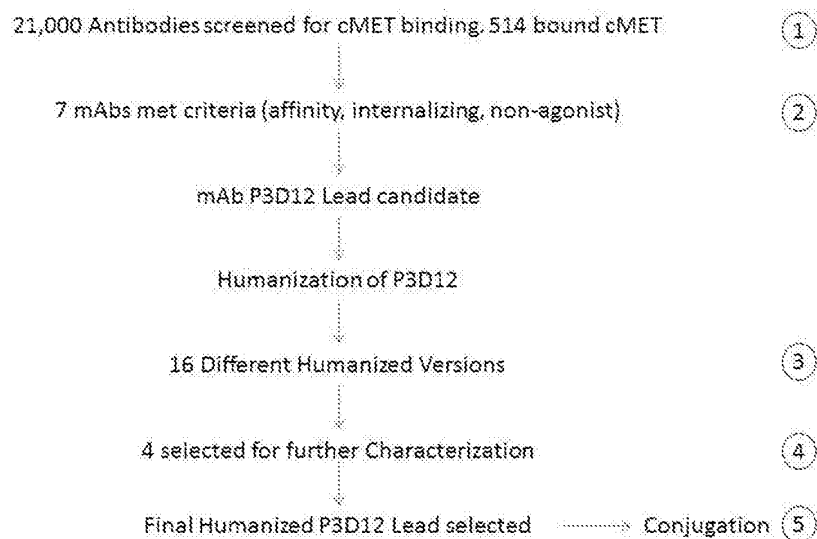
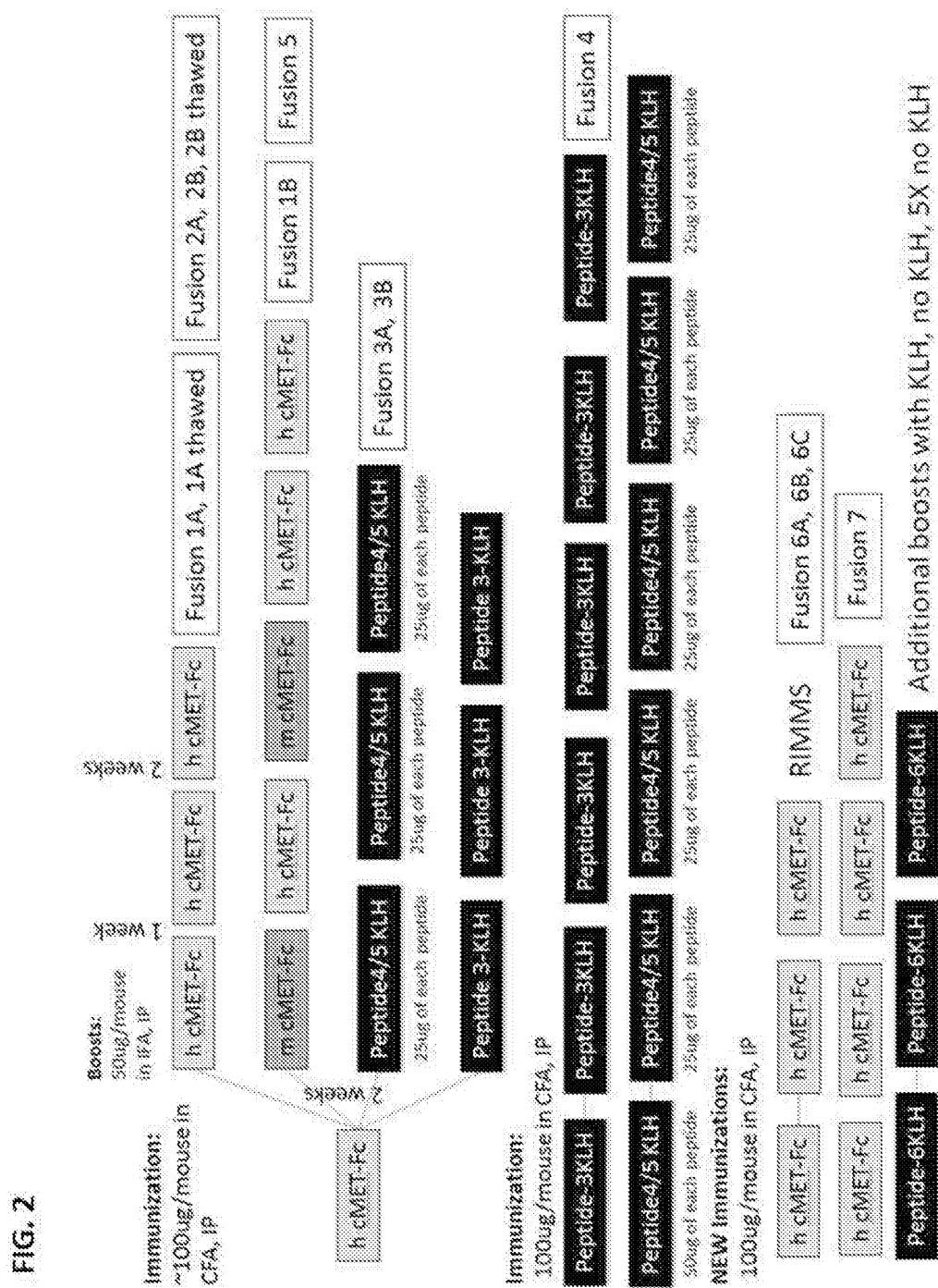


[Fig. 1]

