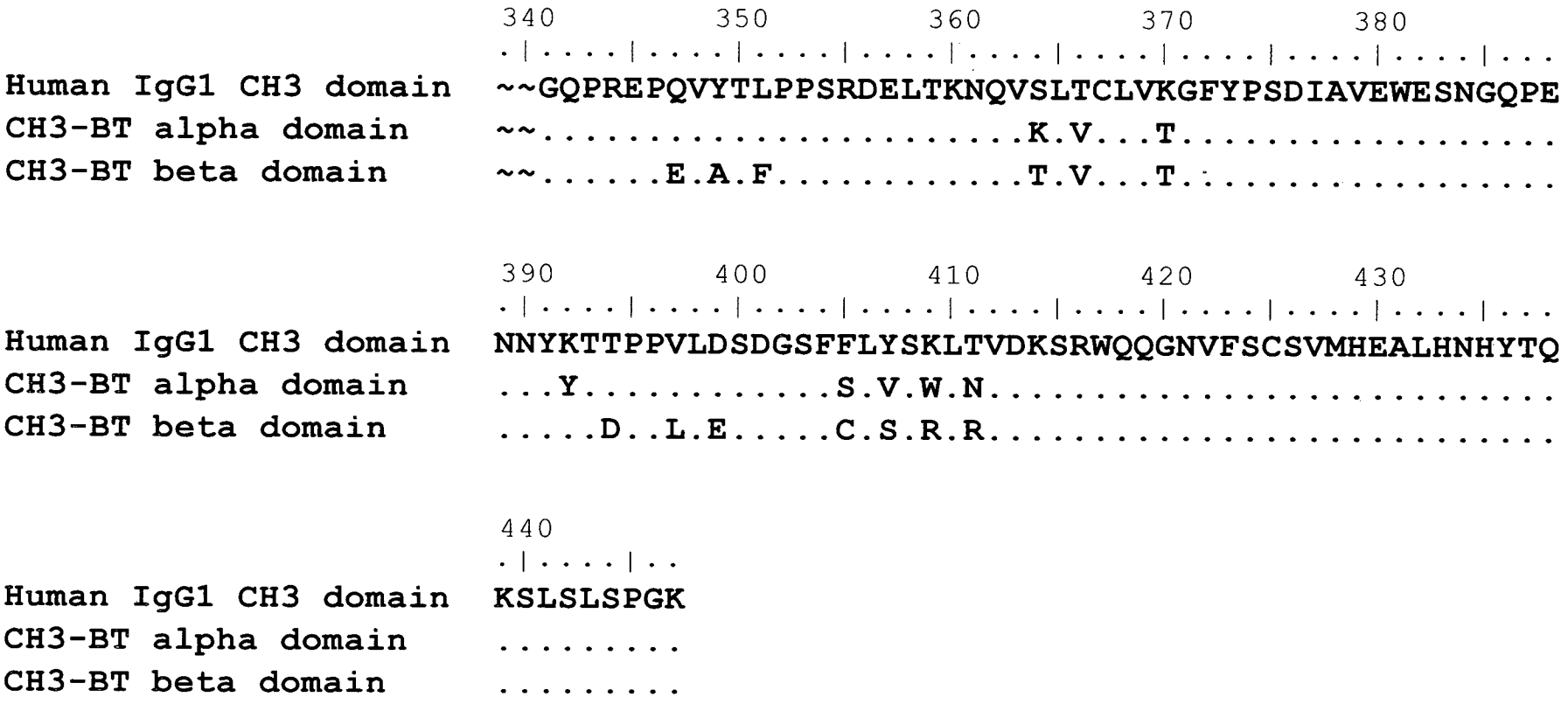
FIG. 1



SUBSTITUTE SHEET (RULE 26)

FIG. 2

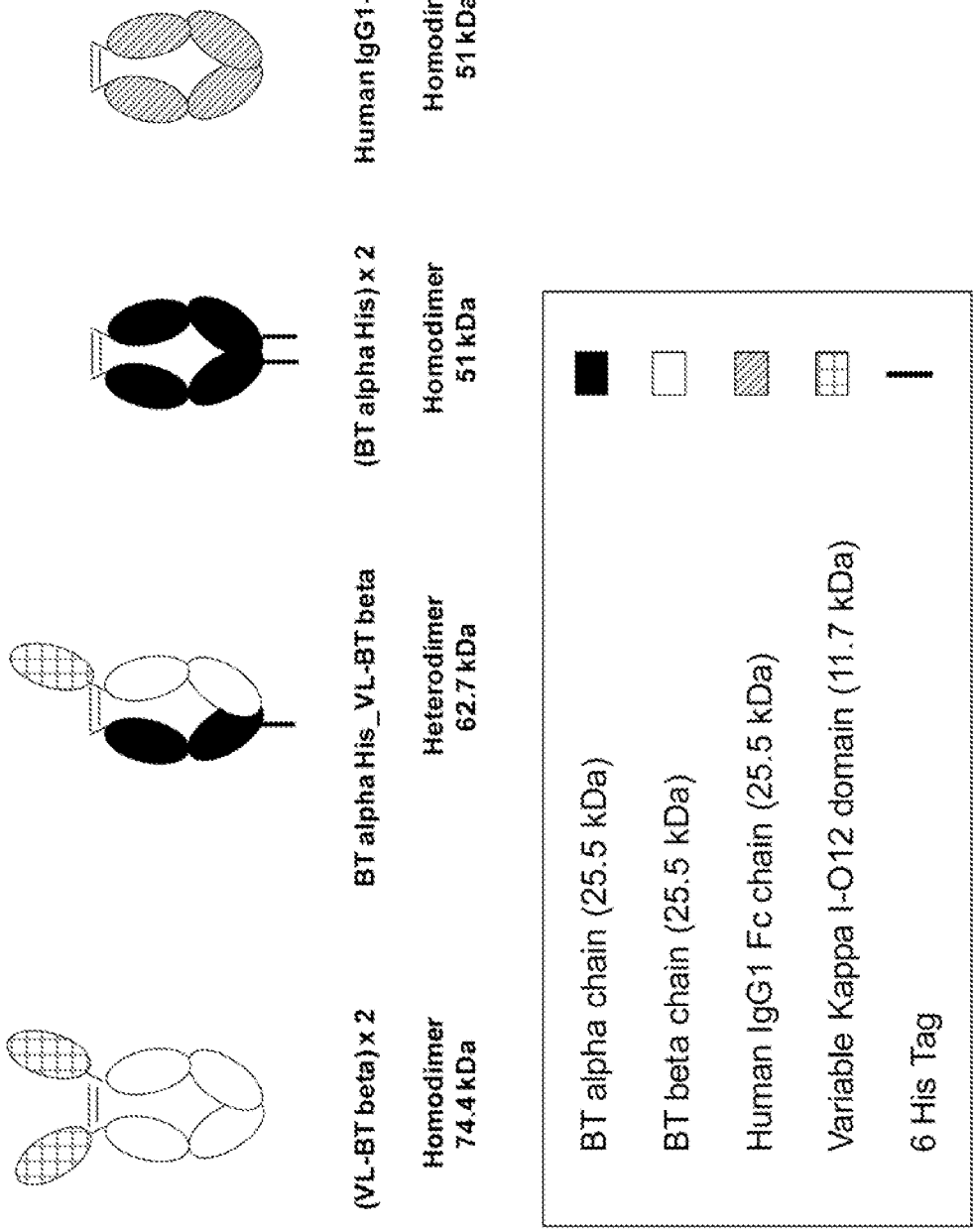

(VL-BT beta) x 2
Homodimer
74.4 kDa
BT alpha His_VL-BT beta
Heterodimer
62.7 kDa
(BT alpha His) x 2
Homodimer
51 kDa
Human IgG1-Fc
Homodimer
51 kDa
BT alpha chain (25.5 kDa)
BT beta chain (25.5 kDa)
Human IgG1 Fc chain (25.5 kDa)
Variable Kappa I-O12 domain (11.7 kDa)
6 His Tag

