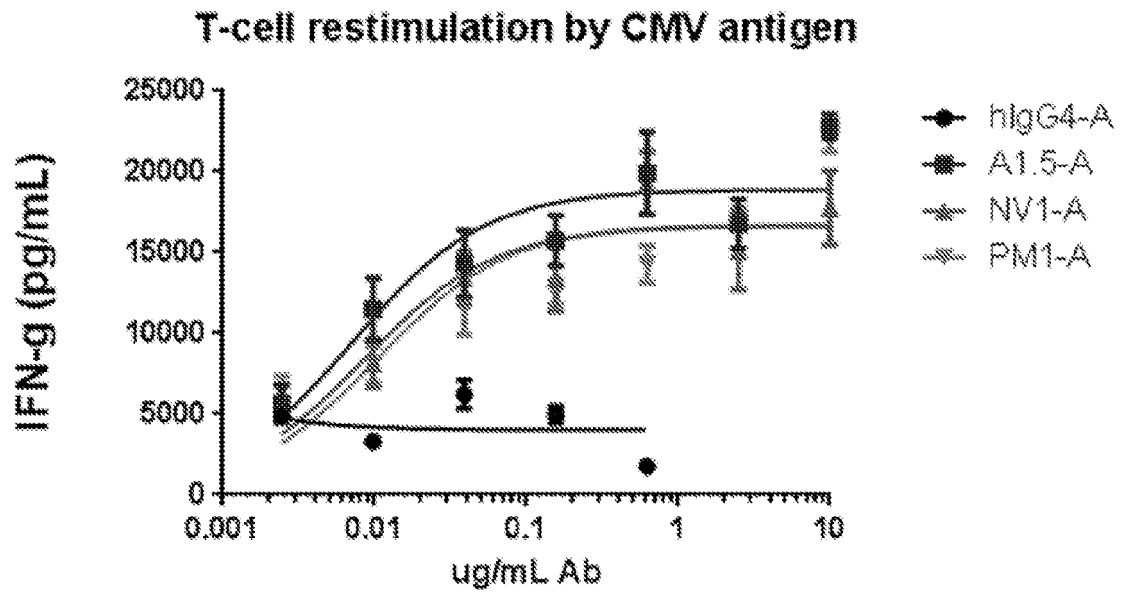


FIGURE 11b

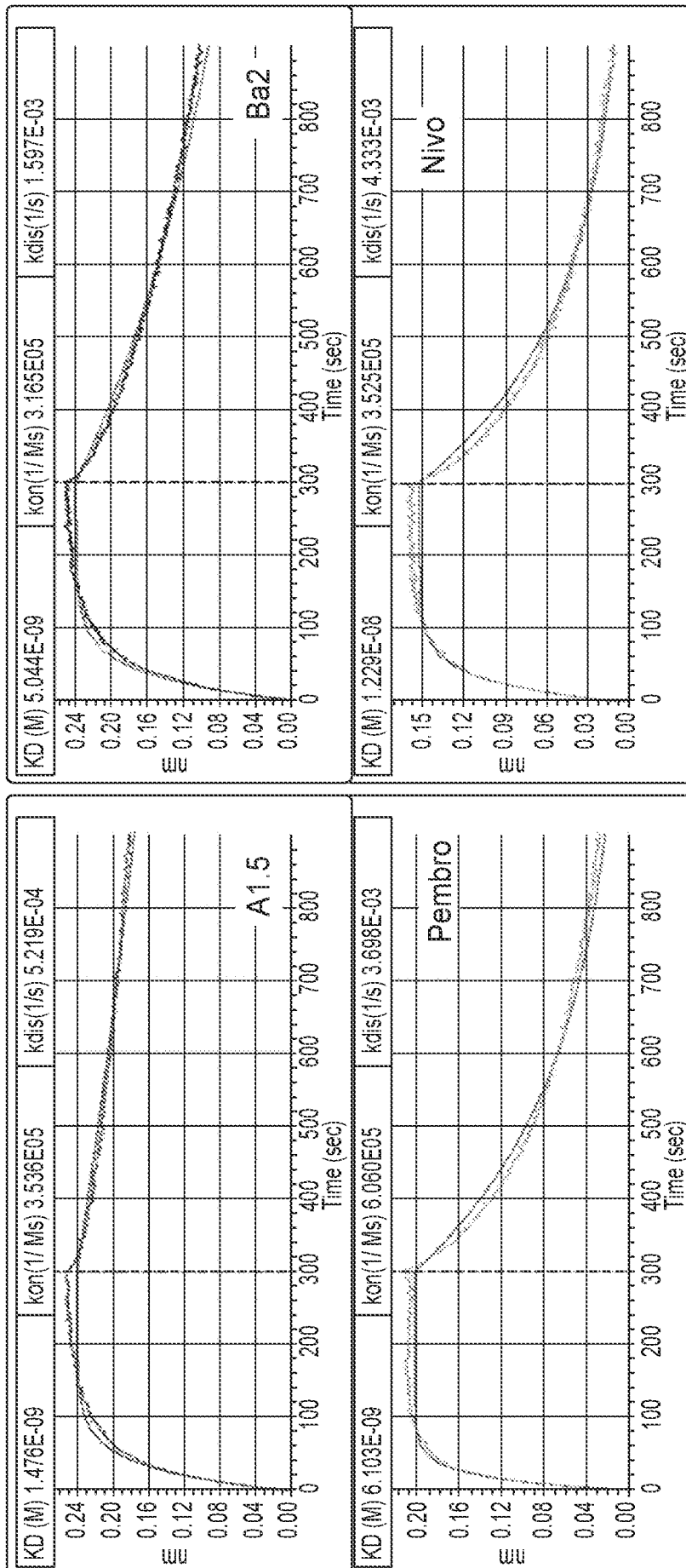
	A 1.5	A 1.7	C 1.5	C 1.7	NV1	PM1
log(inhibitor) vs. response (three parameters)						
Best-fit values						
Bottom	0.1491	0.1158	0.2335	0.2426	0.09043	1.140
Top	2.065	1.891	2.000	1.924	1.902	1.985
LogIC50	0.5390	0.9690	0.8798	0.8837	0.8130	0.6772
IC50	3.459	9.310	7.582	7.651	6.501	4.756

FIGURE 12



11/37

FIGURE 13



Sample ID	Conc. (nM)	Response	KD (M)	kon(1/Ms)	kon Error	kdis(1/s)	kdis Error	Full R^2	Loading Sample ID
hPD1-His	90	0.2526	1.48E-09	3.54E+05	1.82E+03	5.22E-04	2.75E-06	0.996503	A1.5 hlgG4
hPD1-His	90	0.1904	6.37E-09	2.96E+05	1.52E+03	1.89E-03	3.84E-06	0.997158	Ba2 hlgG4
hPD1-His	90	0.1981	1.24E-08	3.33E+05	2.08E+03	4.11E-03	7.81E-06	0.996302	Ct1.1 hlgG4
hPD1-His	90	0.2069	6.10E-09	6.06E+05	4.99E+03	3.70E-03	8.84E-06	0.994474	PM1 hlgG4
Sample ID	Conc. (nM)	Response	KD (M)	kon(1/Ms)	kon Error	kdis(1/s)	kdis Error	Full R^2	Loading Sample ID
hPD1-His	90	0.2487	5.04E-09	3.17E+05	2.16E+03	1.60E-03	4.65E-06	0.994442	Ba2 hlgG4
hPD1-His	90	0.1956	5.52E-09	1.62E+05	8.22E+02	8.95E-04	3.09E-06	0.997361	D4 hlgG4
hPD1-His	90	0.1577	1.23E-08	3.53E+05	3.27E+03	4.33E-03	1.19E-05	0.993183	NV1 hlgG4