FIG. 1

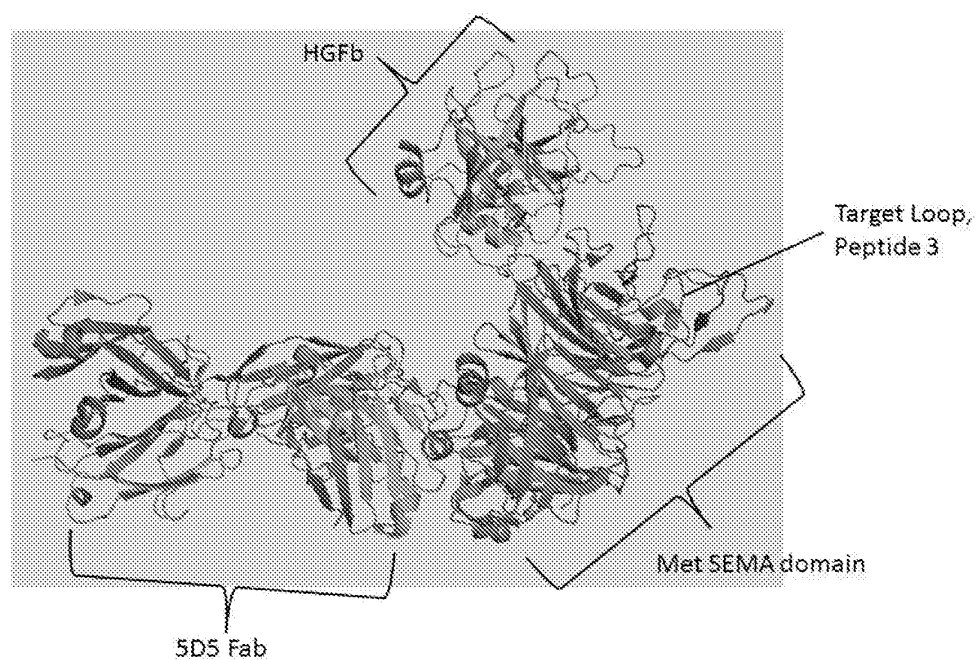
MET Antibodies Work Flow

[Fig. 2]



[Fig. 3]

FIG. 3



[Fig. 4]

2

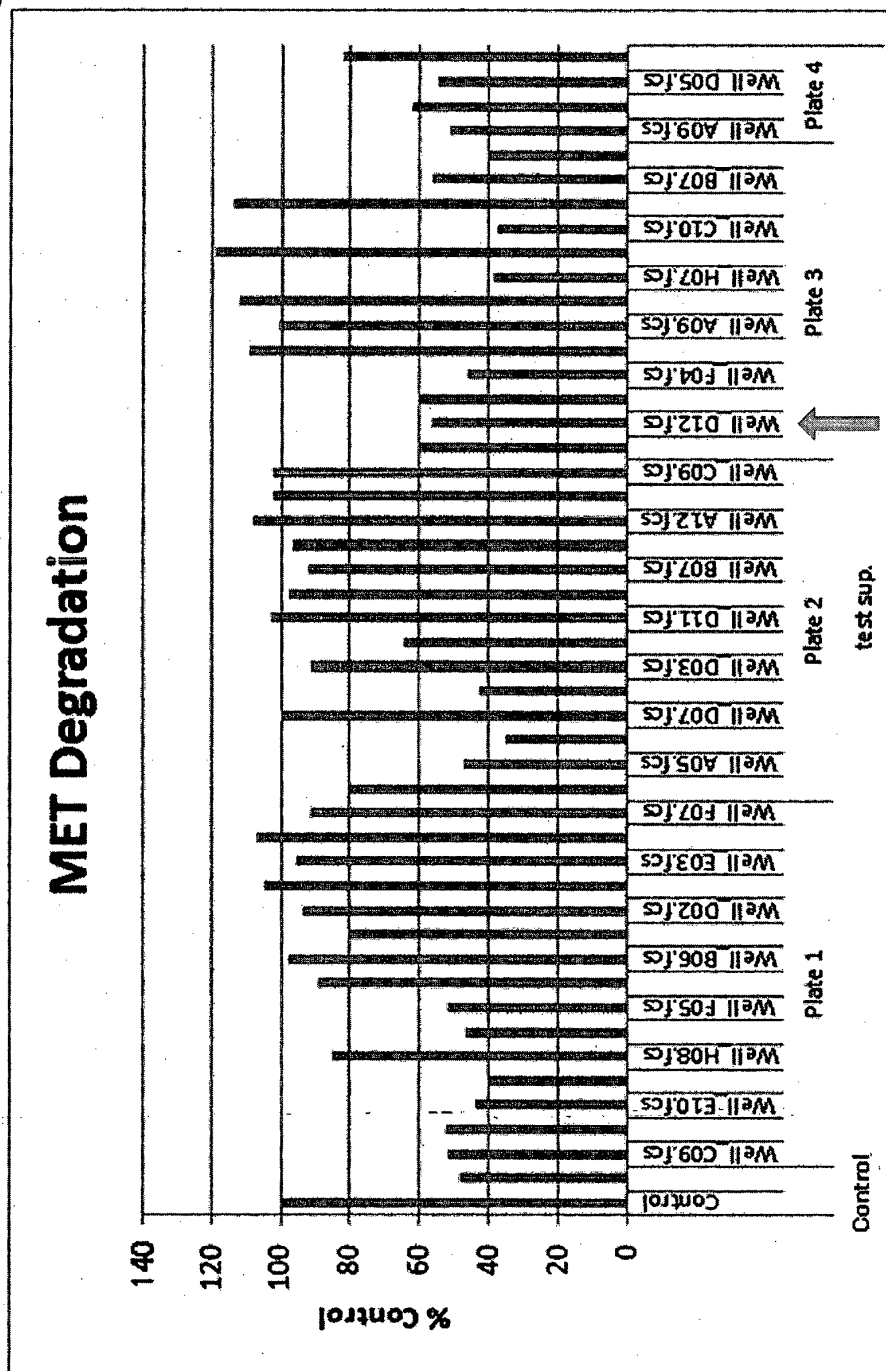
FIG. 4

Well name	Description	FACS Geom. Mean	MET BINDING ELISA OD450nm
Well_E01.fcs	negative control	103000	NA
Well_F01.fcs	negative control	116000	NA
Well_A01.fcs	positive control	83500	NA
Well_B01.fcs	positive control	81800	NA
Well_E05.fcs	hybridoma	61000	4.5319
Well_D12.fcs	hybridoma	67200	4.5803
Well_E02.fcs	hybridoma	69800	4.596
Well_F04.fcs	hybridoma	77300	3.0301
Well_B10.fcs	hybridoma	97200	0.0211
Well_H05.fcs	hybridoma	97600	6.1209
Well_F05.fcs	hybridoma	97800	-0.0005
Well_G10.fcs	hybridoma	98800	0.0078
Well_E07.fcs	hybridoma	98900	0.0011
Well_H10.fcs	hybridoma	98900	1.7238
Well_E06.fcs	hybridoma	99300	5.7925
Well_F10.fcs	hybridoma	99700	0.0185
Well_F07.fcs	hybridoma	99800	0.235