FIG. 3

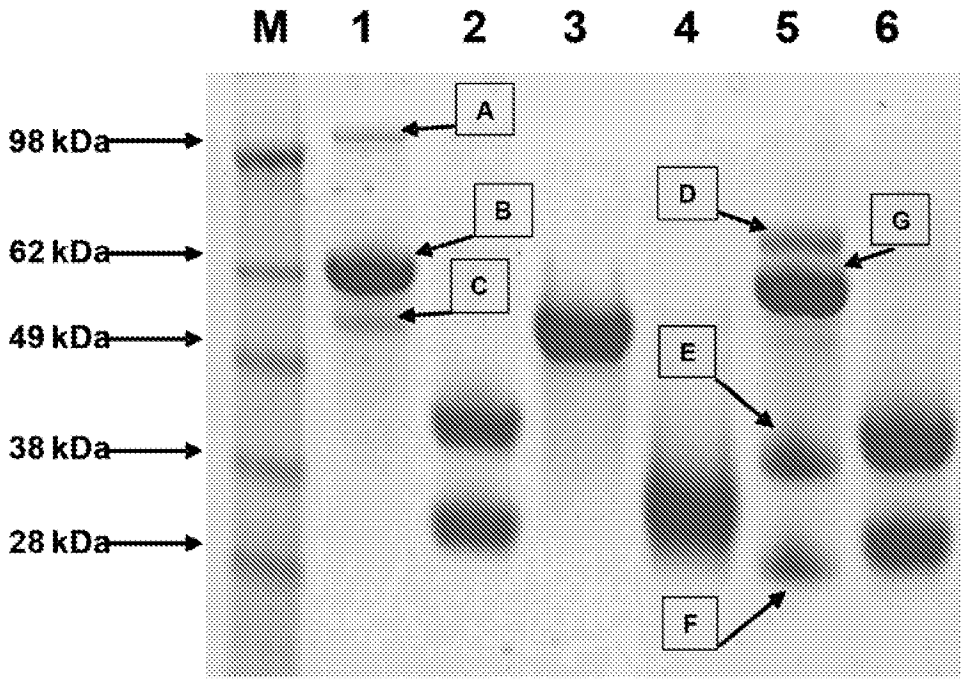
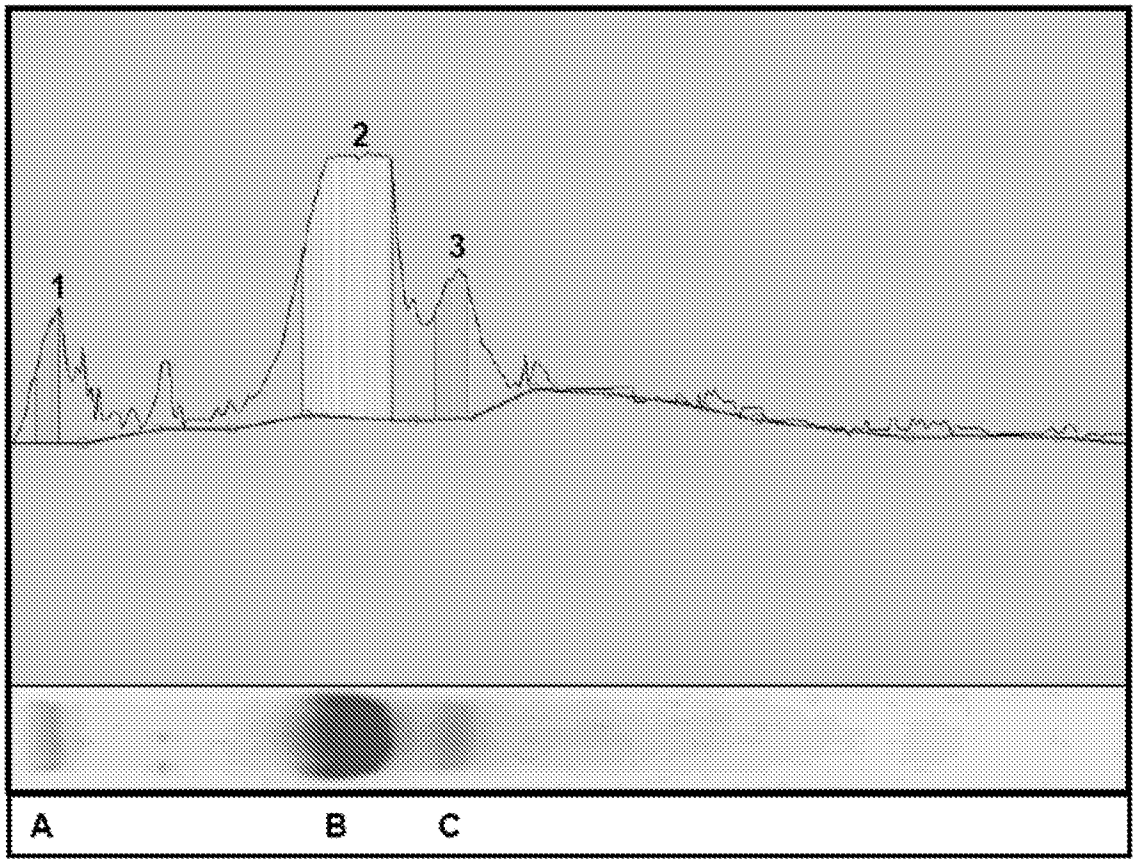
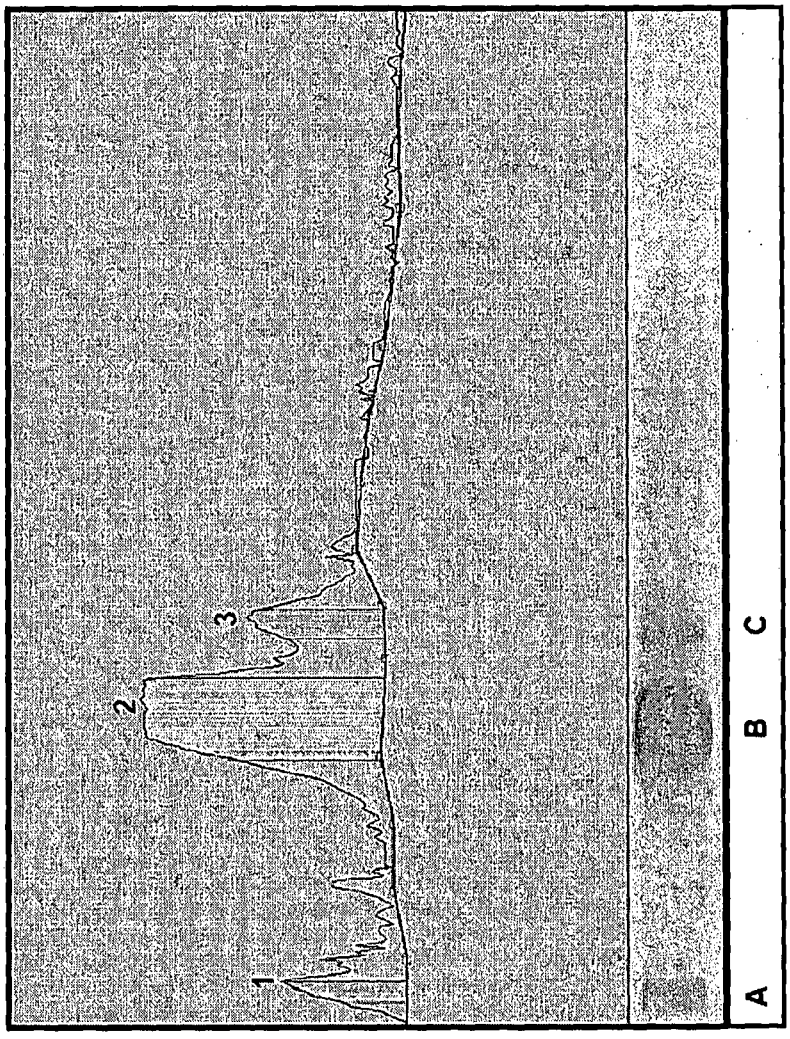