FIGURE 14

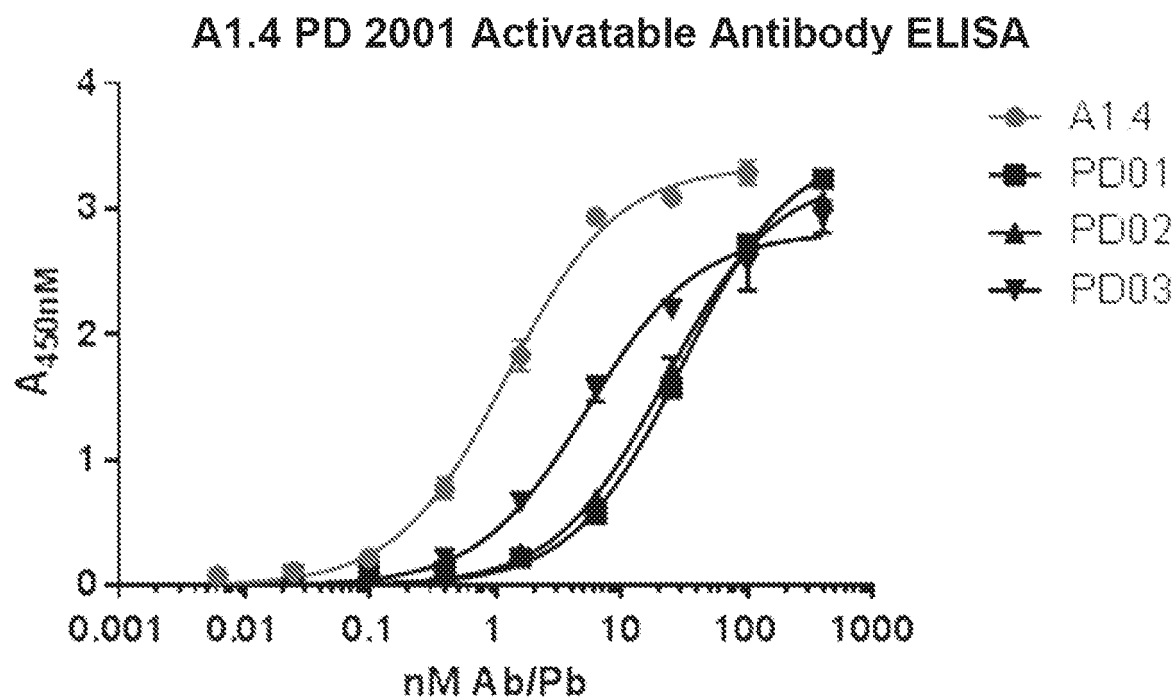
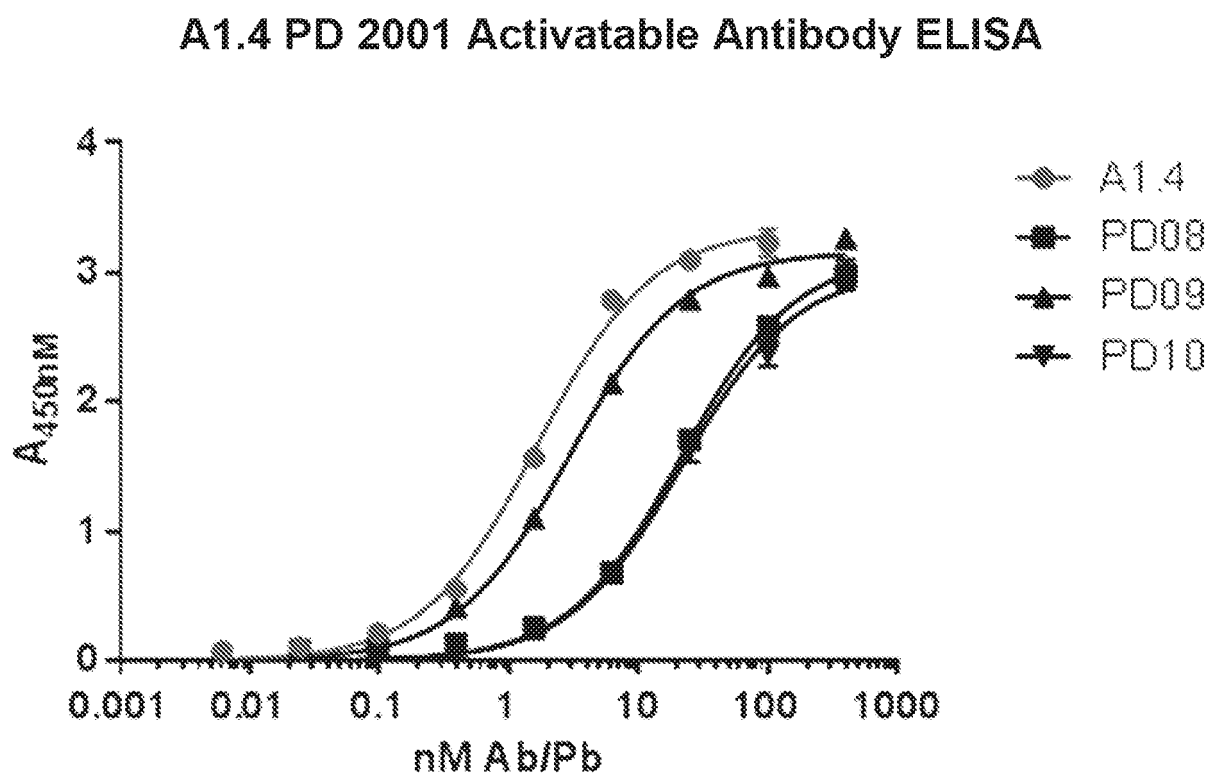
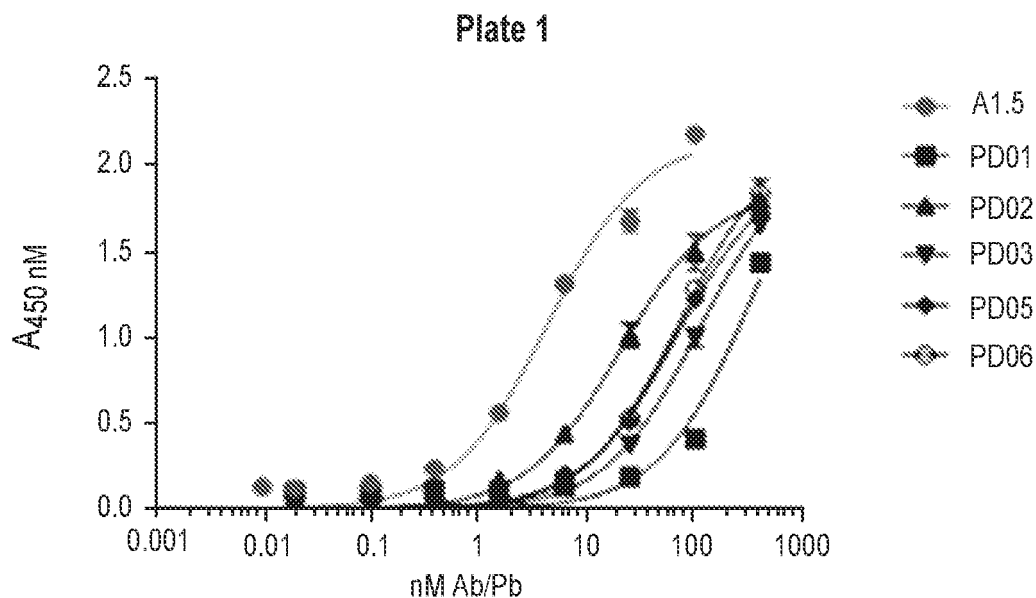
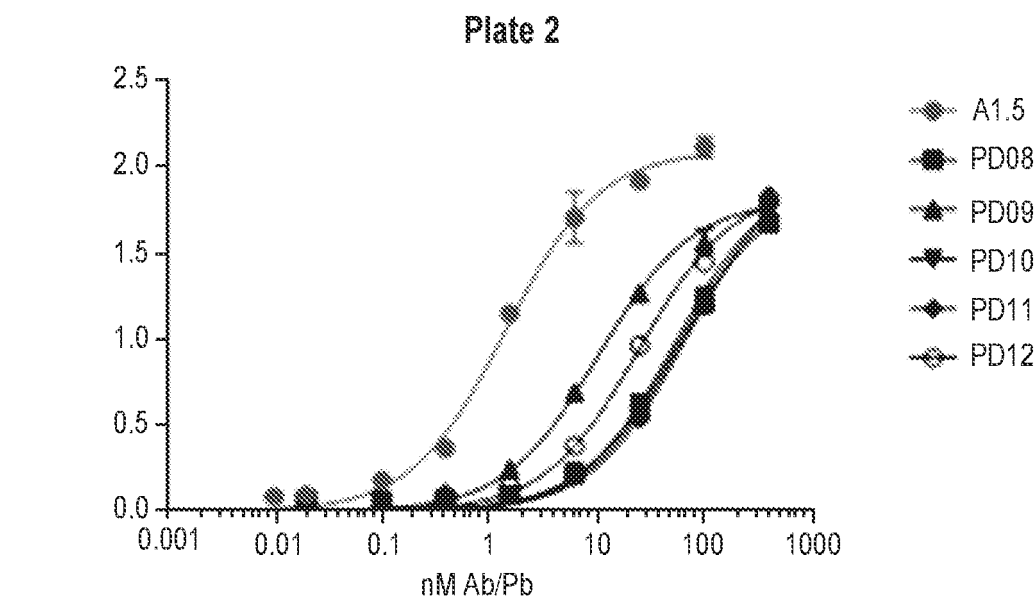


FIGURE 15



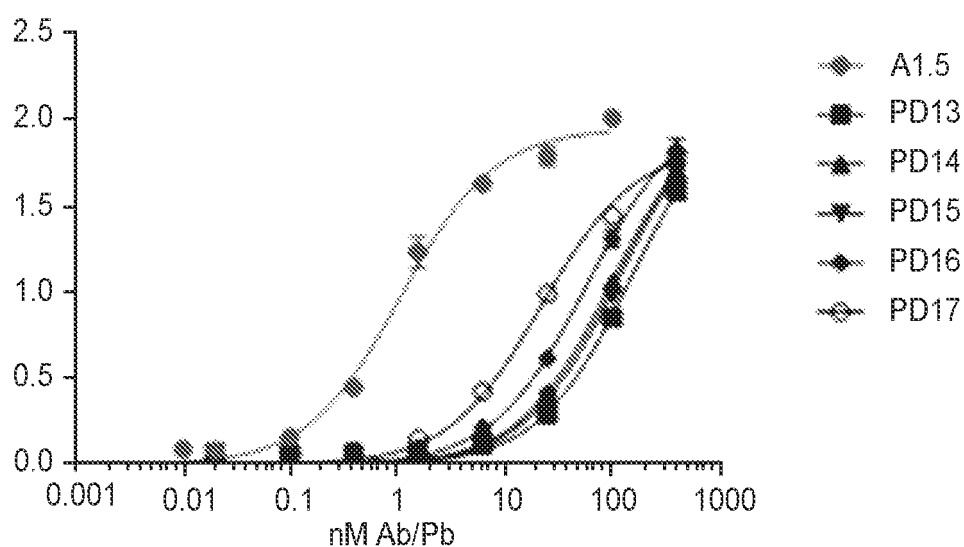
	A1.5	PD01	PD02	PD03	PD05	PD06
One site -- Specific binding		Hit constraint				
Best-fit values						
Bmax	2.145	-2.500	1.852	2.091	2.009	2.158
Kd	4.259	347.8	20.64	109.3	67.61	T1.01



	A1.5	PD08	PD09	PD10	PD11	PD12
One site -- Specific binding						
Best-fit values						
Bmax	2.100	1.899	1.815	1.910	2.017	1.859
Kd	1.417	55.20	10.25	52.97	67.47	24.69

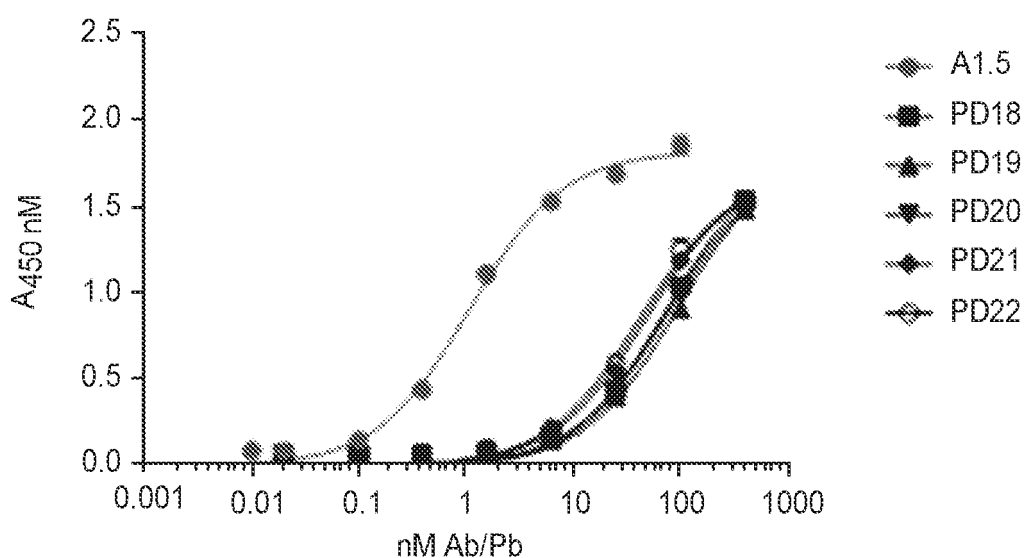
FIGURE 15, cont'd

Plate 3



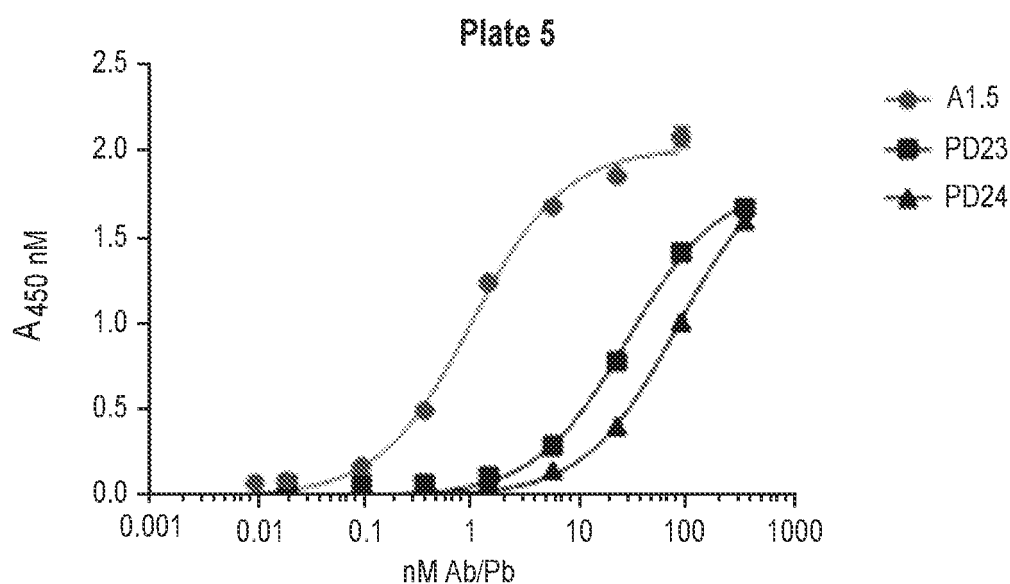
	A1.5	PD13	PD14	PD15	PD16	PD17
One site -- Specific binding		Hit constraint		Hit constraint		
Best-fit values						
Bmax	1.987	-2.200	2.162	-2.200	2.084	1.885
Kd	1.073	151.2	100.3	117.0	58.81	21.95