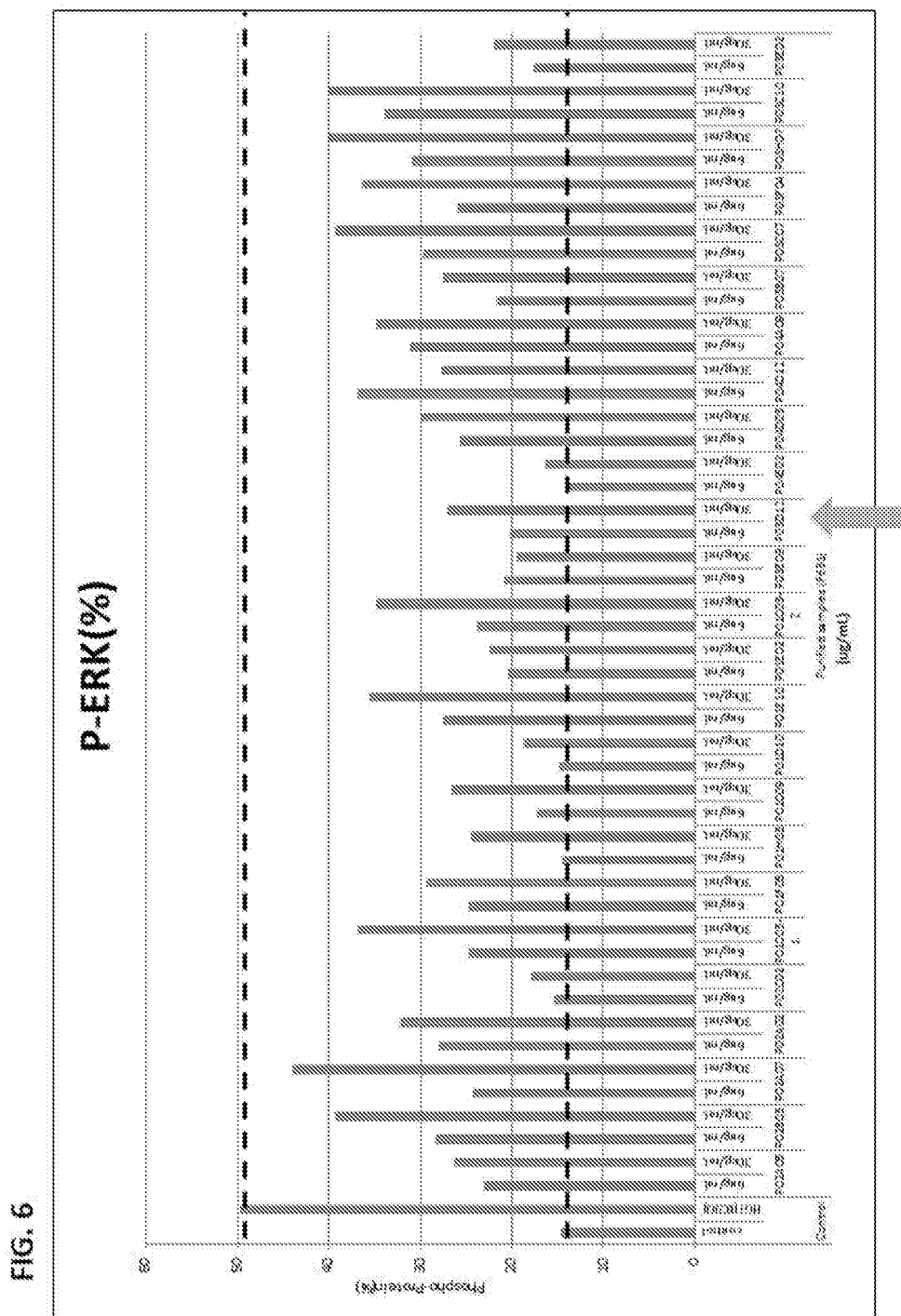
[Fig. 5]