FIG. 4A



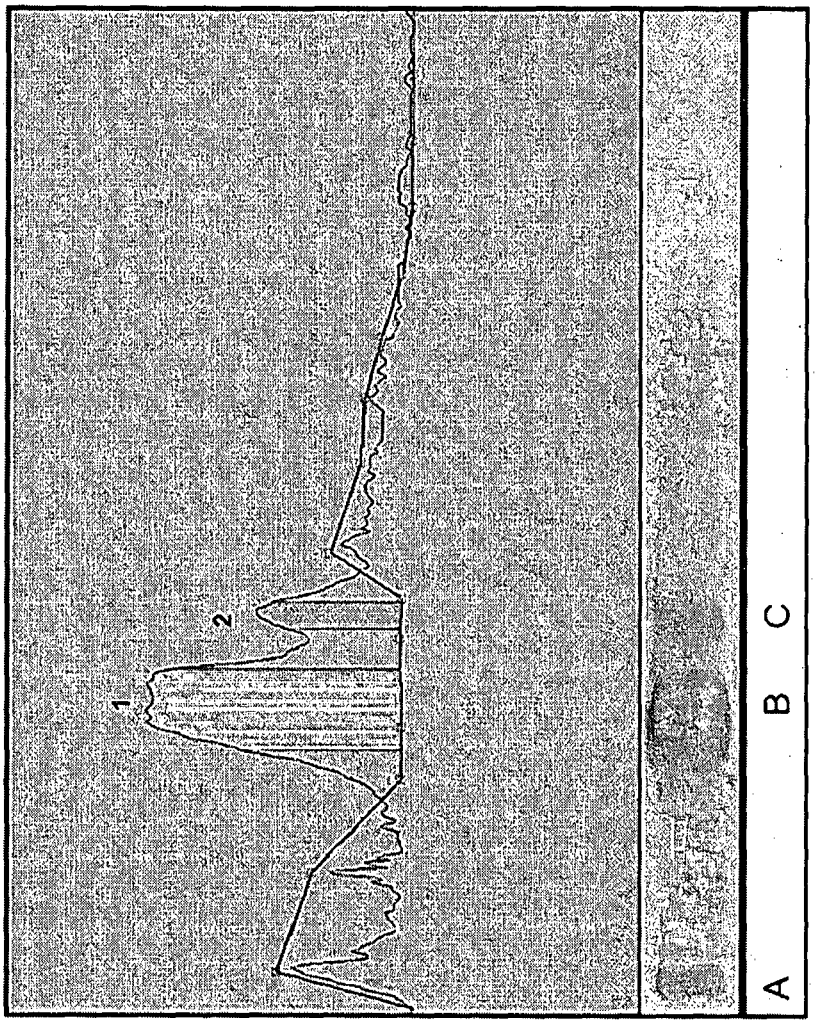
LANE	PEAK	DIST	WIDTH	HEIGHT	AREA	%
1	1	12	8	40	479	8.5
1	2	130	34	130	4286	76.1
1	3	169	12	74	869	15.4