Plate 4



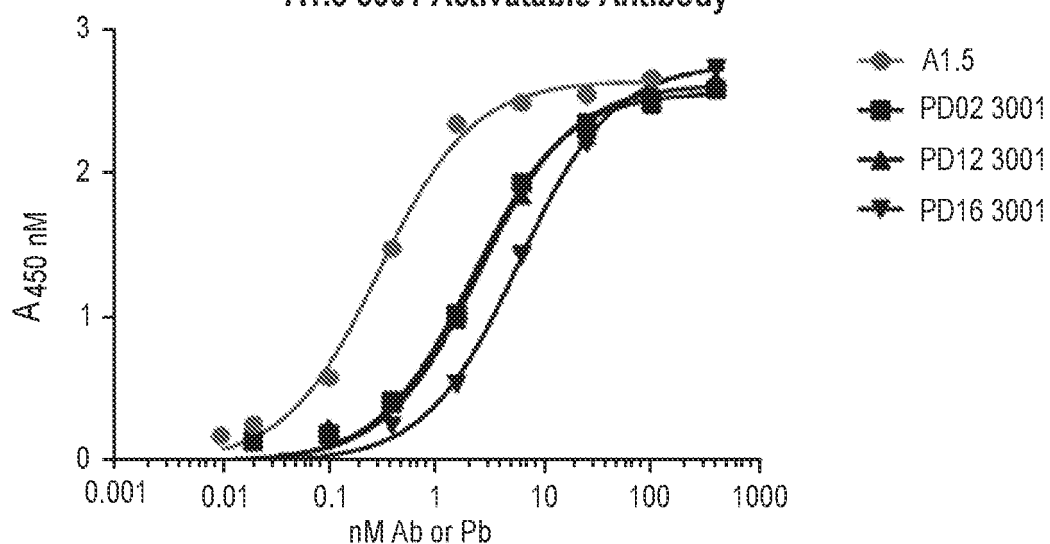
	A1.5	PD18	PD19	PD20	PD21	PD22
One site -- Specific binding						
Best-fit values						
Bmax	1.839	1.844	1.837	1.802	1.750	1.720
Kd	1.087	75.12	93.27	77.46	48.96	41.33

FIGURE 15, cont'd



	A 1.5	PD23	PD24
One site – Specific binding			
Best-fit values			
Bmax	2.049	1.847	1.993
Kd	1.086	31.37	92.42

FIGURE 16

A1.5 3001 Activatable Antibody

	A 1.5	PD02 3001	PD12 3001	PD16 3001
One site – Specific binding				
Best-fit values				
Bmax	2.660	2.580	2.632	2.777
Kd	0.2938	2.160	2.483	5.874

FIGURE 17

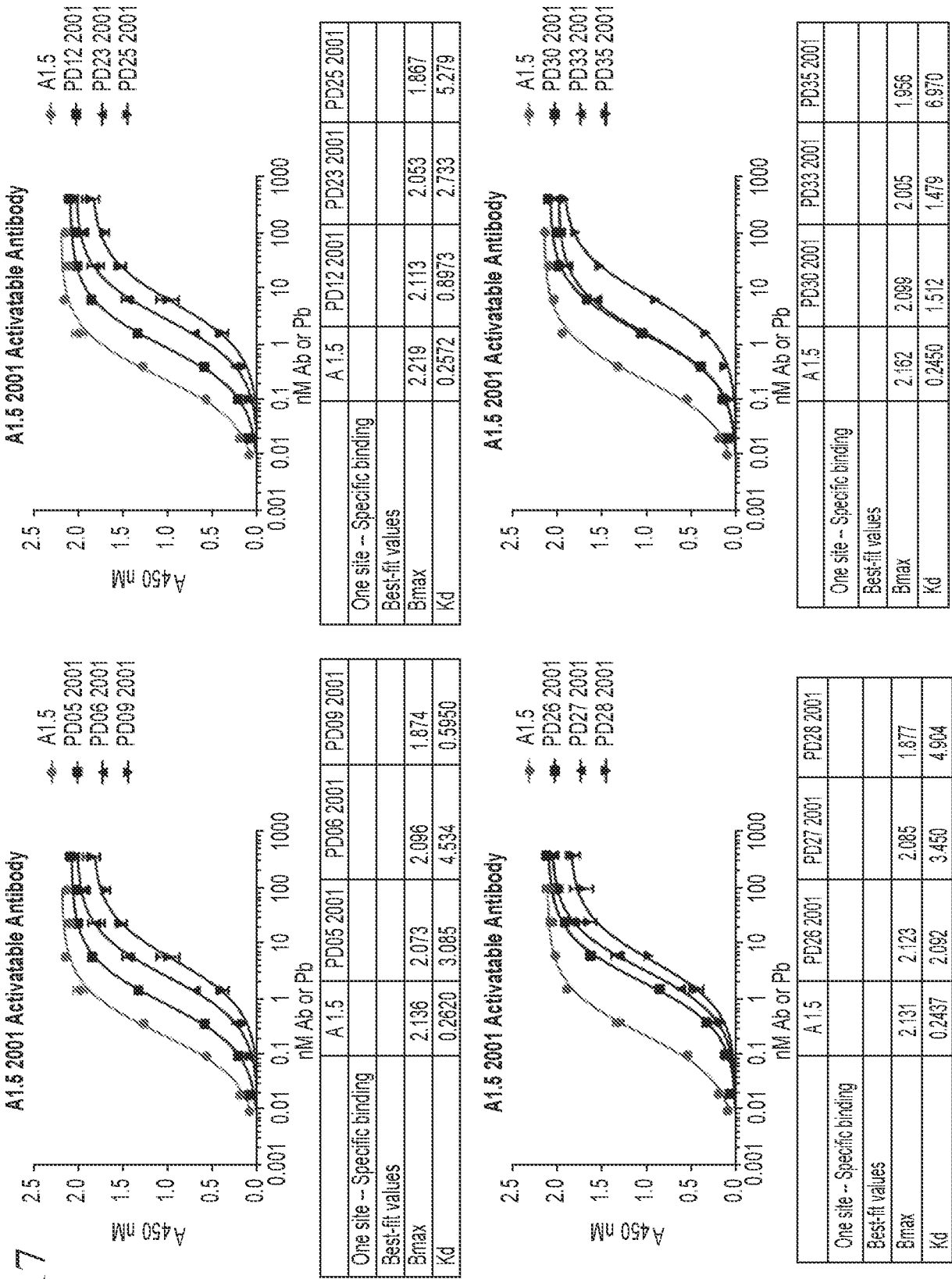
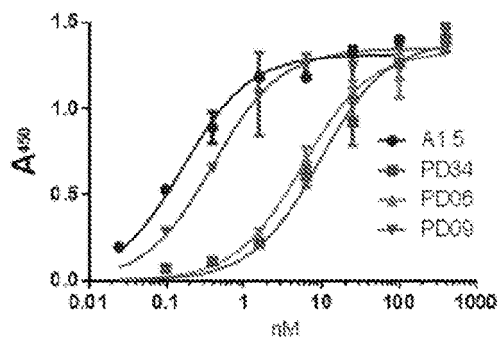
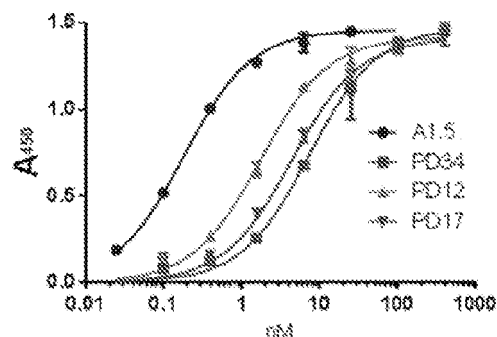


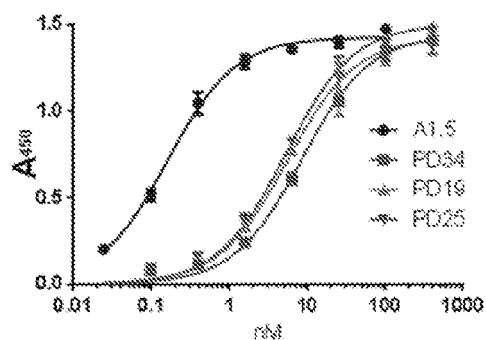
FIGURE 18



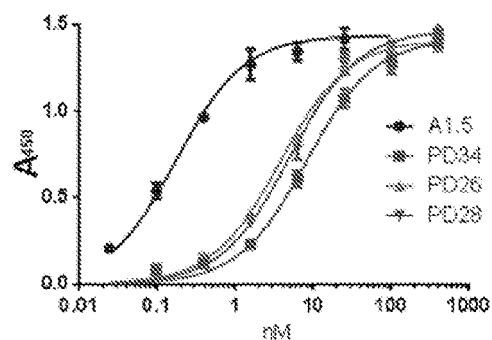
	A1.5	PD34	PD06	PD09
Kd	0.1620	9.372	5.663	0.3990



	A1.5	PD34	PD12	PD17
Kd	0.1803	7.297	1.738	4.364



	A1.5	PD34	PD19	PD25
Kd	0.1595	8.401	5.449	5.224



	A1.5	PD34	PD26	PD28
Kd	0.1762	8.123	3.754	4.970

FIGURE 19

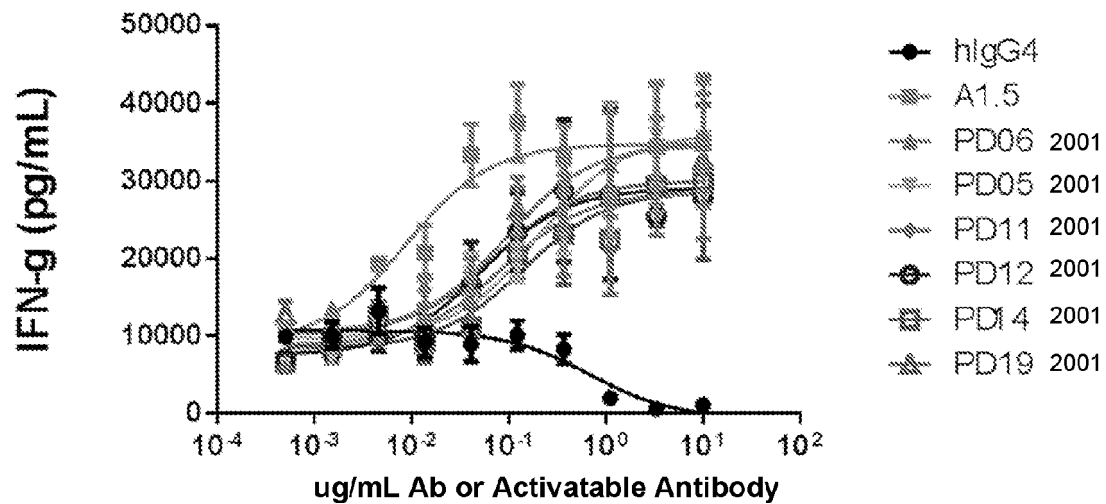
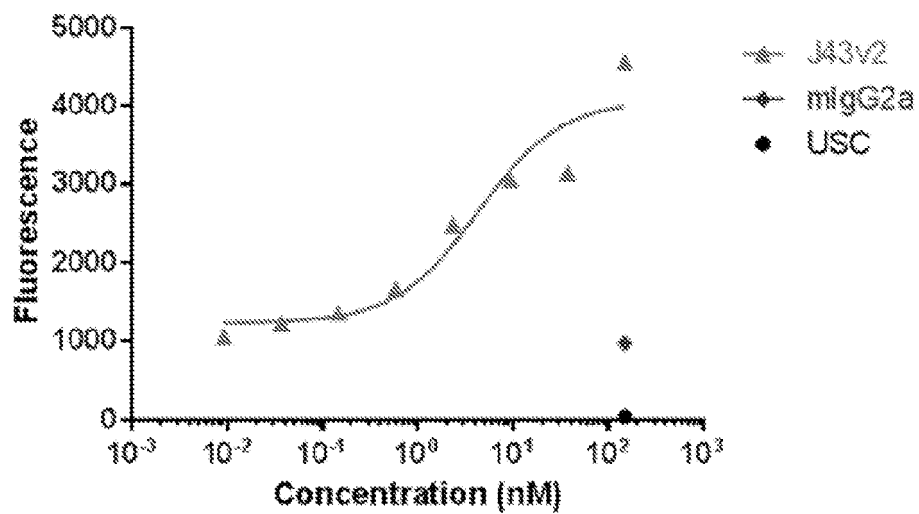


FIGURE 20A

Binding:



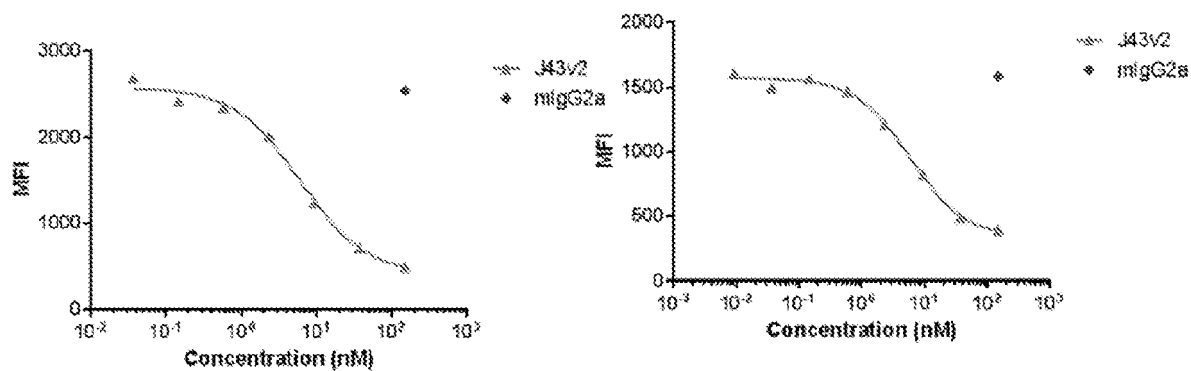
Binding EC50 (nM)

J43v2

4.308

FIGURE 20B

Blocking:



Blocking EC 50 (nM)	
J43v2	
biotin-mPD-L1	6.057
biotin-mPD-L2	5.899

FIGURE 21

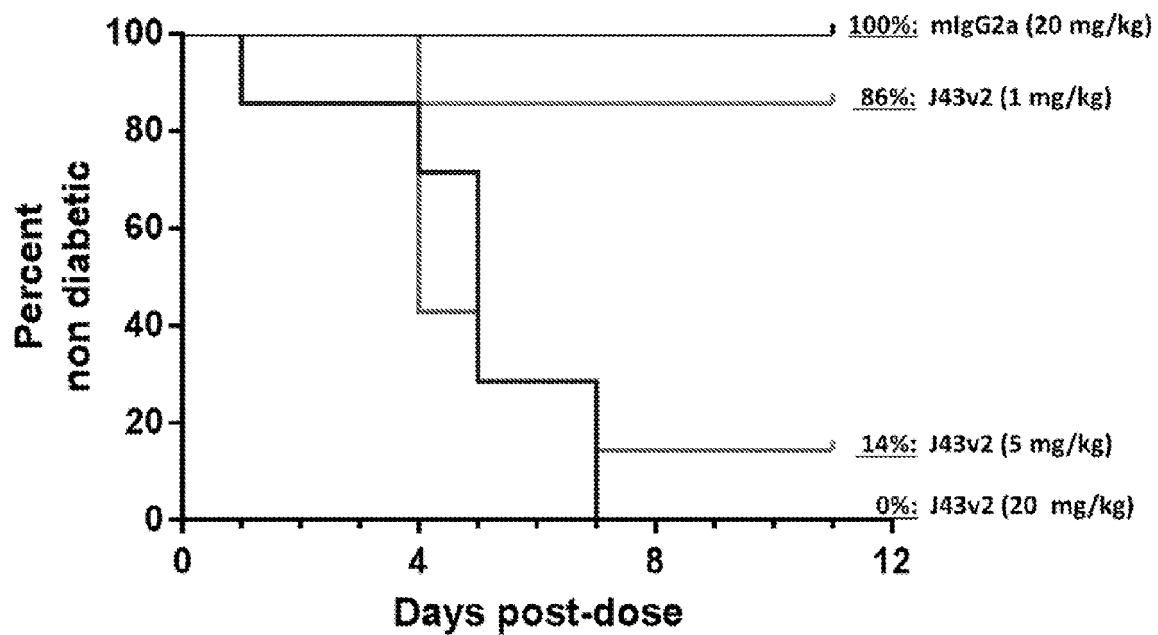


FIGURE 22

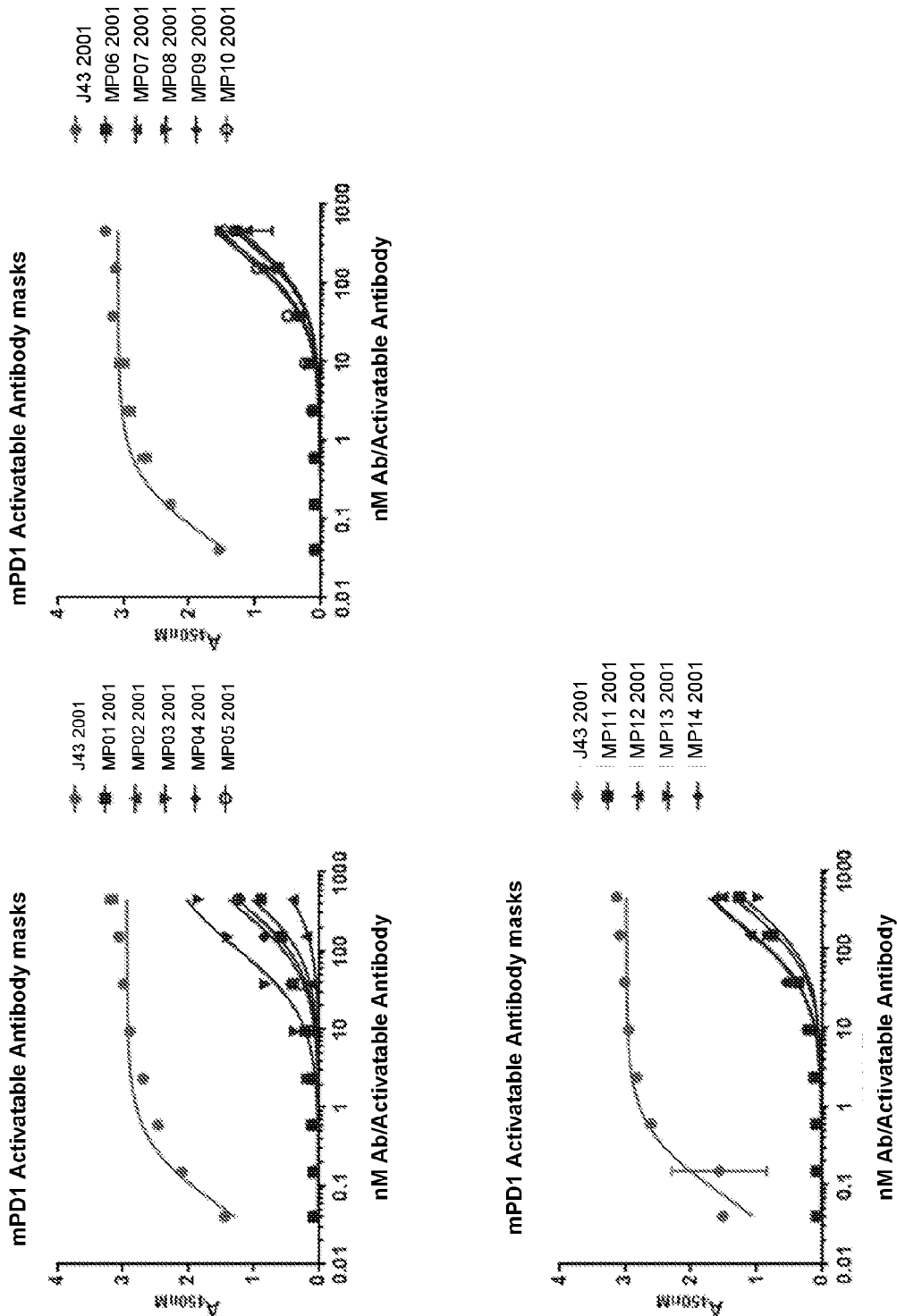
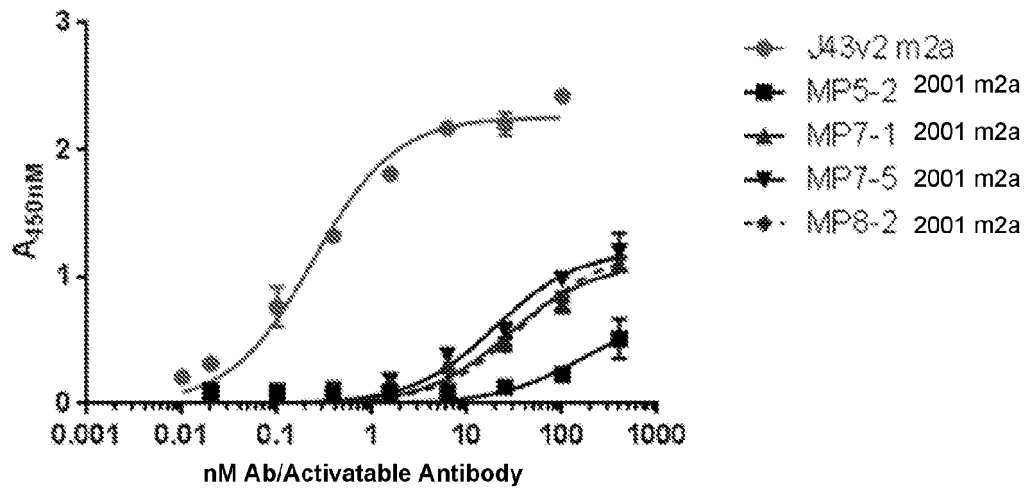