2

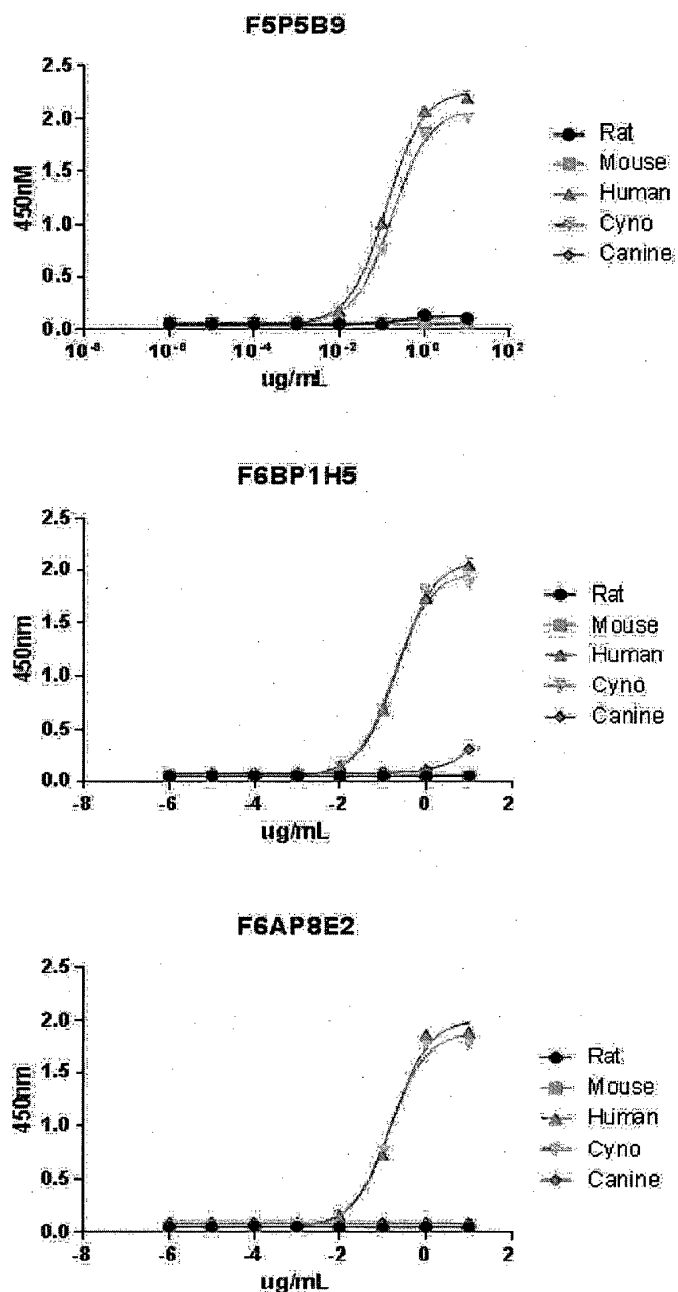
FIG. 5

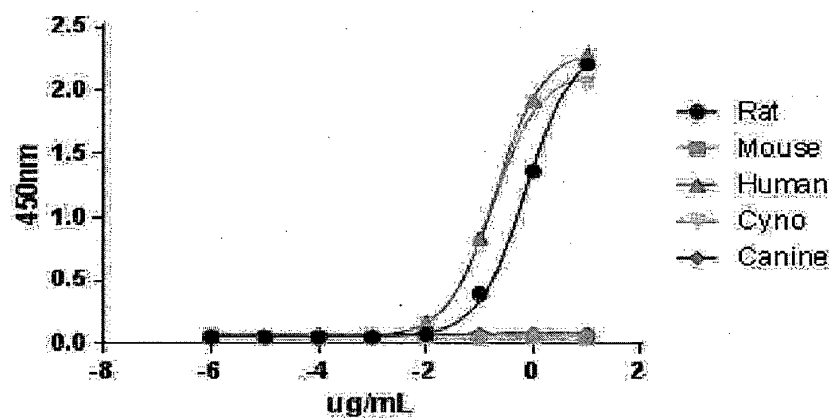
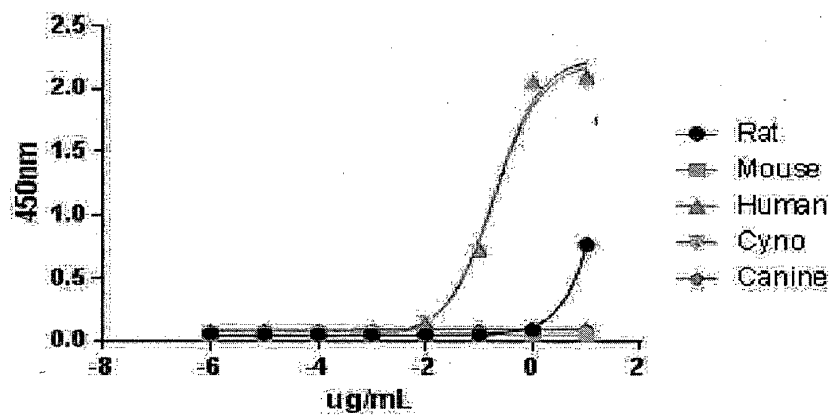
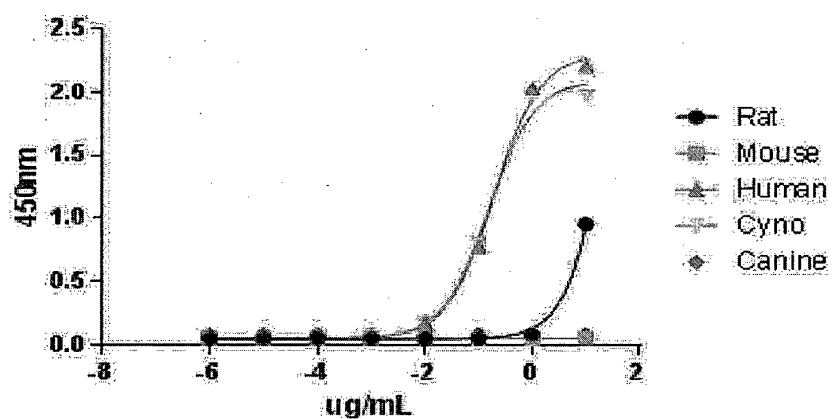


[Fig. 6]



[Fig. 7]

FIG. 7 Anti-Met antibodies species cross-reactivity

F6B1P3D12**F6B1P1E2****F6BP2D4**

[Fig. 8]

Sequences of top 9 murine clones. Two clones were duplicated giving 7 unique sequences

FIG. 8A

Light Chain		SEQ ID NO:	117	100	90	80	70	60	50	40	30	20	10	1
F5_P5_B9_L (1)	--DVLMTQPLSLVSLGGQASISCRSSQ--YIVHGNT--VLEWYLRFGQSPKLLIYKSNRFPDPSGSGGDTFLTKISVRAEDLGVYICFGSHHPYTFGGGKLEIKR	37												
F6A_P8_E2_L (1)	--DIVMTQSPKSMNSMGERTLSCKAS-----ENVGT--YVSMYQKQEDQSPKLLIYGASNRVTGVPDRFTGSGSATDFTLTSSVQAEDLADYHCQSYSPYELTFGAGTKLEIKR	38												
F6A_P12_F12_L (1)	--DIVMTQSPSLAVSVGEKVTMSQSLIYASINQRN--YLAWYQQIPGQSPKLLIYMASTRESGVPPDRFTGSGGDTFLTKISVRAEDLADYICQYQYTYPLTFGAGTKLEIKR	39												
F6B_P1_H5_L (1)	RCDIQMTQSPASLSVSGEFTVITCRASENIYN-----FLAWYLRQKQSPKLLIYVAAVNLADGVPSRFGSGSGGTQYSLKINLSQSEDFGSSYVCQHFPGTFYTFGGGKLEIKR	40												
F6B1_P3_D12_L (1)	--QIVLTQSPAIMSASGEGEKVTILTCASASSVTS-----NLYWYQKQFGSPKLLIYSTNLASGVPARFSGSGGTSLTISMEADAASYFCHQWSSYPPTFGSGTKLEIKR	41												
F6B1_P3_E9_L (1)	--QIVLTQSPAIMSASGEGEKVTILTCASASSVTS-----NLYWYQKQFGSPKLLIYSTNLASGVPARFSGSGGTSLTISMEADAASYFCHQWSSYPPTFGSGTKLEIKR	41												
F6B_P2_D4_L (1)	--QIVLTQSPAIMSASGEGEKVTILTCASASSVSS-----NLYWYQKQFGSPKLLIYSTNLASGVPARFSGSGGTSLTISMEADAASYFCHQWSSYPPTFGSGTKLEIKR	42												
F6B1_P1_E2_L (1)	--QIVLTQSPAIMSASGEGEKVTILTCASASSVSS-----NLYWYQKQFGSPKLLIYSTNLASGVPARFSGSGGTSLTISMEADAASYFCHQWSSYPPTFGSGTKLEIKR	43												
F6B_P3_E2_L (1)	--QIVLTQSPAIMSASGEGEKVTILTCASASSVSS-----NLYWYQKQFGSPKLLIYSTNLASGVPARFSGSGGTSLTISMEADAASYFCHQWSSYPPTFGSGTKLEIKR	43												