FIG. 4A



LANE	PEAK	DIST	WIDTH	HEIGHT	AREA	%
1	1	12	8	40	479	8.5
1	2	130	34	130	4286	76.1
1	3	169	12	74	869	15.4

FIG. 4B



LANE	PEAK	CST	WIDTH	HEIGHT	AREA	%
1	1	122	34	136	4404	83.8
1	2	170	11	76	852	16.2

FIG. 5

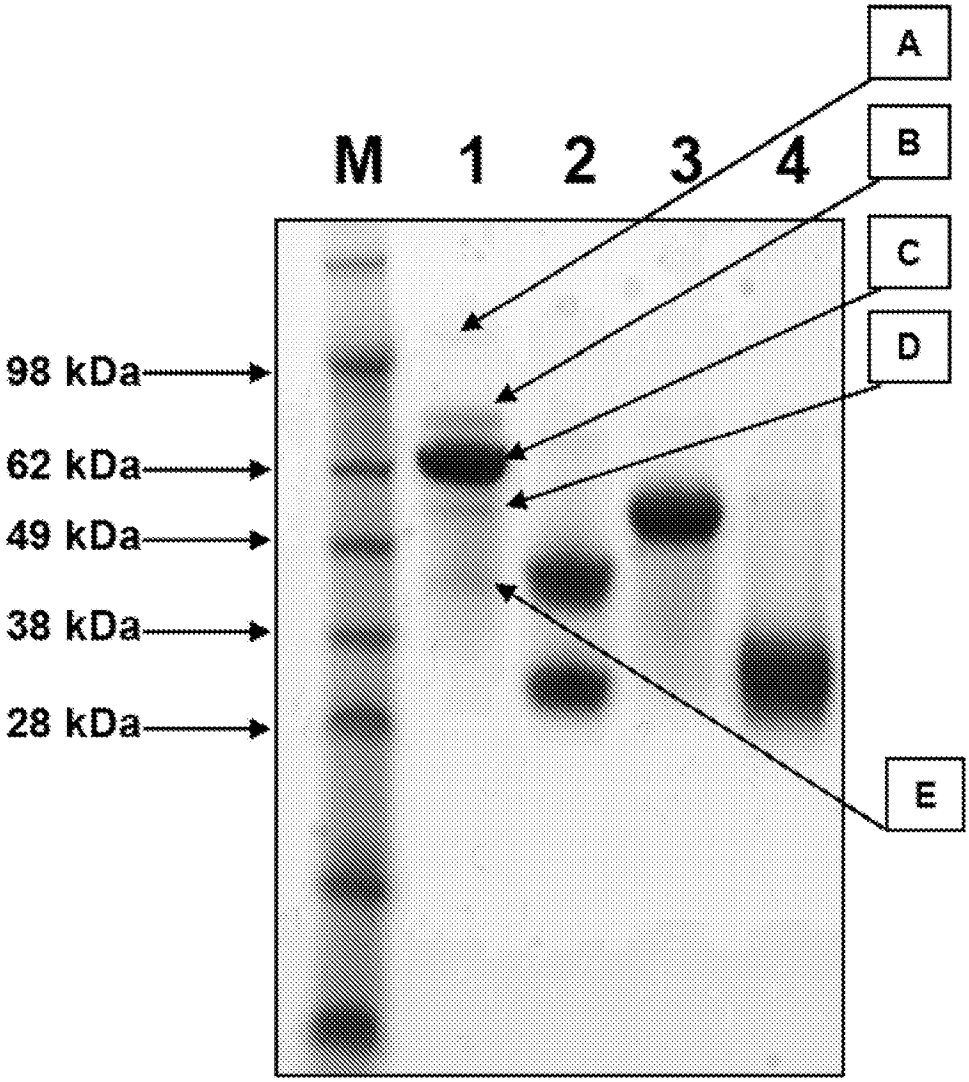


FIG. 6

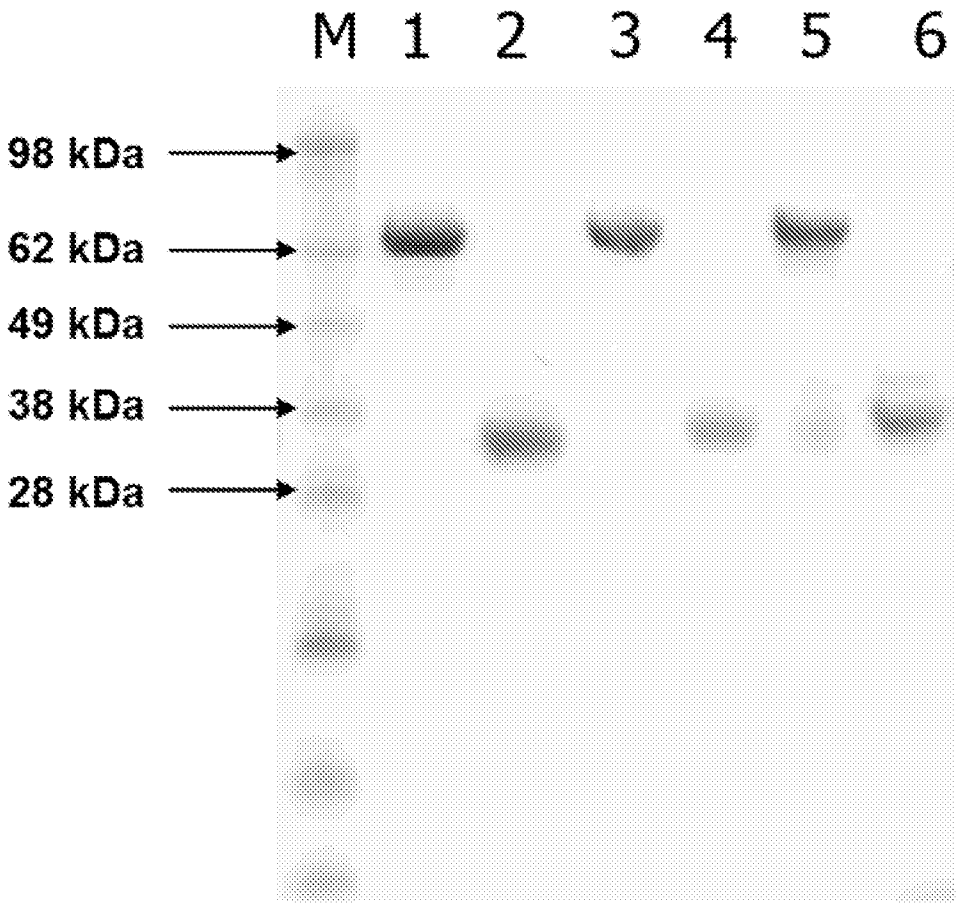
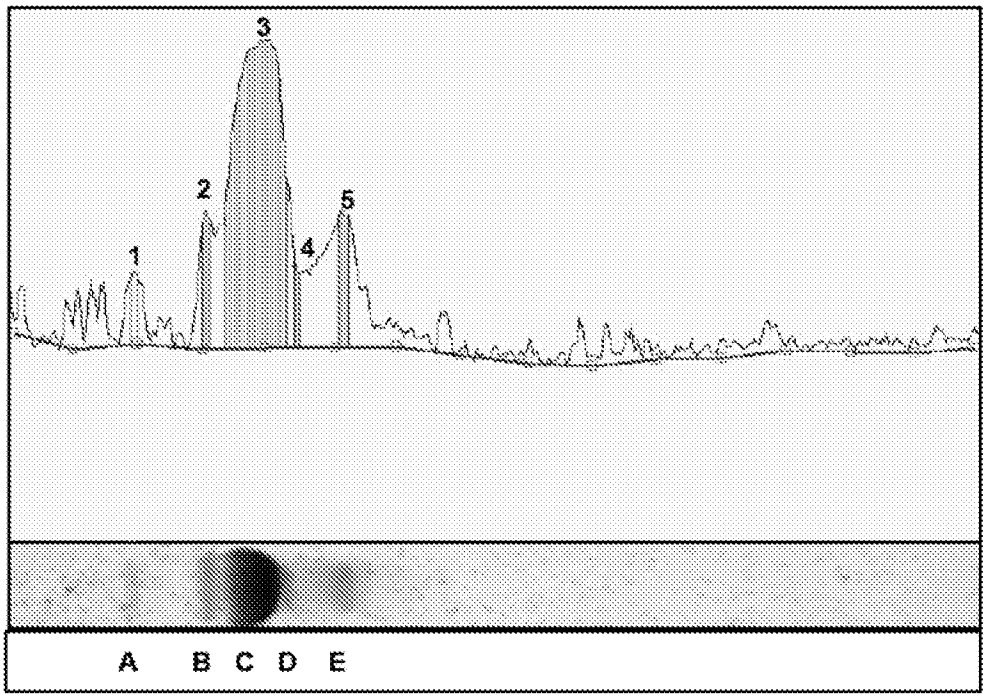
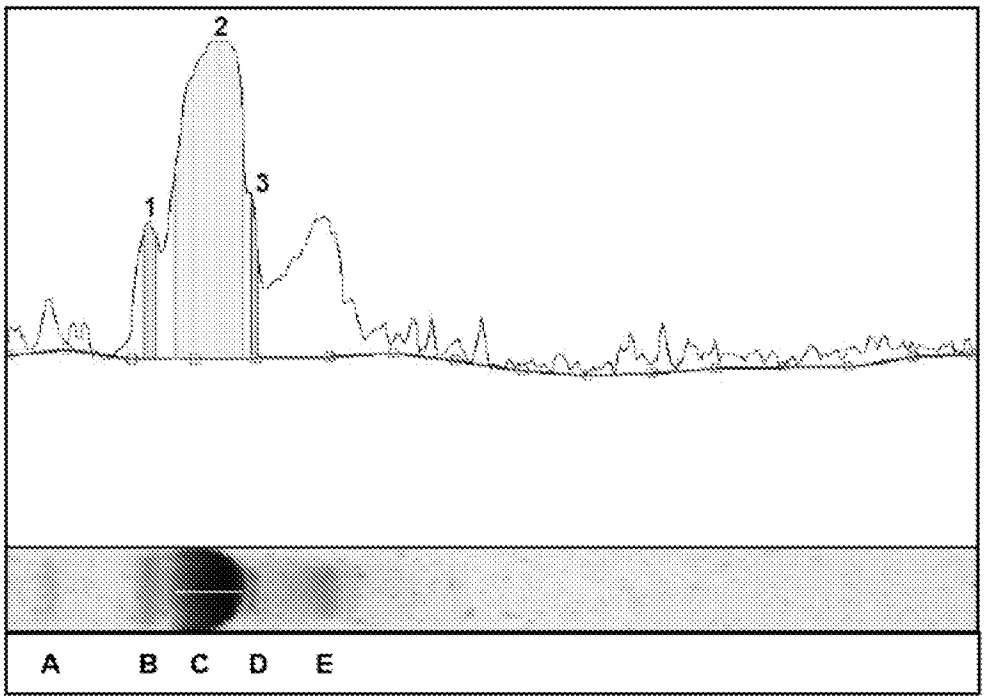


FIG. 7A



LANE	PEAK	DIST	WIDTH	HEIGHT	AREA	%
1	1	70	4	11859	56192	3.0
1	2	110	5	22496	128398	6.8
1	3	146	35	49916	1534196	80.7
1	4	162	3	11971	51389	2.7
1	5	187	5	22496	129985	6.8