FIGURE 23



J43 MP 2001 Activatable Antibody ELISA

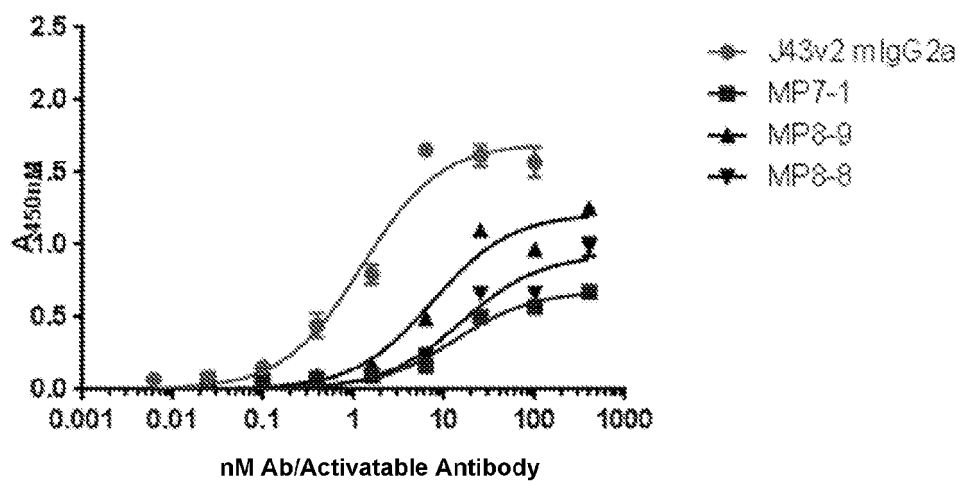


FIGURE 24

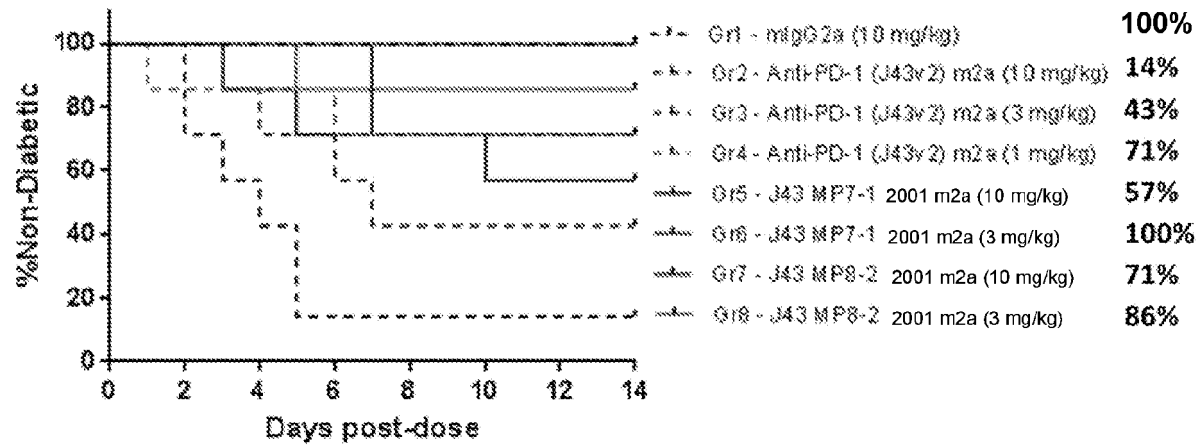


FIGURE 25A

PD34 substrate variants (2001, 2006, 2007, 2008, 2009) ME

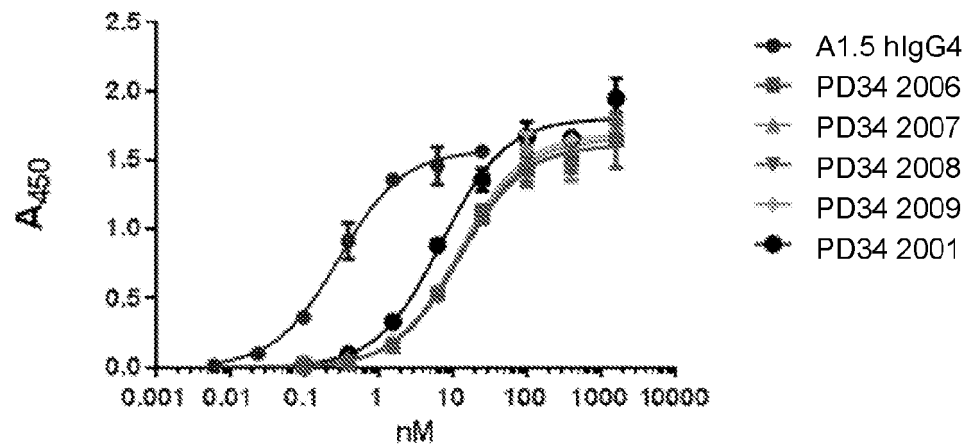


FIGURE 25B

PD34 substrate variants (2001, 2003, 2009, 2011, 2012) ME

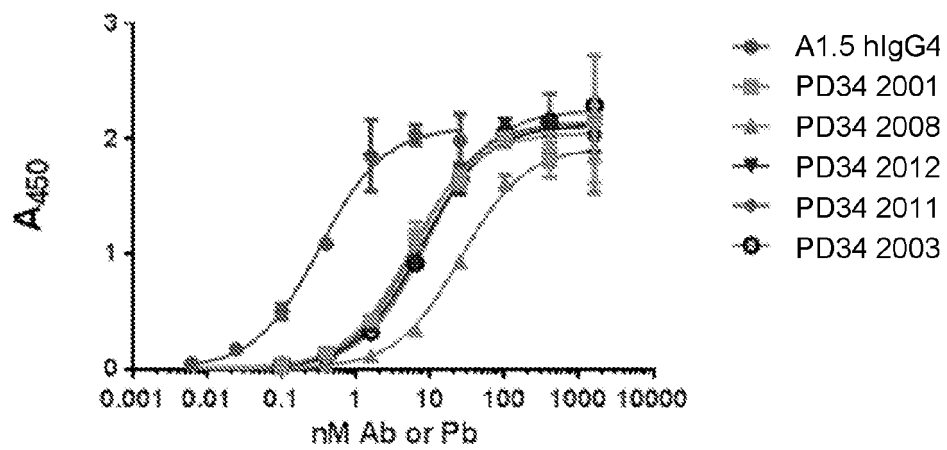


FIGURE 26A

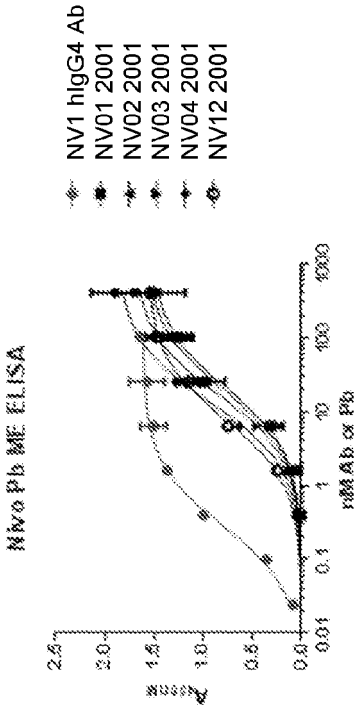


FIGURE 26B

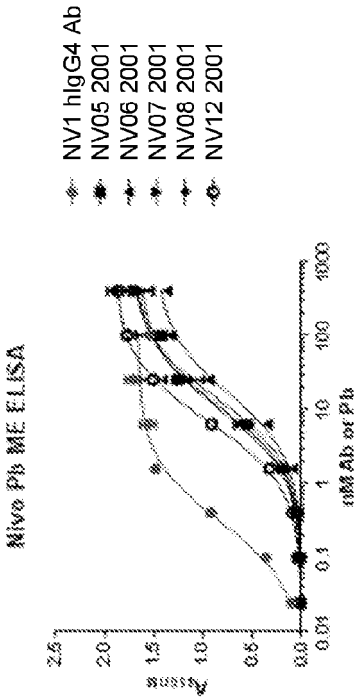


FIGURE 26C

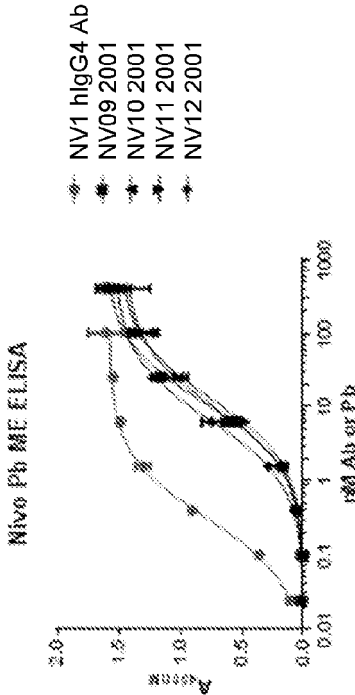


FIGURE 27B