LC F6B1P1E2=F6BP3E2

LC F6B1P3D12=F6B1P3E9

FIG. 8B

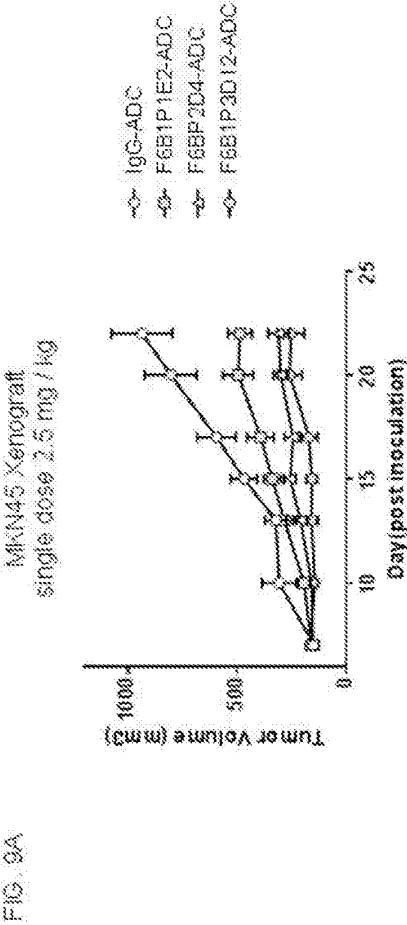
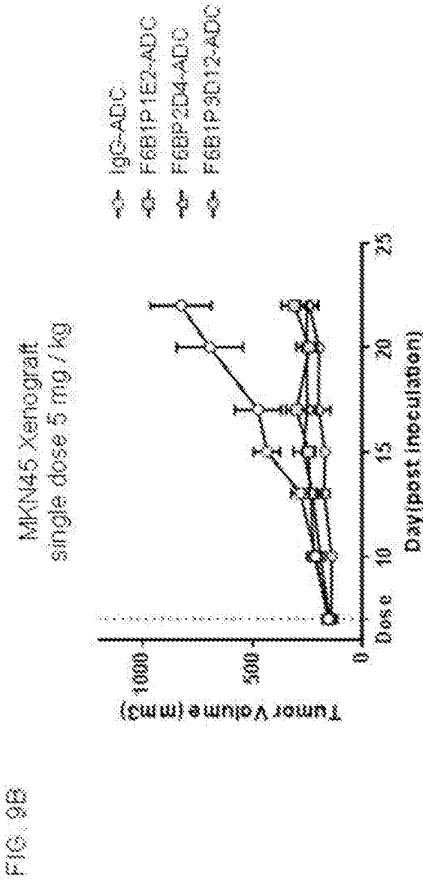
Heavy Chain

SEQ ID NO:	131	120	110	100	90	80	70	60	50	40	30	20	10	1
F5_P5_B9_H (1)	---	QVQLKESGRLVAPSGQSLISLTCTVSGFSLPTNIGVNWVRPQPKGLEWLGILWG---	GGDTPNSALKSRSLISKDNKSKQVFLKQETNLSIQDDTARYICAR---	DYGFQDWGQGTTLTVSS	94									
F6A_P12_F12_H (1)	---	EVLKVESGGGLVQPGGSLRLSCATSGFTFDYINMSWRPQPKALEWLGIFINNKANGYITTKYSASVKGRFTISRDNQS---	LLYLQNTLRAEDSATYICSK---	DRGYFDYWGQGTTLTVSS	95									
F6A_P8_E2_H (1)	VNSEYQIQSQGAELVRPGALVKLSCKASGFNINDYFHHWYKQRPQGLEWIGWIDP--ENGNTIYDFKQKASTADTSSN--TAYLQLSLTSEDTAVYICARGENYLRSEYIYAMDYWGQGTSTVSS	96												
F6B_P1_H5_H (1)	VISEVILQQSGPELVKPGASVKIPCKASGYTFDYNMDWKQSHKSLIEWIGDINP--NNGGTIYNQKFKGKATLTVDKSSS--TAYMELRLSITSEDTAVYICARGDYIGSSRIYIYAMDYWGQGTSTVSS	97												
F6B1_P3_D12_H_79_13 (1)	---	QVQLQQSGAELAKPGASVRMSCRASGYTFTSYHHWVKQRPQGLEWIGYIKP--STDNTEYNQKFKDKATLTADKSSS--TAYMQLSLTSEDSAVYICAR---	SYGN---	YPLMDYWGQGTSTVSS	98									
F6B1P3E9_H (1)	---	QVQLQQSGAELAKPGASVRMSCRASGYTFTSYHHWVKQRPQGLEWIGYIKP--STDNTEYNQKFKDKATLTADKSSS--TAYMQLSLTSEDSAVYICAR---	SYGN---	YPLMDYWGQGTSTVSS	99									
F6B1_P1_E2_H_78_19 (1)	---	QVQLQQSGAELAKPGASVRMSCKASGYTFTSYHHWVKQRPQGLEWIGYINP--STDYTEYNQKFKDKATLTADKSSS--TAYMQLSLTSEDSAVYICVR---	SYGN---	YPLMDYWGQGTSTVSS	100									
F6B_P3_E2_H' (1)	---	QVQLQQSGAELAKPGASVRMSCKASGYTFTSYHHWVKQRPQGLEWIGYINP--STDYTEYNQKFKDKATLTADKSSS--TAYMQLSLTSEDSAVYICVR---	SYGN---	YPLMDYWGQGTSTVSS	101									
F6B_P2_D4_H (1)	---	QVQLQQSGAELAKPGASVRMSCKASGYTFTSYHHWVKQRPQGLEWIGYINP--STDYIEYNQKFKDKATLTADKSSS--TAYMQLSLTSEDSAVYICAR---	SYGN---	YPLMDYWGQGTSTVSS	102									