FIG. 7B



LANE	PEAK	DIST	WIDTH	HEIGHT	AREA	%
1	1	64	5	22053	126843	7.8
1	2	99	29	51049	1404252	86.9
1	3	109	3	23782	85575	5.3

FIG. 7C

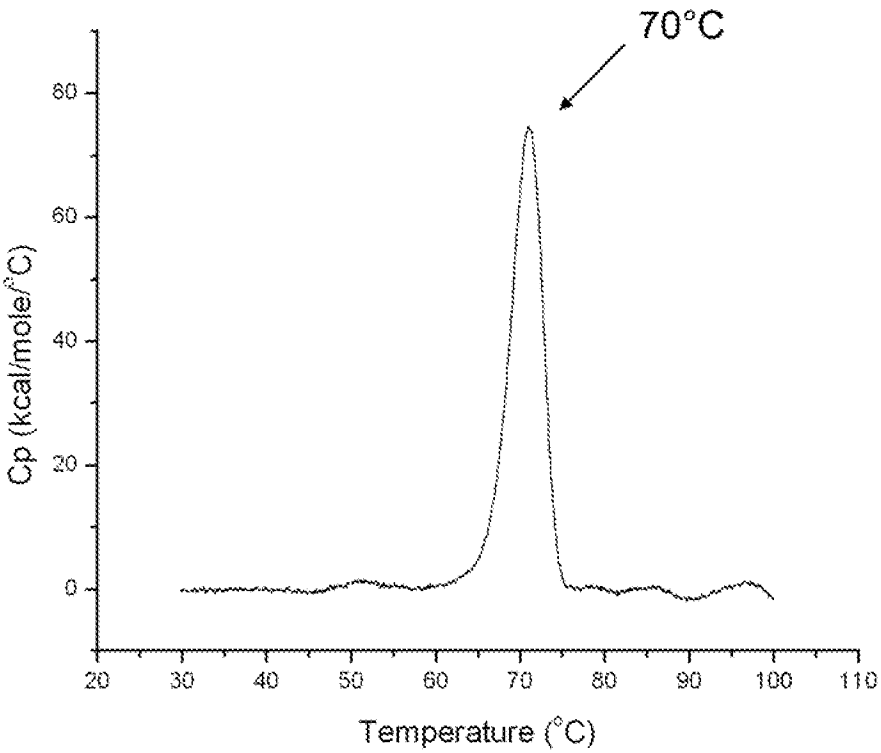

70°C
Cp (kcal/mole/°C)
0
20
40
60
60
20
30
40
50
60
70
80
90
100
110
Temperature (°C)