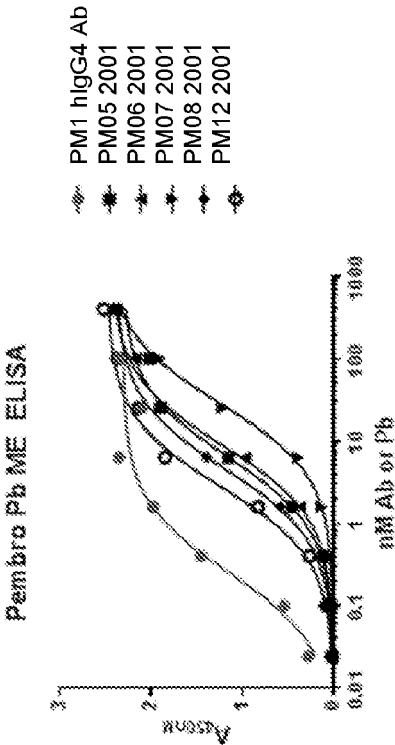


FIGURE 27A

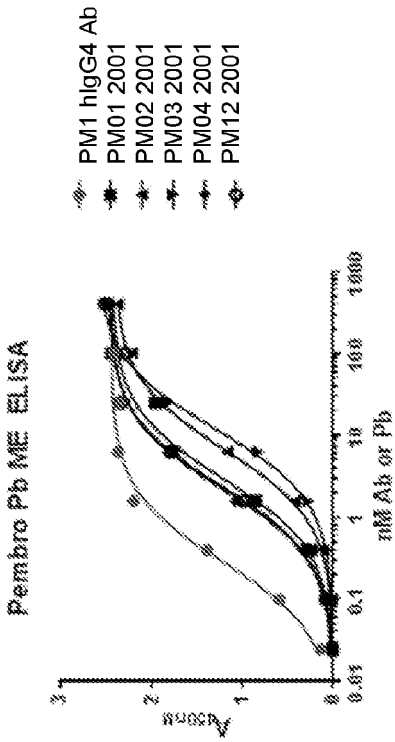


FIGURE 27C

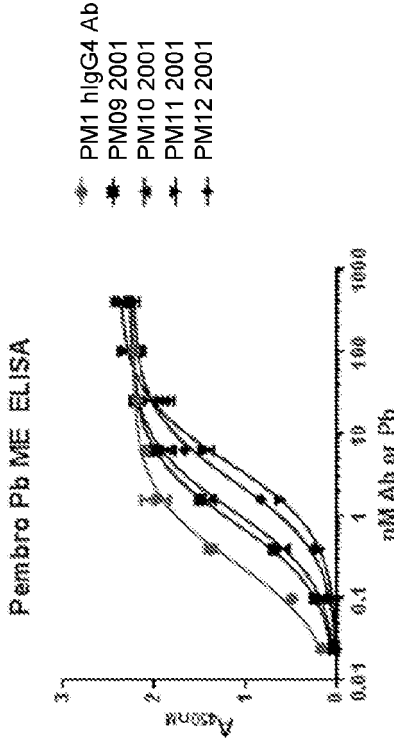


FIGURE 28

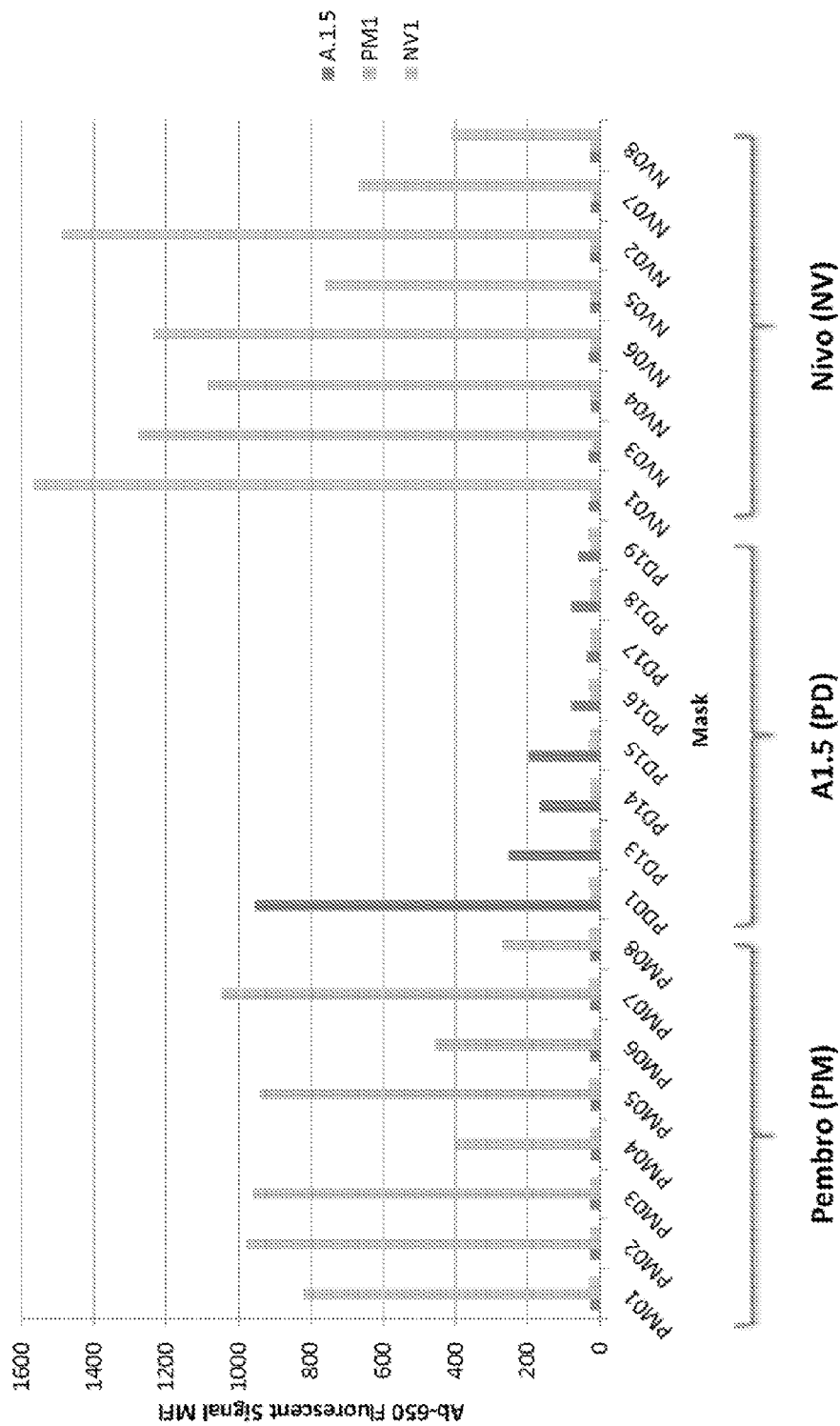


FIGURE 29A

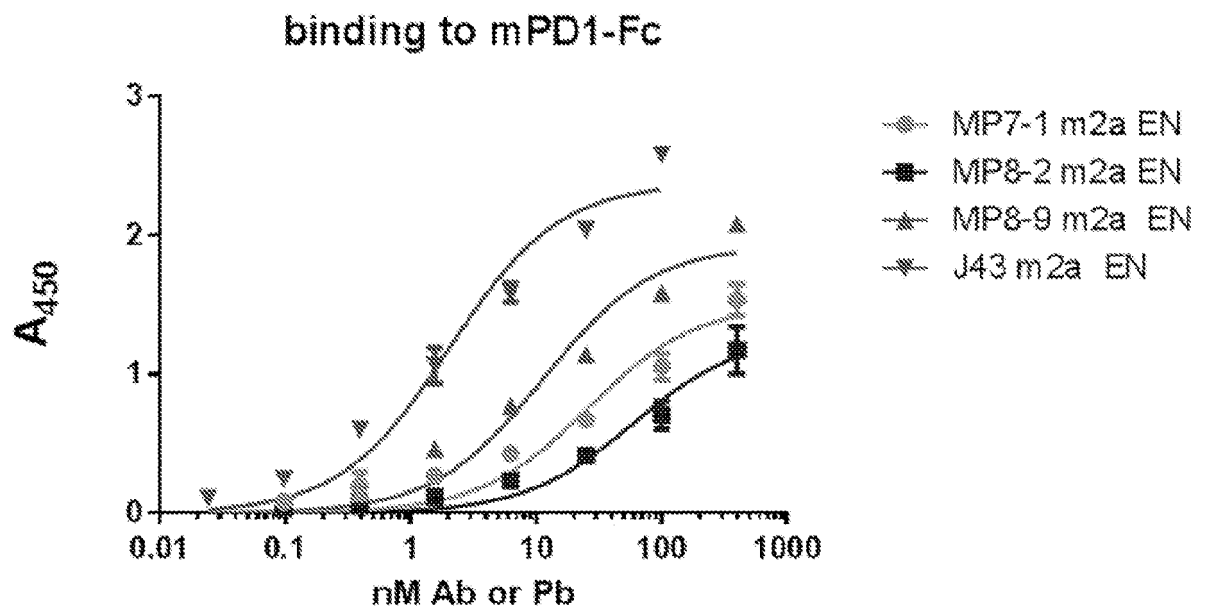


FIGURE 29B

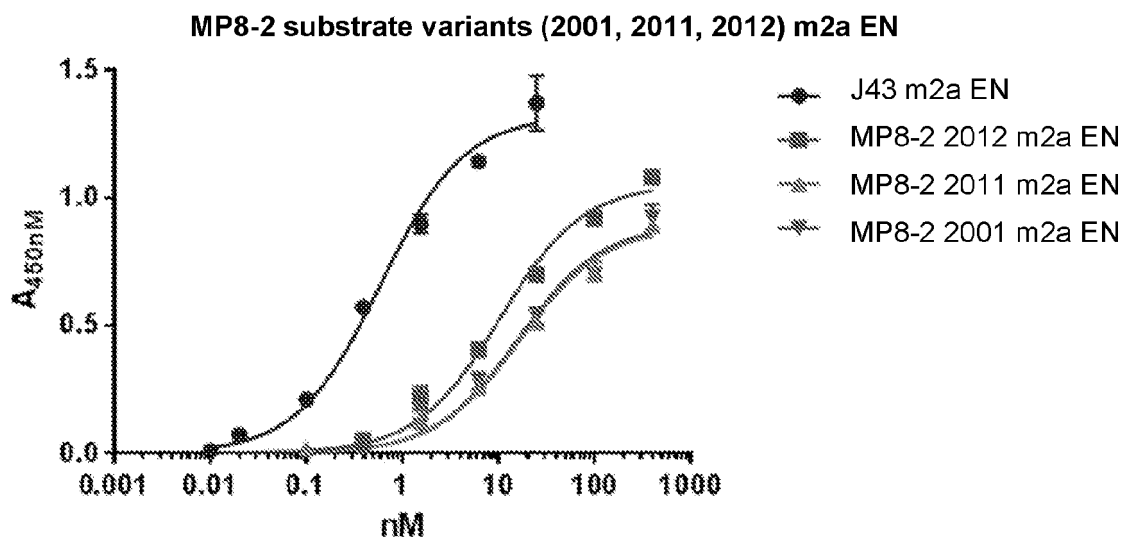


FIGURE 30

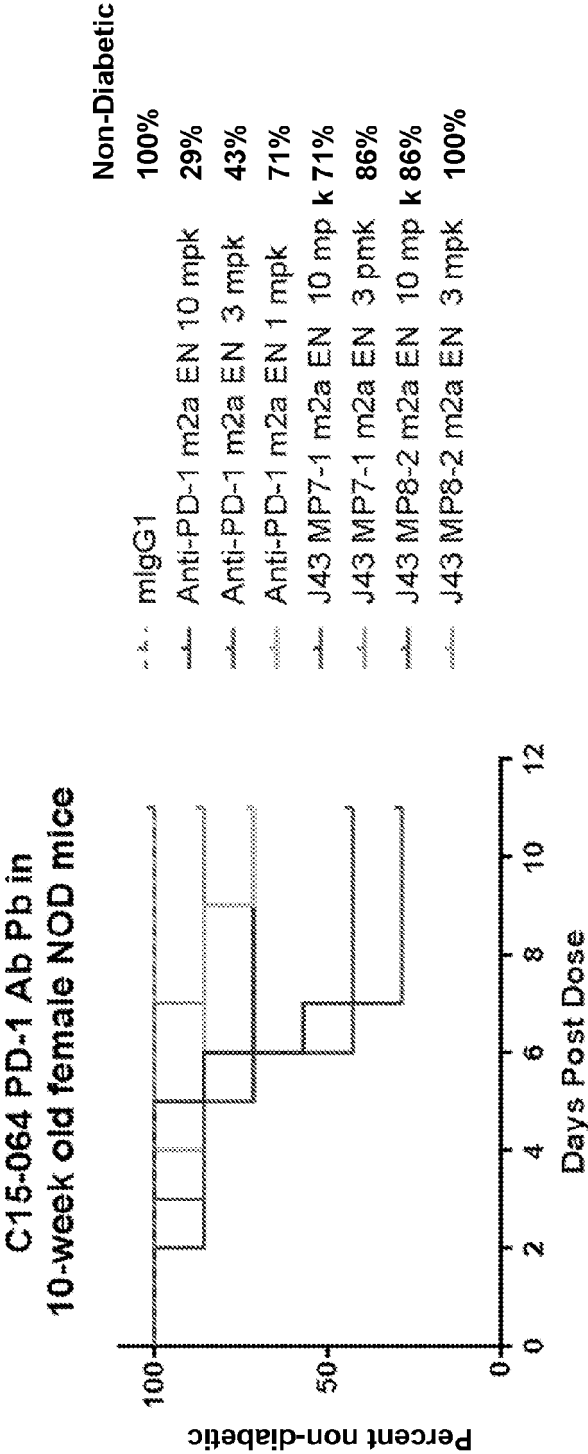


FIGURE 31A

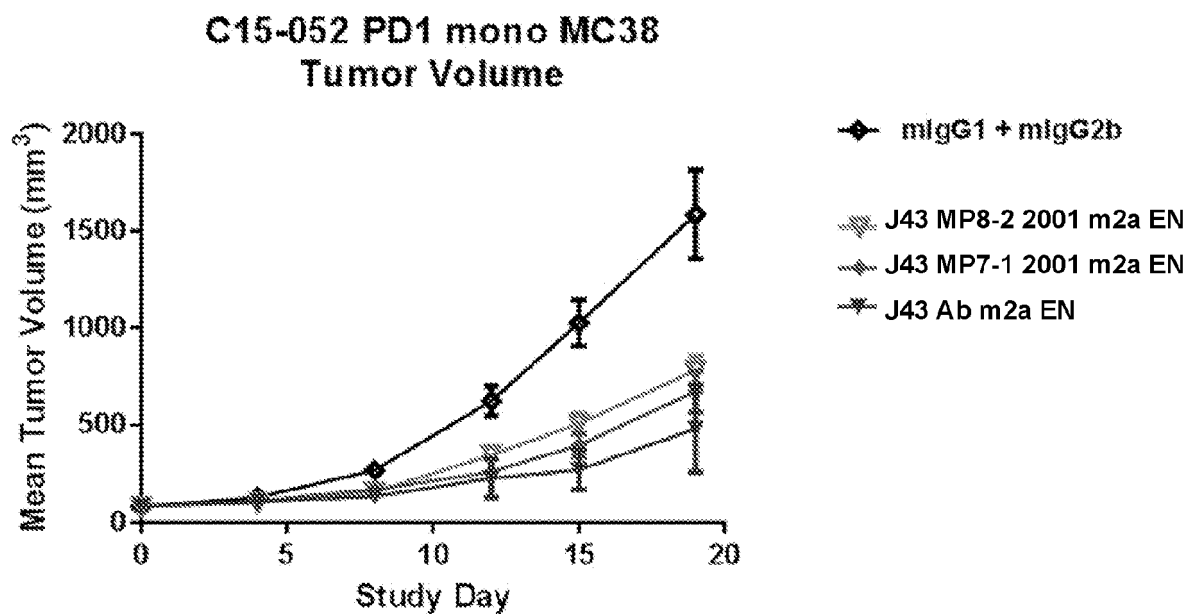


FIGURE 31B

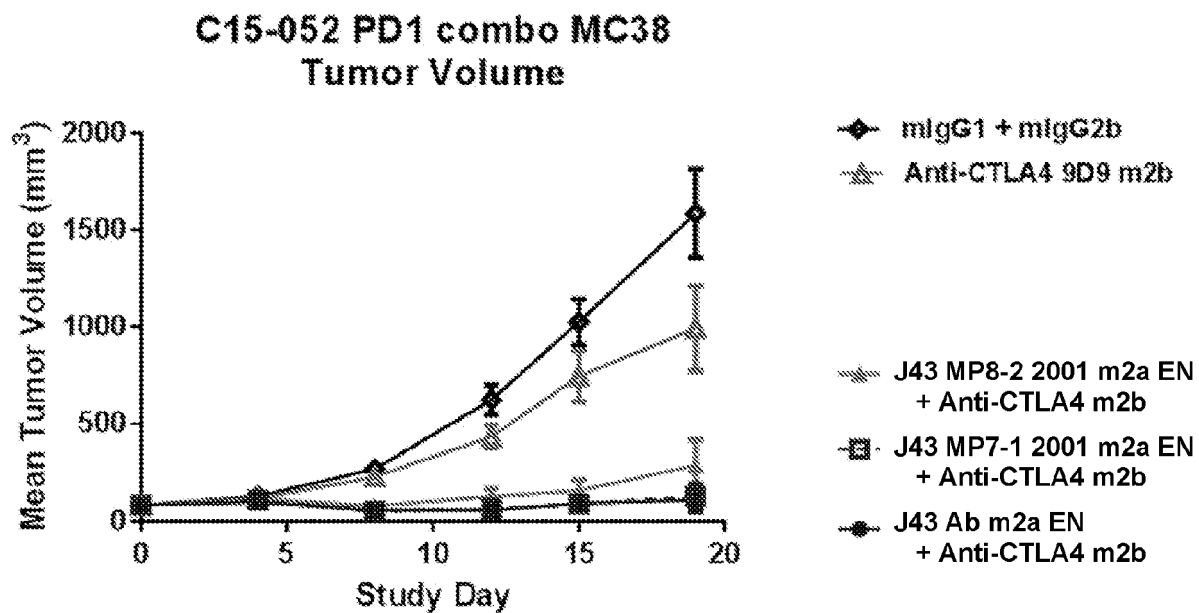


FIGURE 32A

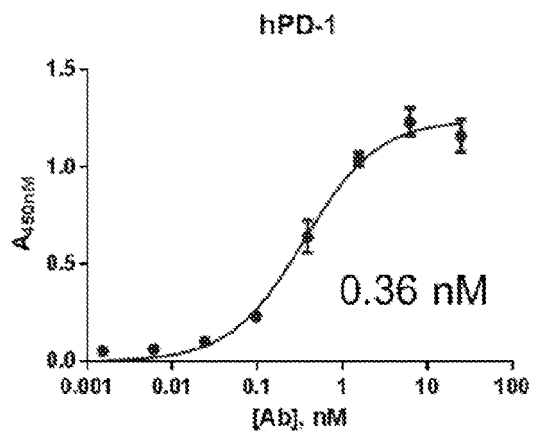


FIGURE 32B