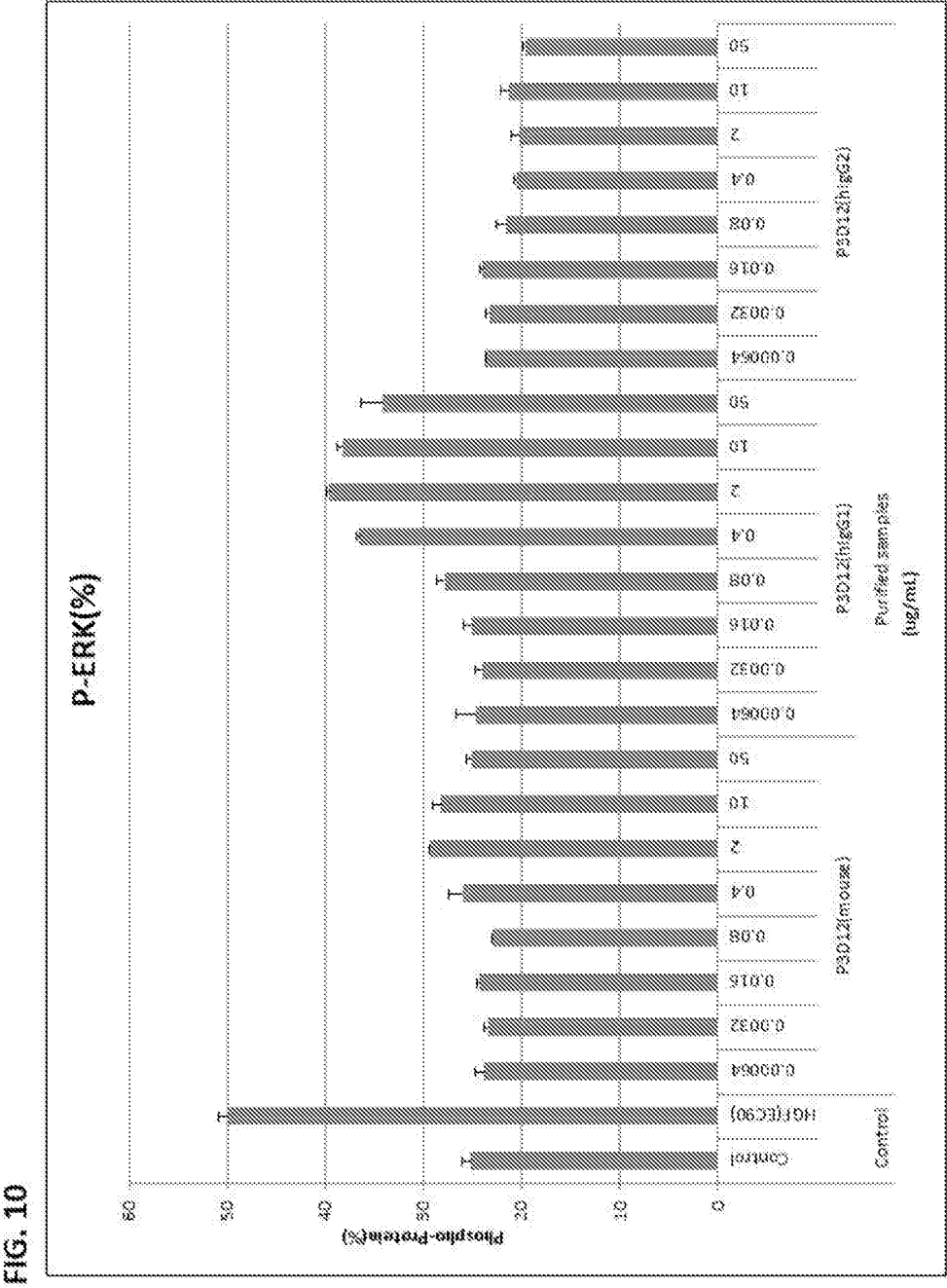
HC F6B1P3D12H7913=F6B1P3E9

HC F6B1P1E2H7819=F6BP3E2

[Fig. 9]