FIG. 8

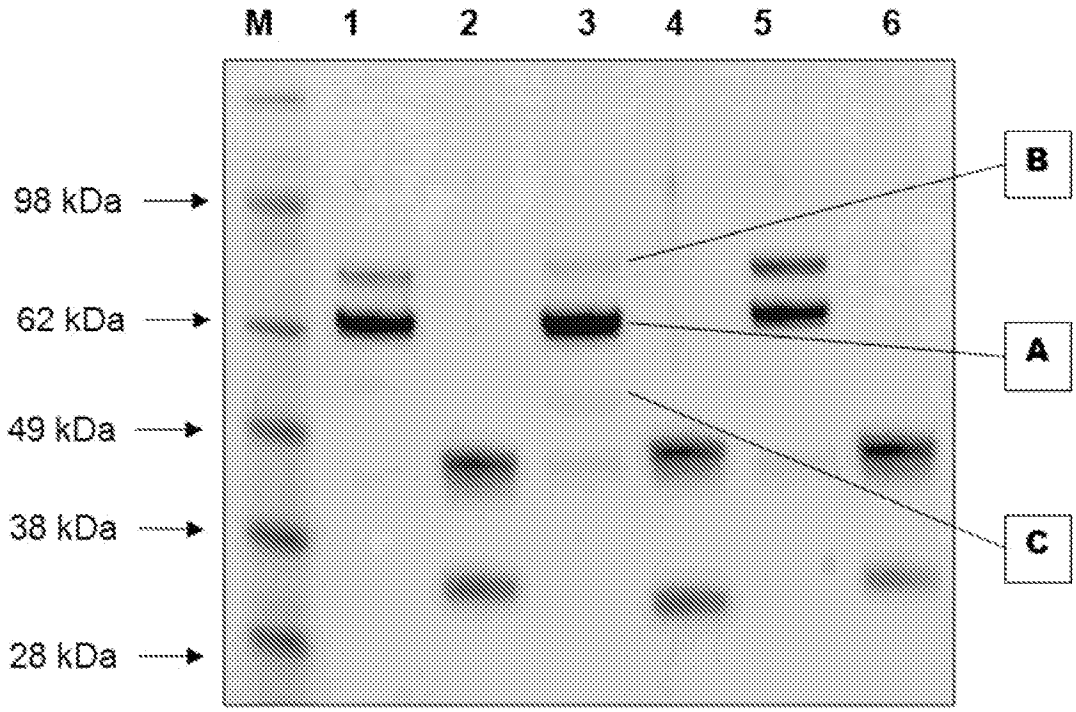


FIG. 9

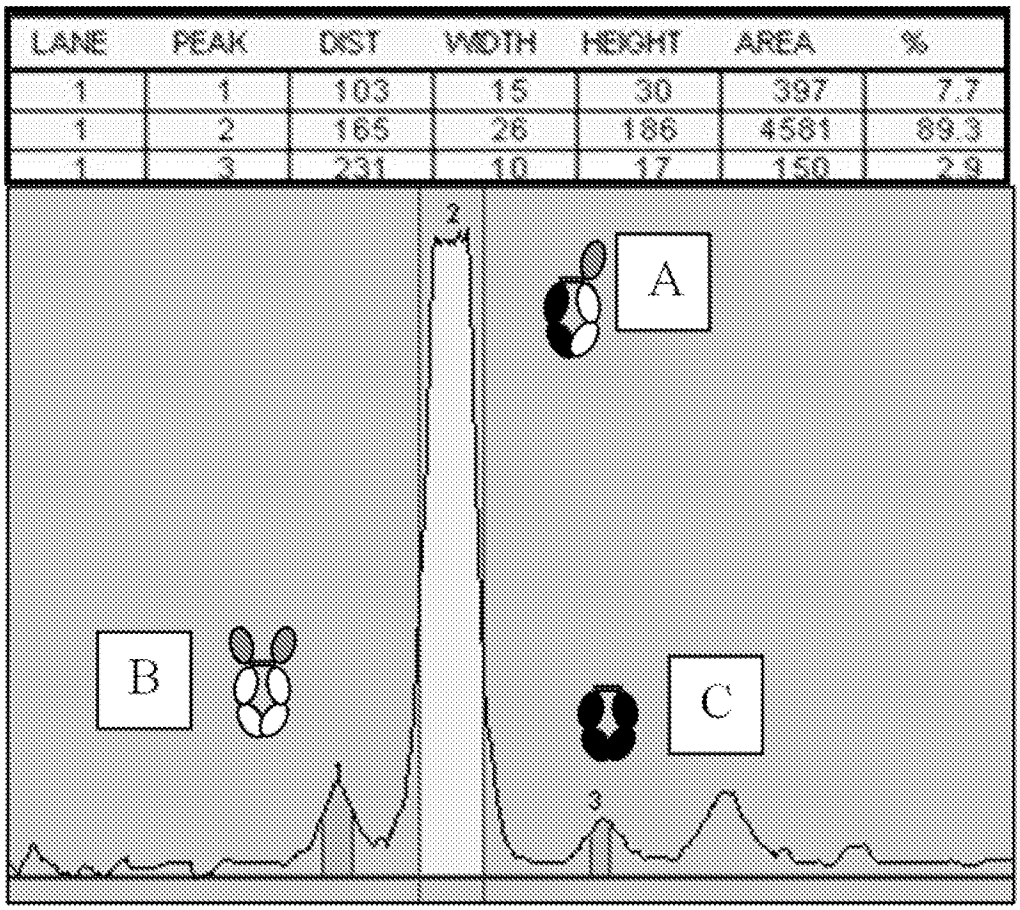

LANE PEAK DST WIDTH HEIGHT AREA %
1 1 103 15 30 397 7.7
1 2 165 26 186 4581 89.3
1 3 231 10 17 150 2.9
2
A
B
C
3

FIG. 10

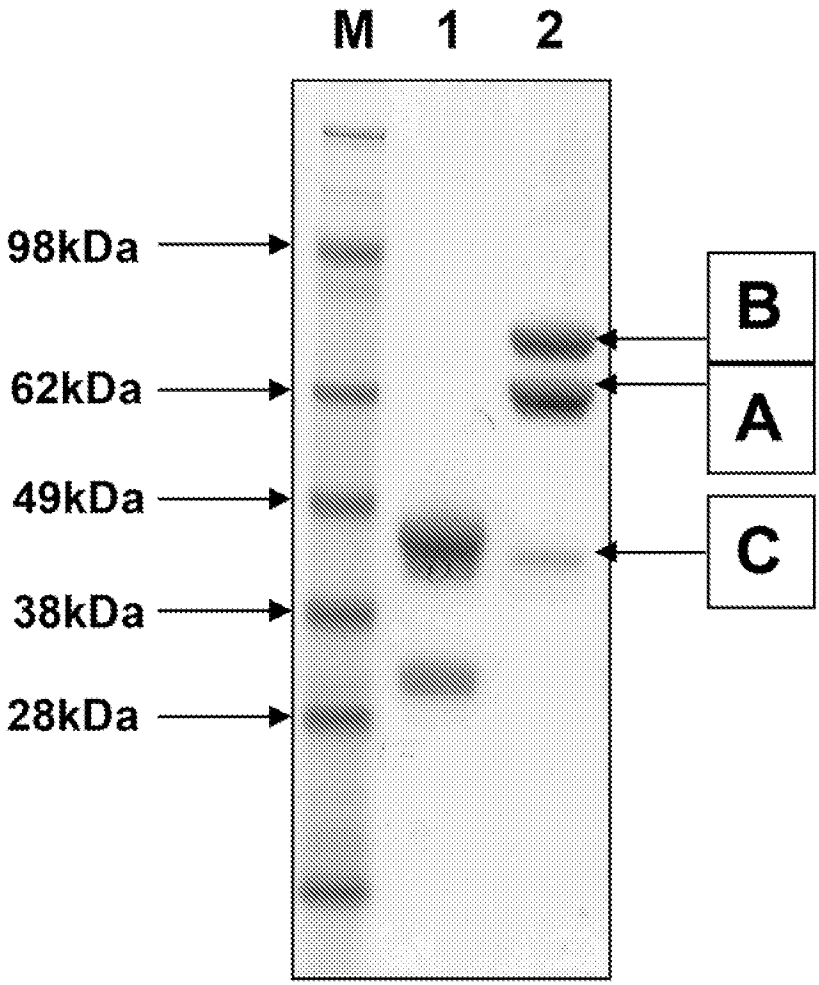


FIG. 11A

PEAK	DIST	WIDTH	HEIGHT	AREA	%
1	355	61	163	7459	38.5
2	450	77	186	10092	52.1
