

Table S1. Amino acid sequences of 10E8 variants.

Heavy chain

	5	15	25	35	45	52C	62	72	82	84
10E8	EVQLVESGGG	LVKPGGSLRL	SCSASGDFD	NAWMTWVRQP	PGKGLEWVGR	ITGPGEGWSV	DYAAPVEGRF	TISRLNSINF	LYLEMNLRM	
HC6	.R.A.
HC6-S74W	.R.A.W.	.	.
HC6-S74Y	.R.A.Y.	.	.
HC6-S74Y-DKTT	.R.A.D.YK.T	.	.T
H6	.R.	.	.N.	D.	.	.S.	.	.ES.K.	.	.V.T
H6-DTKT	.R.	.	.N.	D.	.	.S.	.	.ES.K.	.D.TK.T	.V.T
H6-DTKT-DNTY	.R.ES.K.	.D.TK.T	.V.T
H6-DTKT-DNTY-LS	.R.ES.K.	.D.TK.T	.V.T
H6-DTKT-DNTY-Y100e	.R.ES.K.	.D.TK.T	.V.T
H8	.R.	.R.	.N.S.K.	.M.	F.
H8-DYKT	.R.	.R.	.N.S.K.	.D.YK.T	F.
	89	99	101I	107	117	127	137	147	157	
10E8	EDSGLYFCAR	TGKYDFWSG	YPPGEEYQD	WGRGTLVTS	SASTKGPSVF	PLAPSSKSTS	GGTAALGCLV	KDYFPEPVTV	SWNSGALTS	
HC6I.
HC6-S74WI.
HC6-S74YI.
HC6-S74Y-DKTTI.
H6	.T.Y.	.H.	.	.Q.	.I.
H6-DTKT	.T.Y.	.H.	.	.Q.	.I.
H6-DTKT-DNTY	.T.Y.	.	.	.Q.	.I.
H6-DTKT-DNTY-LS	.T.Y.	.	.	.Q.	.I.
H6-DTKT-DNTY-Y100e	.T.Y.	.	C.	.Q.	.I.
H8	.T.	.H.A.	.G.	.LE.	.Q.	.I.
H8-DYKT	.T.	.H.A.	.G.	.LE.	.Q.	.I.
	167	177	187	197	207	217	227	237	247	
10E8	VHTFFAVLQS	SGLYSLSSVV	TVPSSSLGTQ	TYICNVNHPK	SNTKVDKQVE	PKSCDKTHTC	PPCPAPELLG	GPSVFLFPPK	PKDTLMISRT	
HC6
HC6-S74W
HC6-S74Y
HC6-S74Y-DKTT
H6
H6-DTKT
H6-DTKT-DNTY
H6-DTKT-DNTY-LS
H6-DTKT-DNTY-Y100e
H8
H8-DYKT
	257	267	277	287	297	307	317	327	337	
10E8	PEVTCVVVDV	SHEDPEVKFN	WYVDGVEVHN	AKTKPREEQY	NSTYRVVSVL	TVLHQDWLNG	KEYKCKVSNK	ALPAPIEKTI	SKAKGQPREP	
HC6
HC6-S74W
HC6-S74Y
HC6-S74Y-DKTT
H6
H6-DTKT
H6-DTKT-DNTY
H6-DTKT-DNTY-LS
H6-DTKT-DNTY-Y100e
H8
H8-DYKT
	347	357	367	377	387	397	407	417	427	
10E8	QVYTLPPSRD	ELTKNQVSLT	CLVKGFPYPSD	IAVEWESNGQ	PENNYKTPPP	VLDSDGSFFL	YSKLTVDKSR	WQQGNVFCSS	VMHEALHNHY	
HC6
HC6-S74W
HC6-S74Y
HC6-S74Y-DKTT
H6
H6-DTKT
H6-DTKT-DNTY
H6-DTKT-DNTY-LSL.	.S.
H6-DTKT-DNTY-Y100e
H8
H8-DYKT
	437									
10E8	TQKSLSLSPG	K								
HC6
HC6-S74W
HC6-S74Y
HC6-S74Y-DKTT
H6
H6-DTKT
H6-DTKT-DNTY
H6-DTKT-DNTY-LS
H6-DTKT-DNTY-Y100e
H8
H8-DYKT

Continues...

Table S1 (continued). Amino acid sequences of 10E8 variants*.

Light chain

	5	16	26	36	46	56	66	76	86
10E8	SYELTQETGV	SVALGRTVII	TCRGDSLRS	YASWYQKKPG	QAPILLFYGK	NNRPSGVPDR	FSGSASGNRA	SLTISGAQAE	DDAEYCSSR
L3D...V.....I...T.....	..E.D.....
L3-ASPAKQ	AS...DPA.	...KQ...V.....I...T.....	..E.D.....
L3-ASPAKQ-S30C	AS...DPA.	...KQ...C.V.....I...T.....	..E.D.....
L10	AS...DPA.	...KQ...V.....	..V.V.....I...S..T.A.....D.....

	95A	103	113	123	133	143	153	163	173
10E8	DKSGSRLSVF	GGGTKLTVLS	QPKAAPSVTL	FPPSSEELQA	NKATLVCLIS	DFYPGAVTVA	WKADSSPVKA	GVETTTPSKQ	SNNKYAASSY
L3
L3-ASPAKQ
L3-ASPAKQ-S30C
L10
					
	183	193	203	213					
10E8	LSLTPEQWKS	HRSYSCQVTH	EGSTVEKTVA	PTECS					
L3					
L3-ASPAKQ					
L3-ASPAKQ-S30C					
L10					

* 10E8 variants comprised the following heavy and light chain combinations:
 10E8v1: HC6-S74Y-DKTT/L3
 10E8v4: H6-DTKT-DNTY/L3-ASPAKQ
 10E8v4-LS: H6-DTKT-DNTY-LS/L3-ASPAKQ
 10E8v4-DS: H6-DTKT-DNTY-Y100eC/L3-ASPAKQ-S30C
 10E8v5: HC6-S74Y-DKTT/L3-ASPAKQ