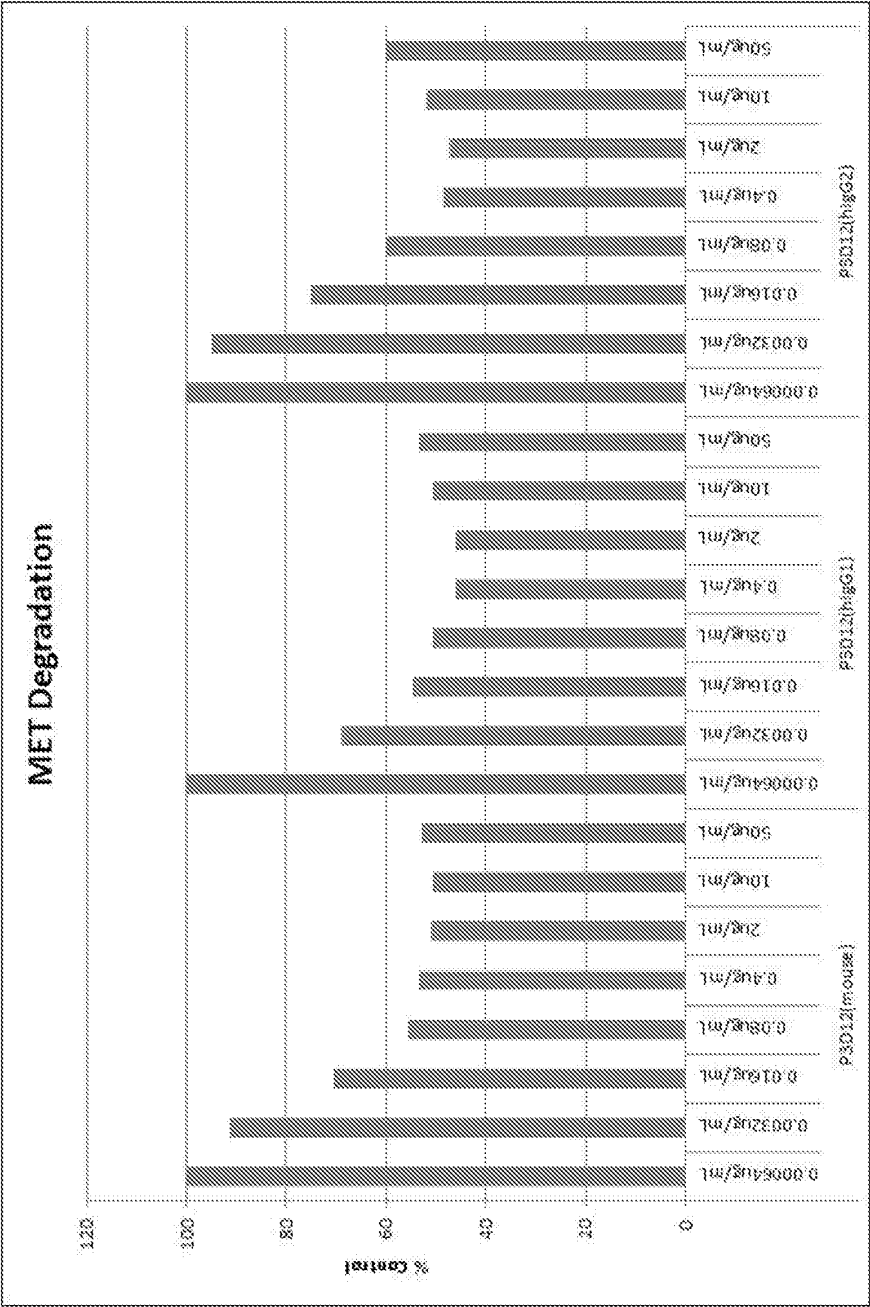


[Fig. 10]



[Fig. 11]

FIG. 11



[Fig. 12]

4

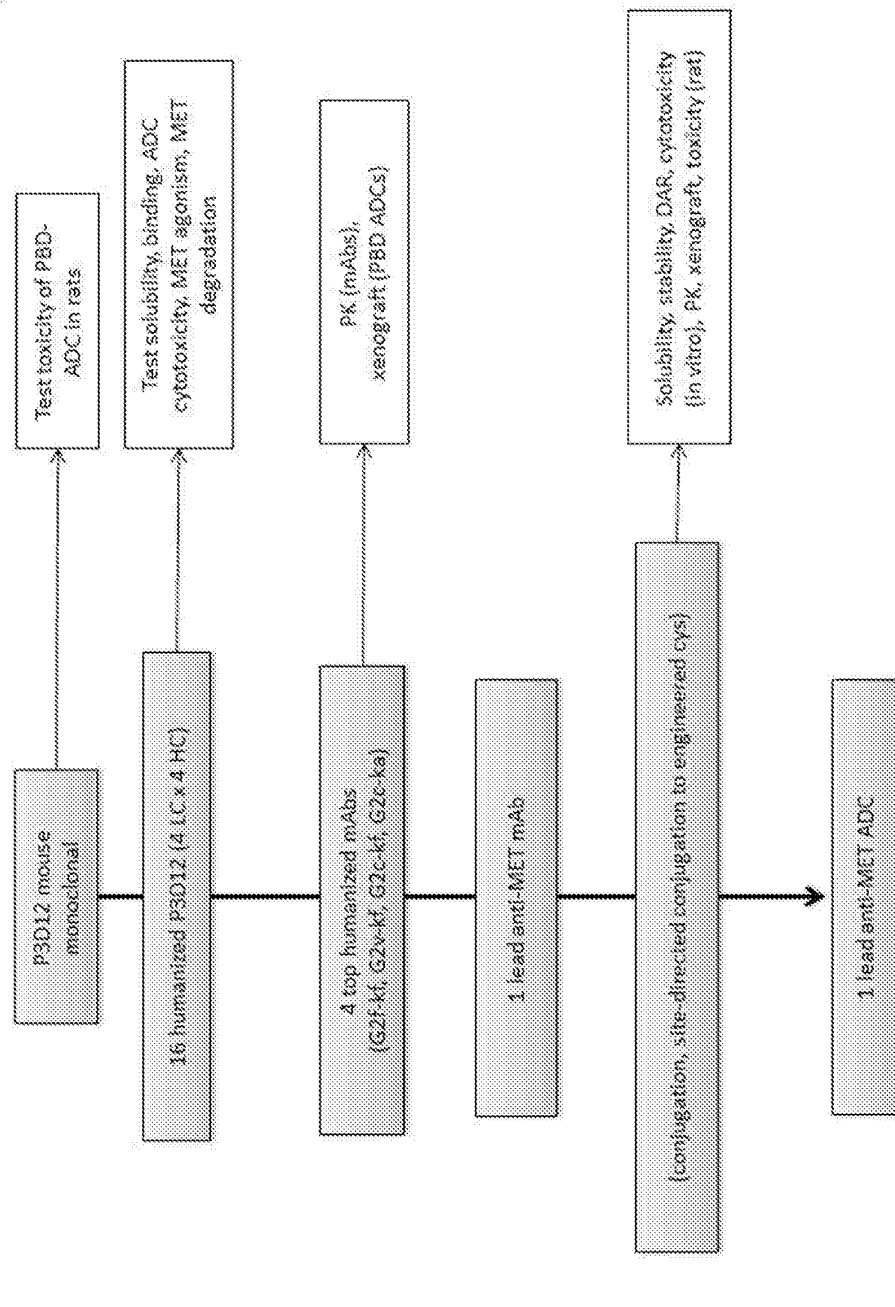


FIG. 12

[Fig. 13]