3	731	34	61	1834	9.5

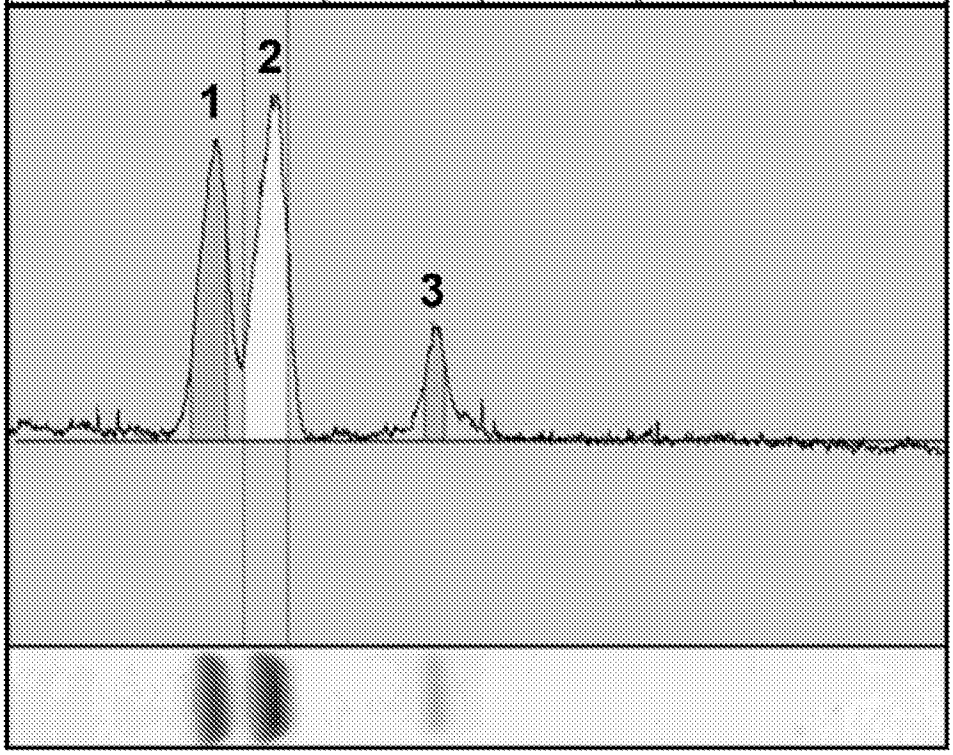


FIG. 11B

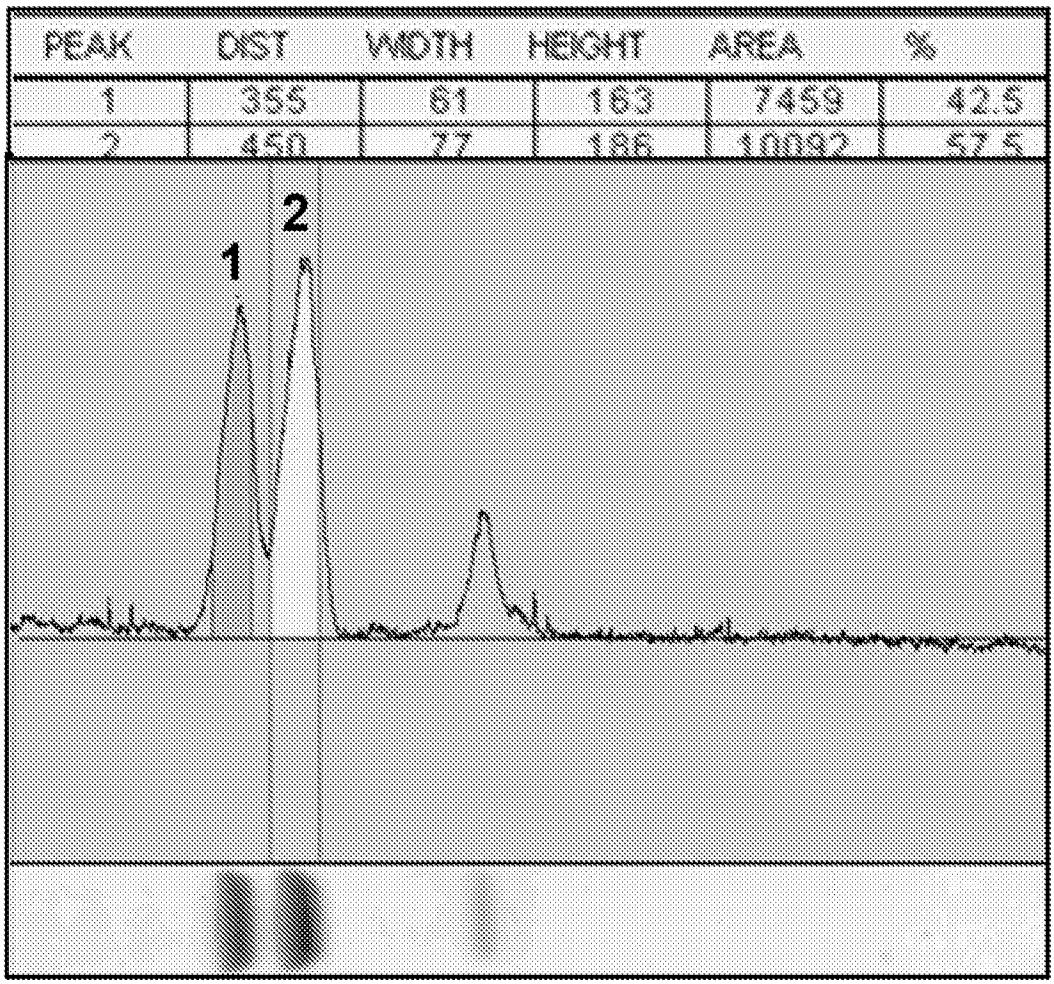

PEAK DIST WIDTH HEIGHT AREA %
1 355 61 163 7459 42.5
2 450 77 186 10092 57.5
1
2

FIG. 12

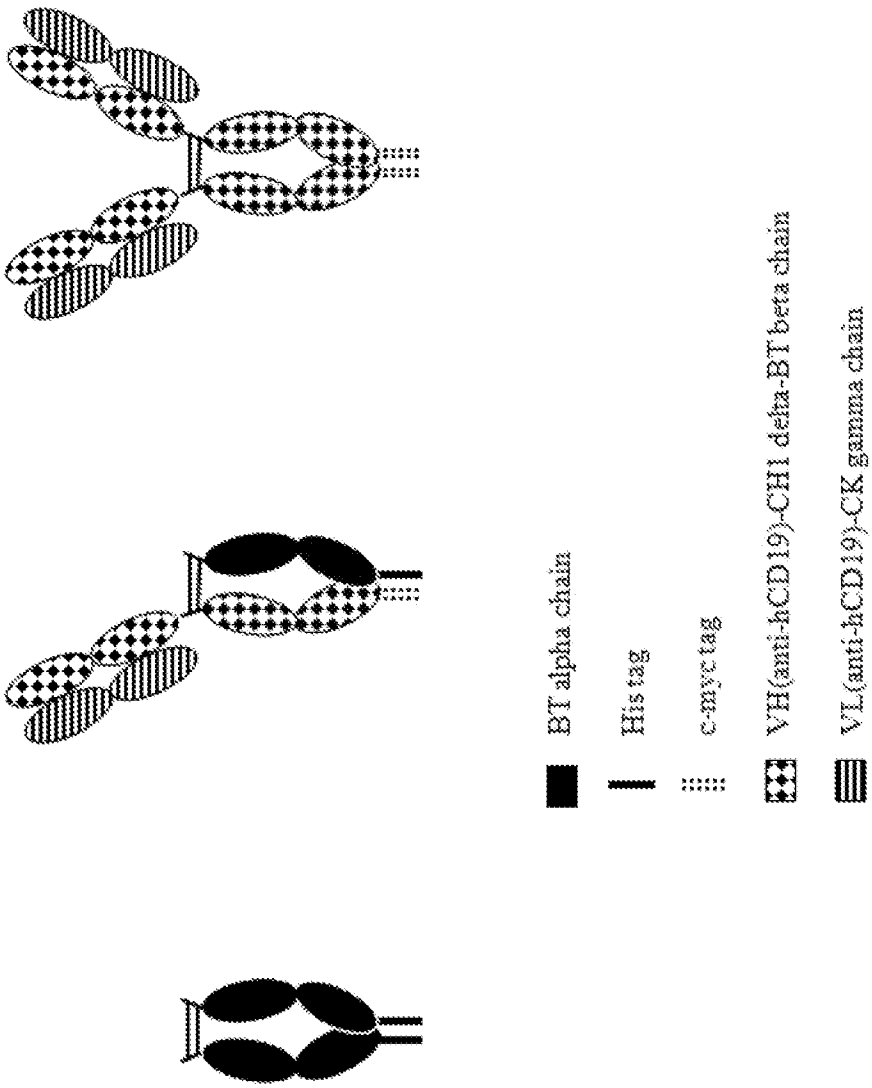

BT alpha chain
His tag
c-myc tag
VH(anti-hCD19)-CH1 delta-BT beta chain
VL(anti-hCD19)-CK gamma chain