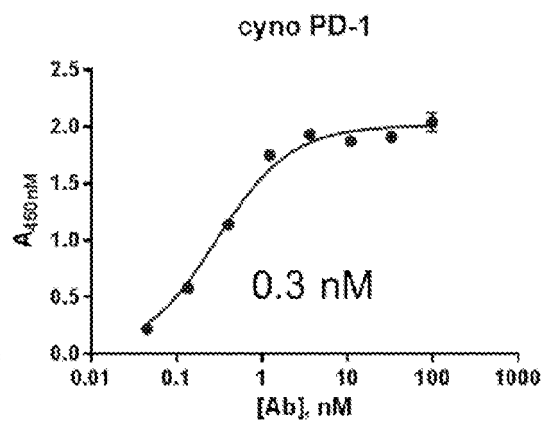


FIGURE 32C

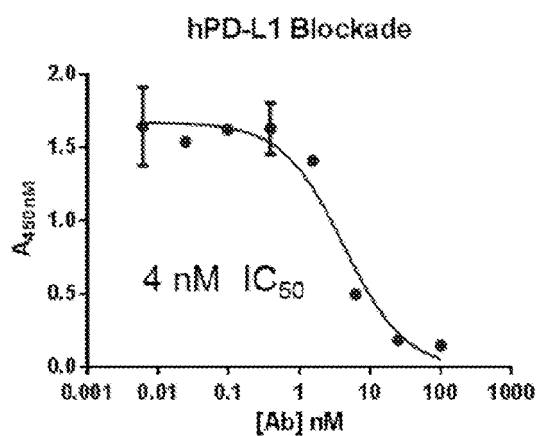


FIGURE 32D

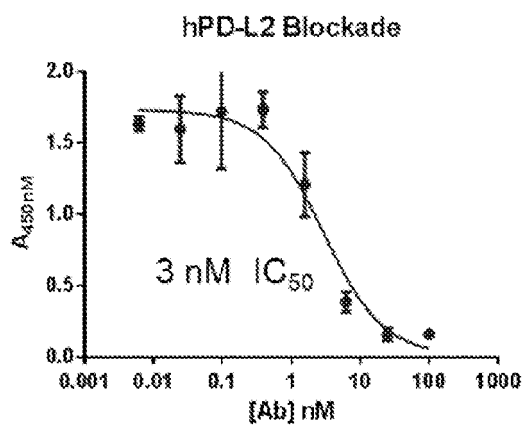


FIGURE 32E

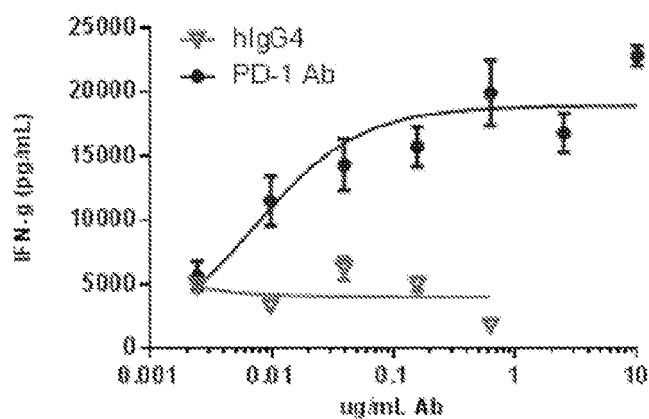


FIGURE 33A

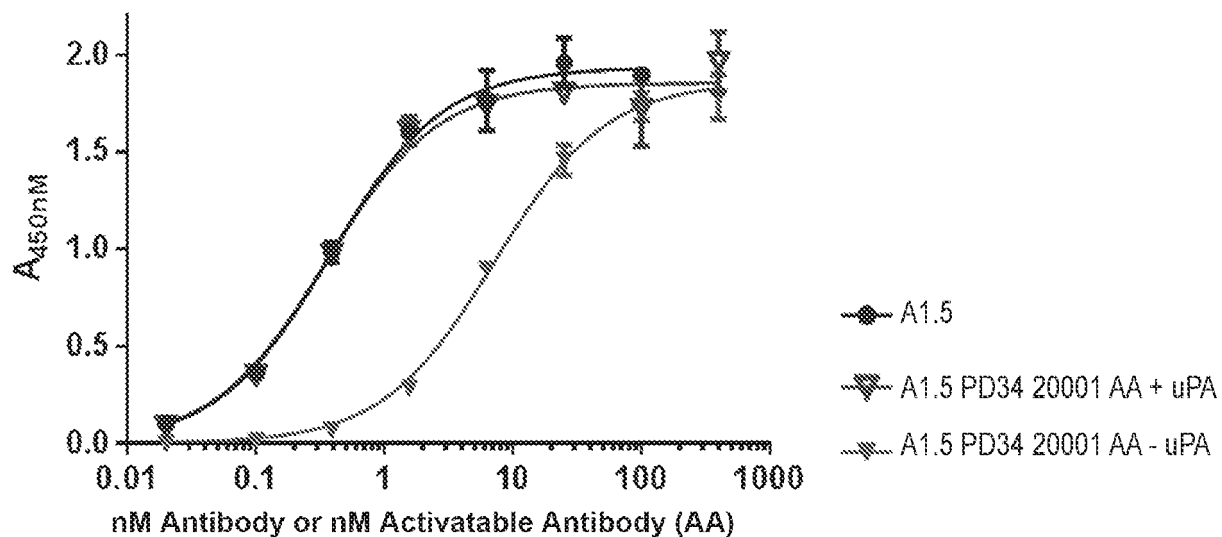


FIGURE 33B

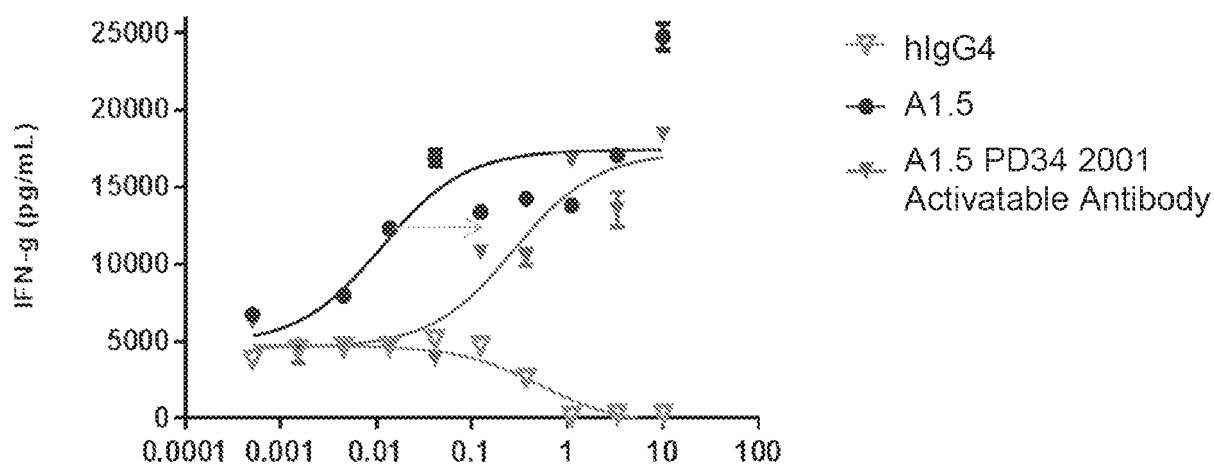
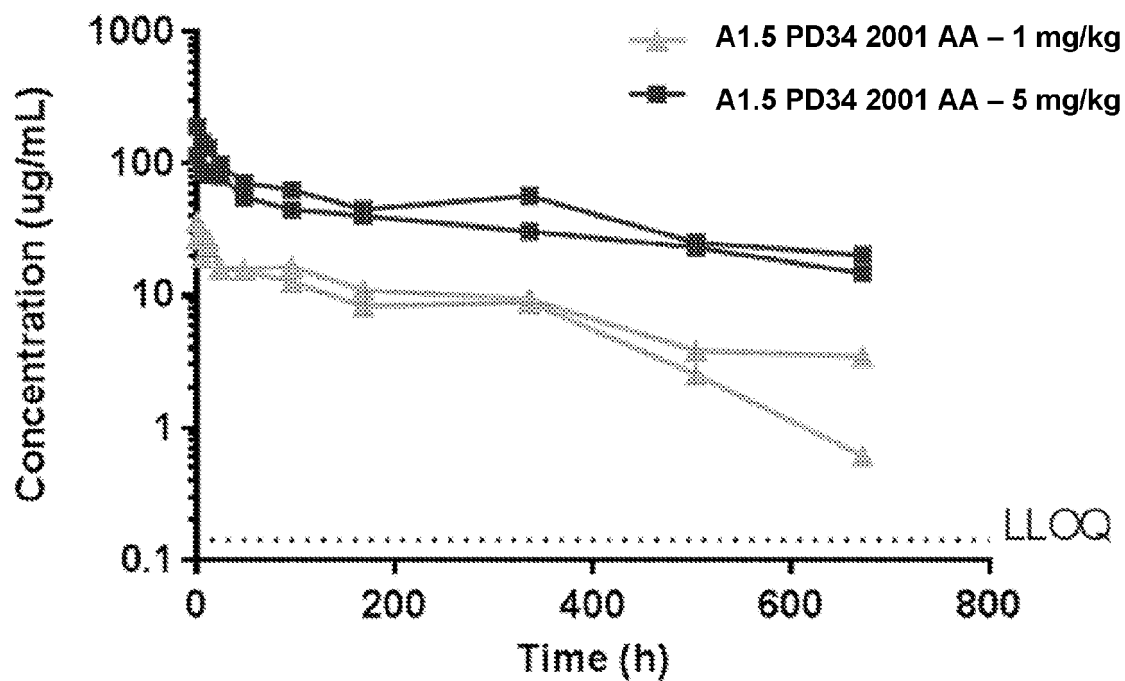


FIGURE 34



Test Article	Dose (mg/kg)	$t_{1/2}$ (days)	C_{max} (ug/mL)	AUC_{last} (day*ug/mL)
A1.5 PD34 2001 Activatable Antibody	1	8.5	32.1	228.8
	5	14.1	153.1	1132.2
A1.5 Antibody	1	8.3	34.1	184.1
	5	8.7	150.8	544.0