3

FIG. 13

HUMANIZED LIGHT CHAIN VARIABLE REGIONS

SEQ ID NO.	1	50
P3D12 VL 45 (1)	QIVLTQSPAIMSASPGEKVTITCSASSVTSTNYLYWYQQKPGSSPKMIY	
P3D12 VL-ven 46 (1)	QIVLTQSPATMSAPGERVTITSCSASSVTSTNYLYWYQQKPGSEPRMIY	
P3D12 VL-fra 47 (1)	QIVLTQSPAILSLSPGERATITSCSASSVTSTNYLYWYQQKPGSSPKLIY	
P3D12 VL-abb/sdr 48 (1)	QIVLTQSPATISLSPGERATITSCRASQSVTSTNYLYWYQQKPGSSPRLLIY	
P3D12 VL-cdr 49 (1)	QIVLTQSPATISLSPGERATITSCSASSVTSTNYLYWYQQKPGSSPRLLIY	
	51	100
P3D12 VL (51)	STSNLASGVPARFSGSGCTSYLTISSMFAKDAASYFCHQWSSYPPTFG	
P3D12 VL-ven (51)	STSNLASGVPARFSGSGCTSYTLTISRMEPEDAASYFCHQWSSYPPTFG	
P3D12 VL-fra (51)	STSNLASGVPARFSGSGCTSYTLTISSEAEADAASYFCHQWSSYPPTFG	
P3D12 VL-abb/sdr (51)	STSNLASGVPARFSGSGCTDYTLTISRLEPEDFASYFCHQWSSYPPTFG	
P3D12 VL-cdr (51)	STSNLASGVPARFSGSGCTSYTLTISRLEPEDFASYFCHQWSSYPPTFG	
	101	
P3D12 VL (101)	SGTKLEIKR	
P3D12 VL-ven (101)	SGTKLEIKR	
P3D12 VL-fra (101)	SGTKLEIKR	
P3D12 VL-abb/sdr (101)	SGTKLEIKR	
P3D12 VL-cdr (101)	SGTKLEIKR	

[Fig. 14]

3

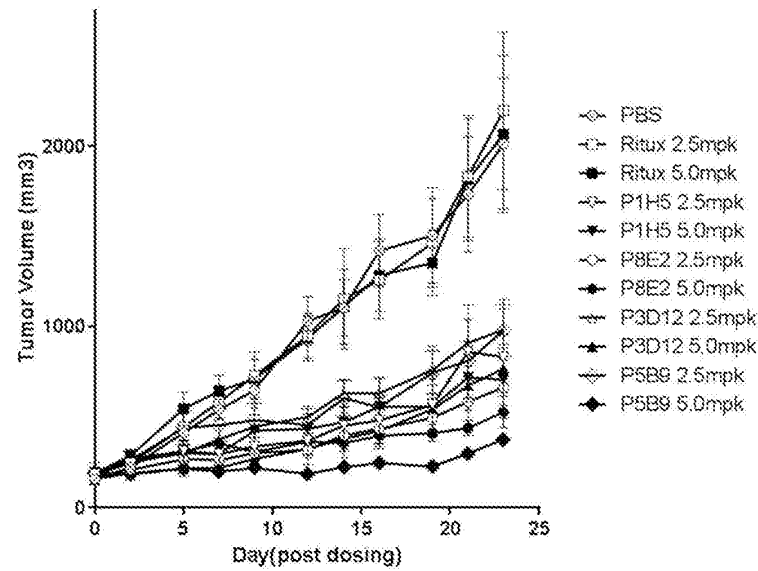
FIG. 14

HUMANIZED HEAVY CHAIN VARIABLE REGIONS

SEQ ID NO.	1	50
P3D12 VH 104 (1)	QVQLQOSGAELAKPGASVMS	CRASGYTFTSYMMHWVKORPCGGLDWIGY
P3D12 VH-fra 105 (1)	QVQLQOSGAELAKPGASVMS	CRASGYTFTSYMMHWVKORPCGGLDWIGY
P3D12 VH-ven 106 (1)	QVQLVQSGAEVAKPGASVMS	CRASGYTFTSYMMHWVKORPCGGLDWIGY
P3D12 VH-abb/sdr 107 (1)	QVQLVQSGAEVAKPGASVMS	CRASGYTFTSYMMHWVKORPCGGLDWIGY
P3D12 VH-cdr 108 (1)	QVQLVQSGAEVAKPGASVMS	CRASGYTFTSYMMHWVKORPCGGLDWIGY
	51	100
P3D12 VH (51)	IKPSTDNTEYNQKFKDKATL	FADKSSSTAYMQLSSLTSEISAVYYCARSY
P3D12 VH-fra (51)	IKPSTDNTEYNQKFKDKATL	FADKSSSTAYMQLSSLTSEISAVYYCARSY
P3D12 VH-ven (51)	IKPSTDNTEYNQKFKDKATL	FADKSSSTAYMQLSSLTSEISAVYYCARSY
P3D12 VH-abb/sdr (51)	IKPSTDNTEYNQKFKDKATL	FADKSSSTAYMQLSSLTSEISAVYYCARSY
P3D12 VH-cdr (51)	IKPSTDNTEYNQKFKDKATL	FADKSSSTAYMQLSSLTSEISAVYYCARSY
	101	119
P3D12 VH (101)	GNYPPLMDYWGQGTSTVTVSS	
P3D12 VH-fra (101)	GNYPPLMDYWGQGTSTVTVSS	
P3D12 VH-ven (101)	GNYPPLMDYWGQGTSTVTVSS	
P3D12 VH-abb/sdr (101)	GNYPPLMDYWGQGTSTVTVSS	
P3D12 VH-cdr (101)	GNYPPLMDYWGQGTSTVTVSS	

[Fig. 15]

FIG. 15 MKN-45 xenograft



[Fig. 16]

FIG. 16