FIGURE 35

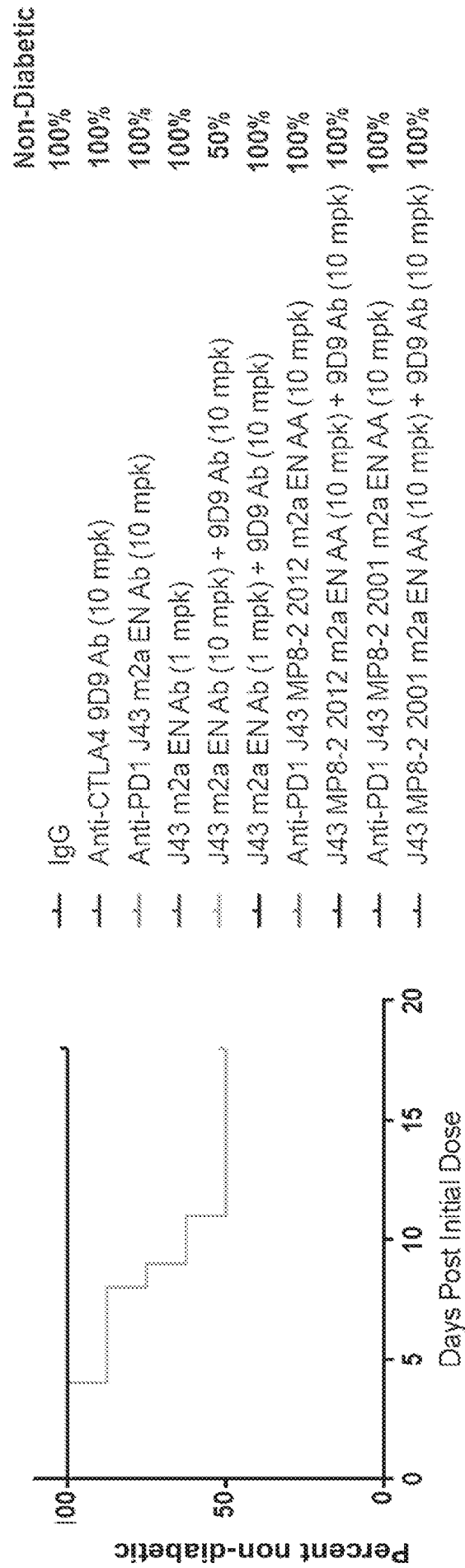
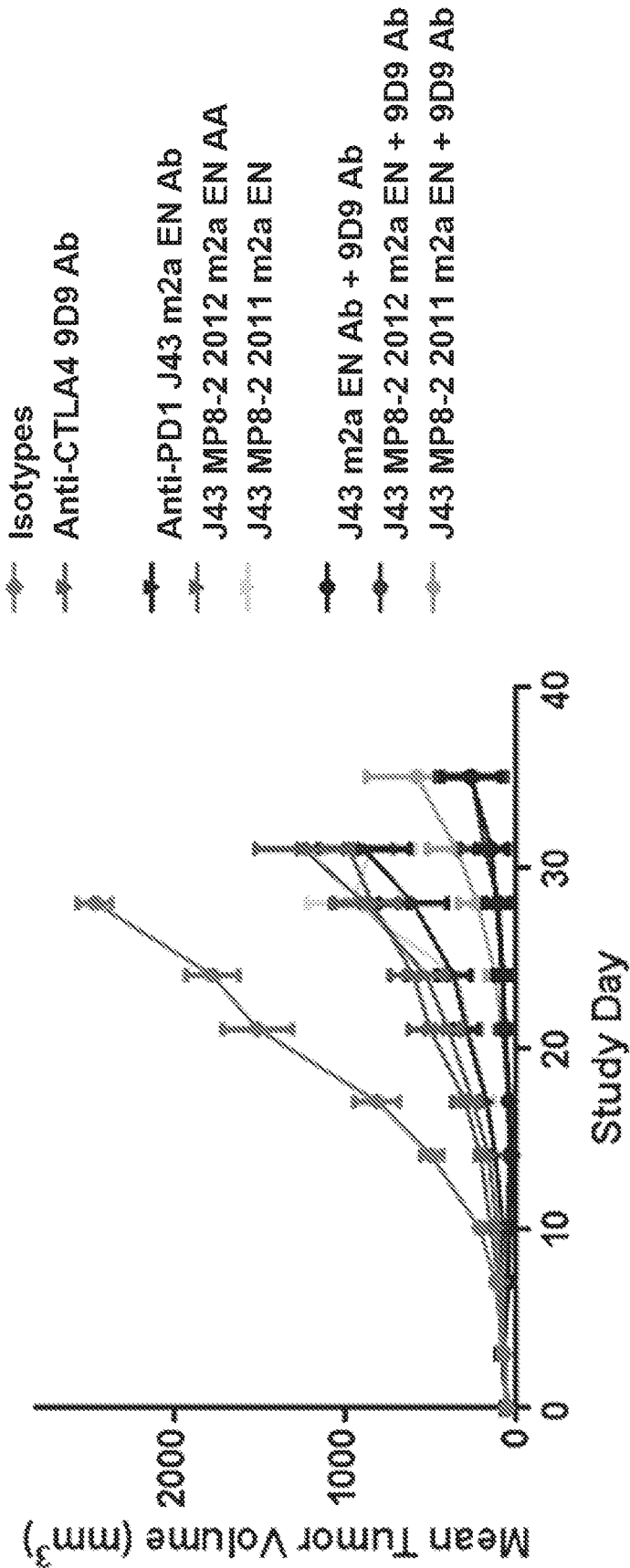


FIGURE 36



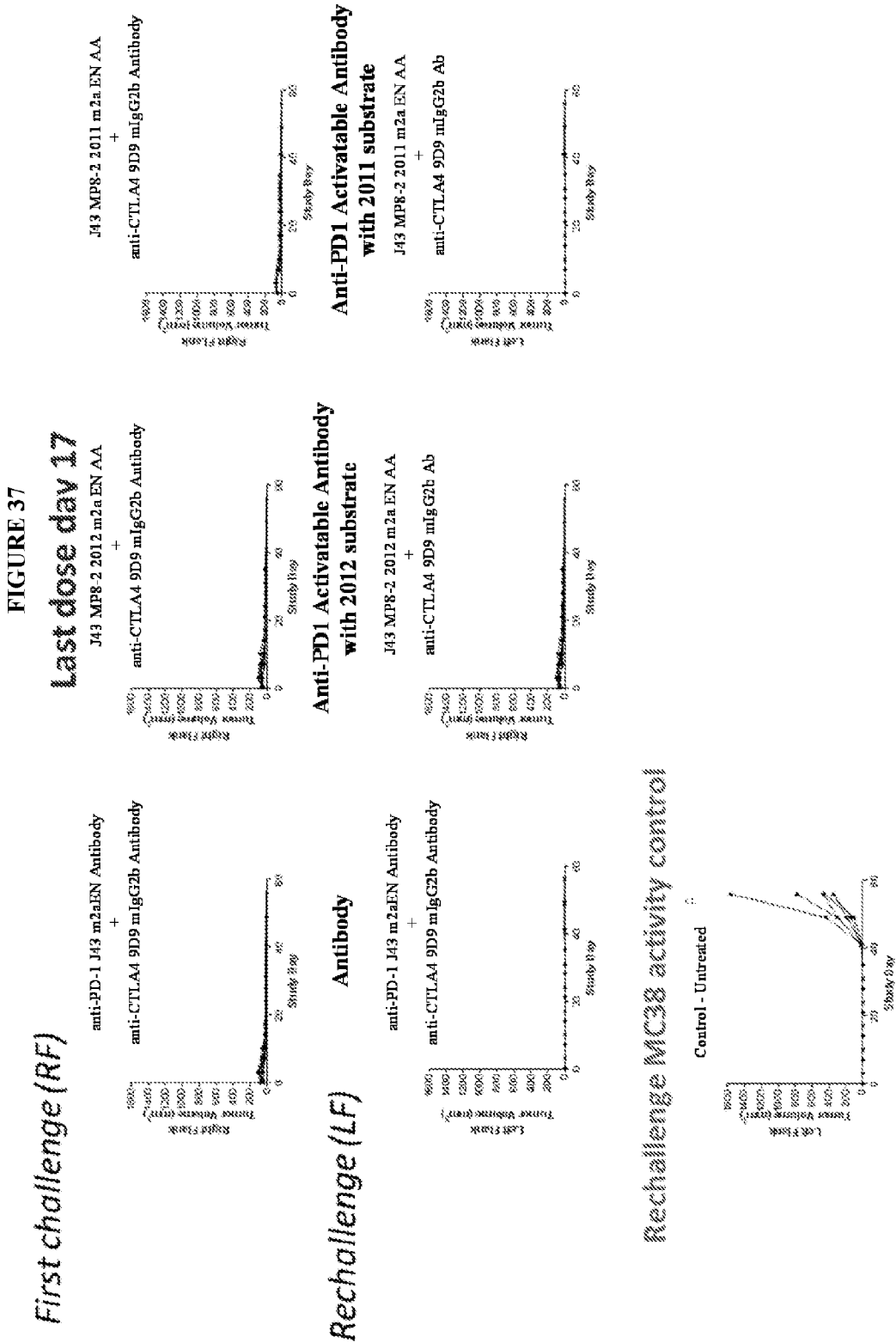


FIGURE 38

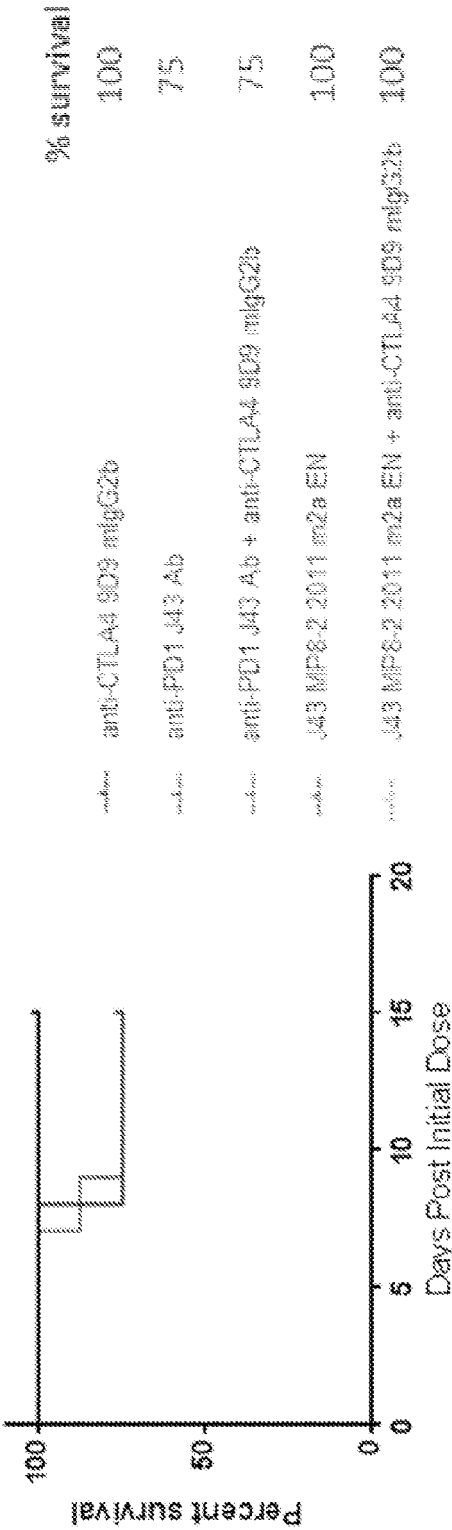


FIGURE 39

Transform of T-cell restimulation by CMV Ag and a-PD1

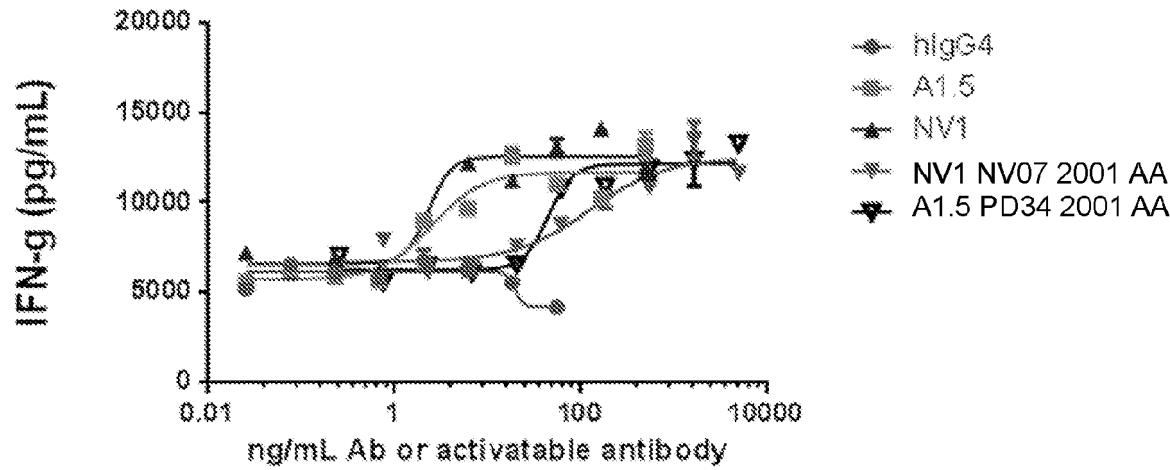


FIGURE 40

Transform of T-cell restimulation by CMV Ag and a-PD1