FIG. 13

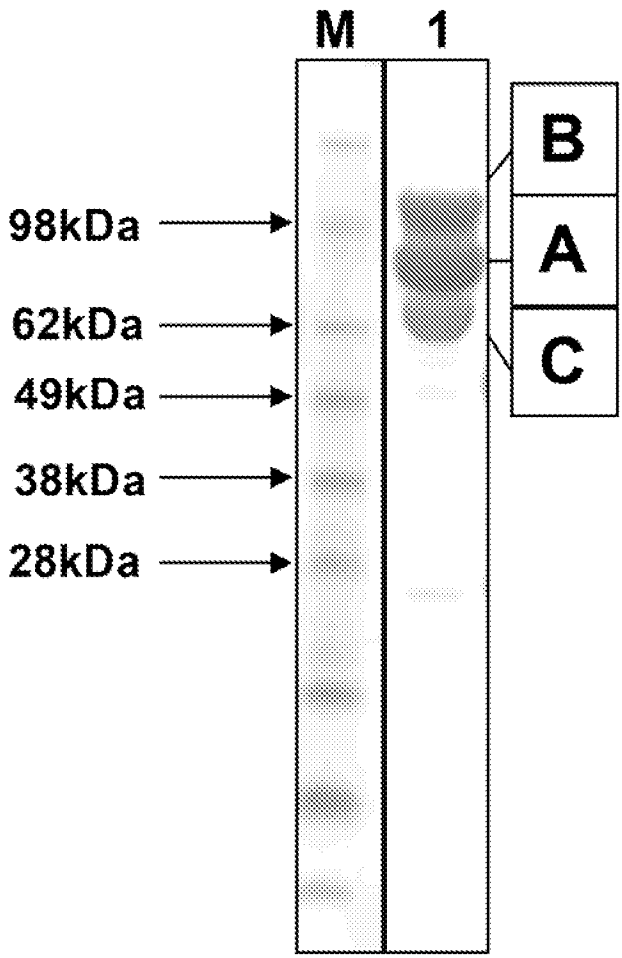


FIG. 14

PEAK	DIST	WIDTH	HEIGHT	AREA	%
1	20	16	118	1920	26.2
2	64	30	129	3684	50.2
3	90	16	107	1734	23.6

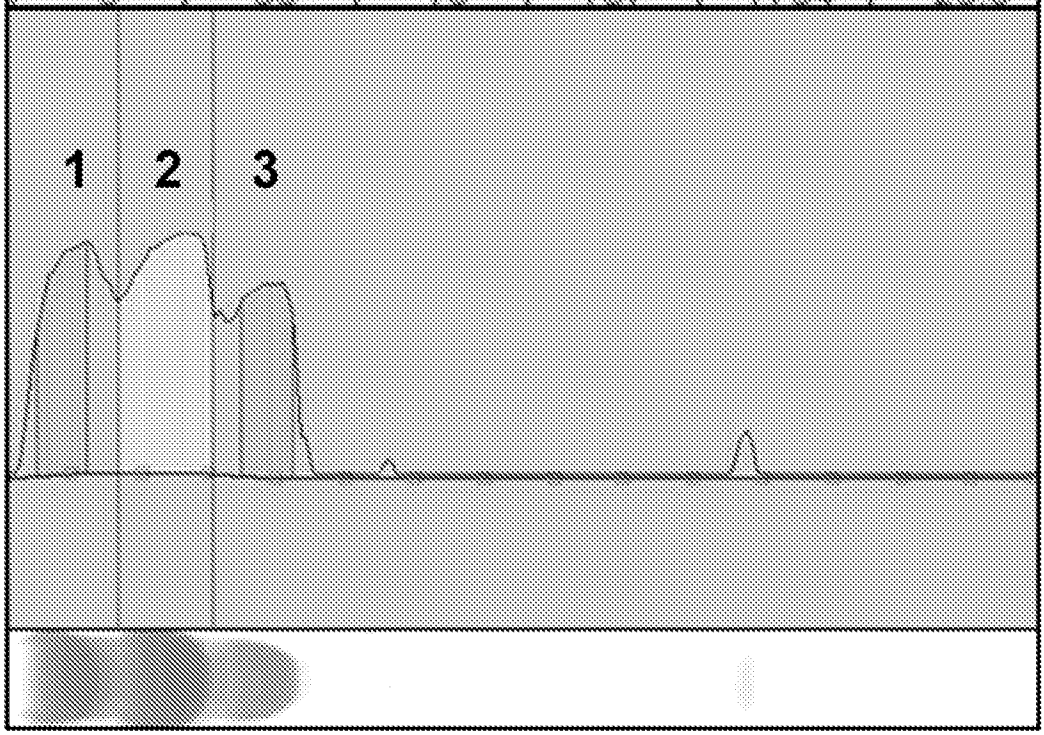


FIG. 15

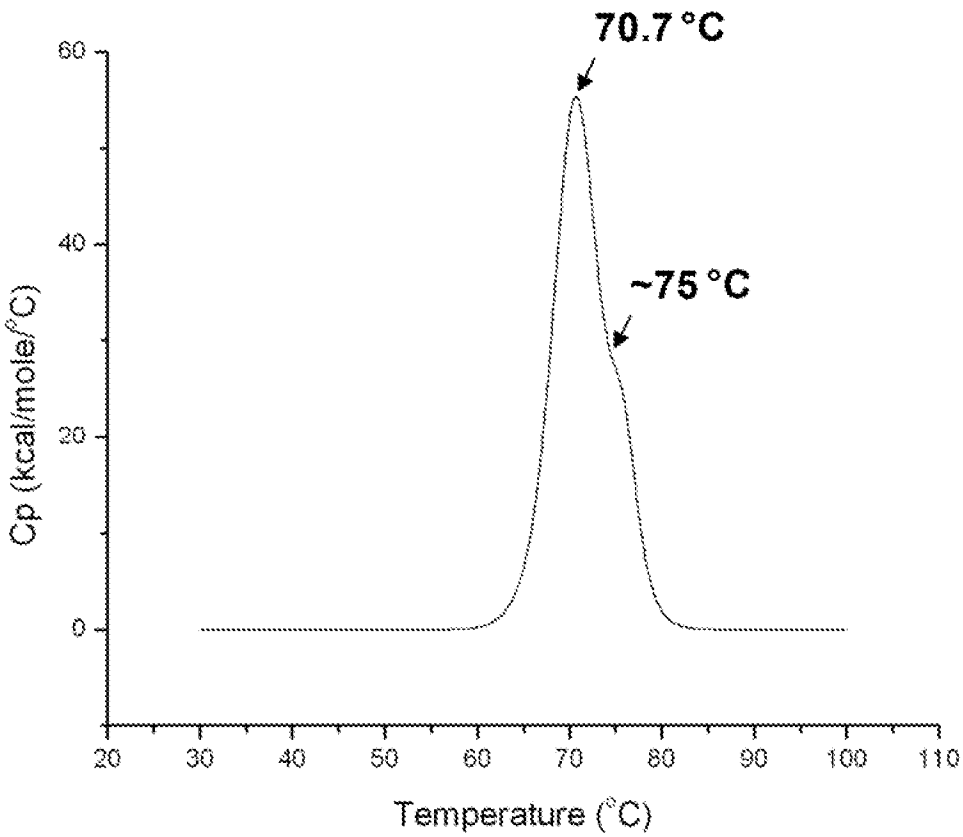

70.7 °C
~75 °C
Cp (kcal/mole/°C)
60
40
20
0
20
30
40
50
60
70
80
90
100
110
Temperature (°C)

FIG. 16

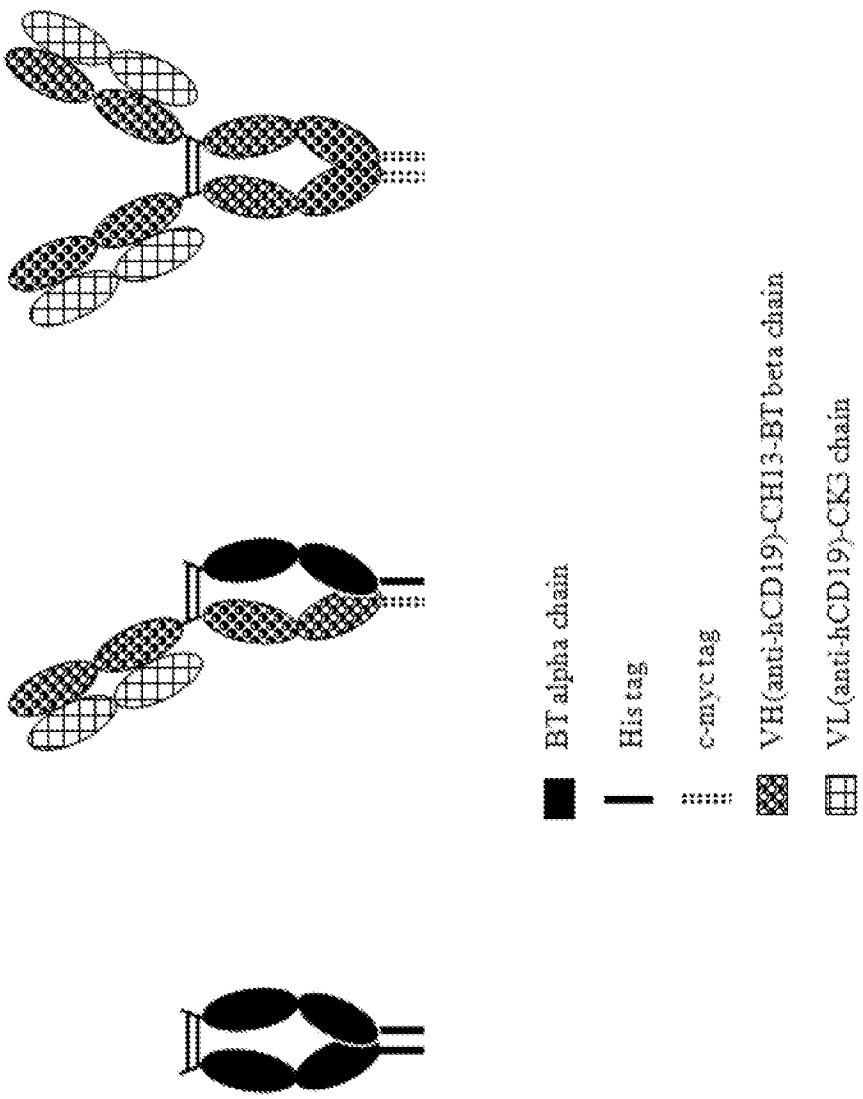

BT alpha chain
His tag
c-myc tag
VH(anti-hCD19)-CH13-BT beta chain
VL(anti-hCD19)-CK3 chain

FIG. 17

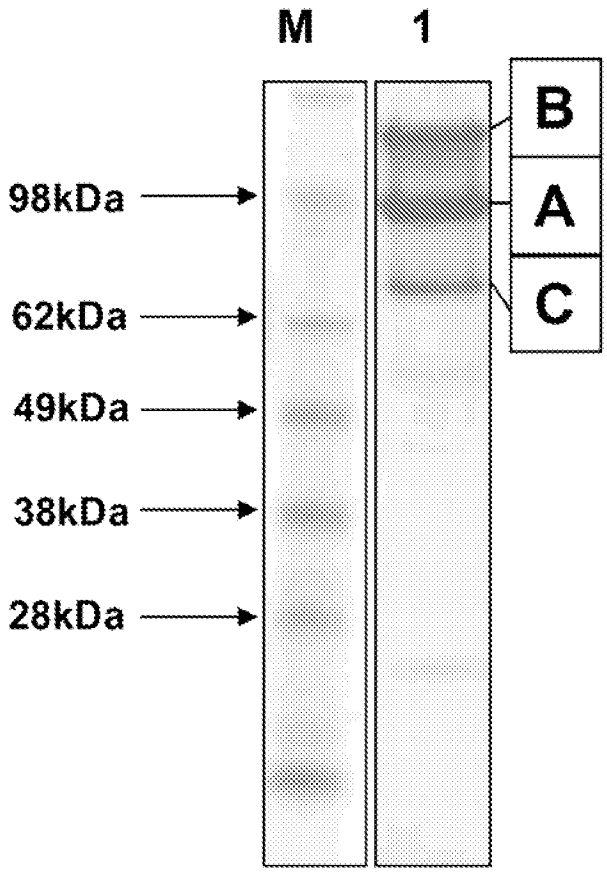


FIG. 18

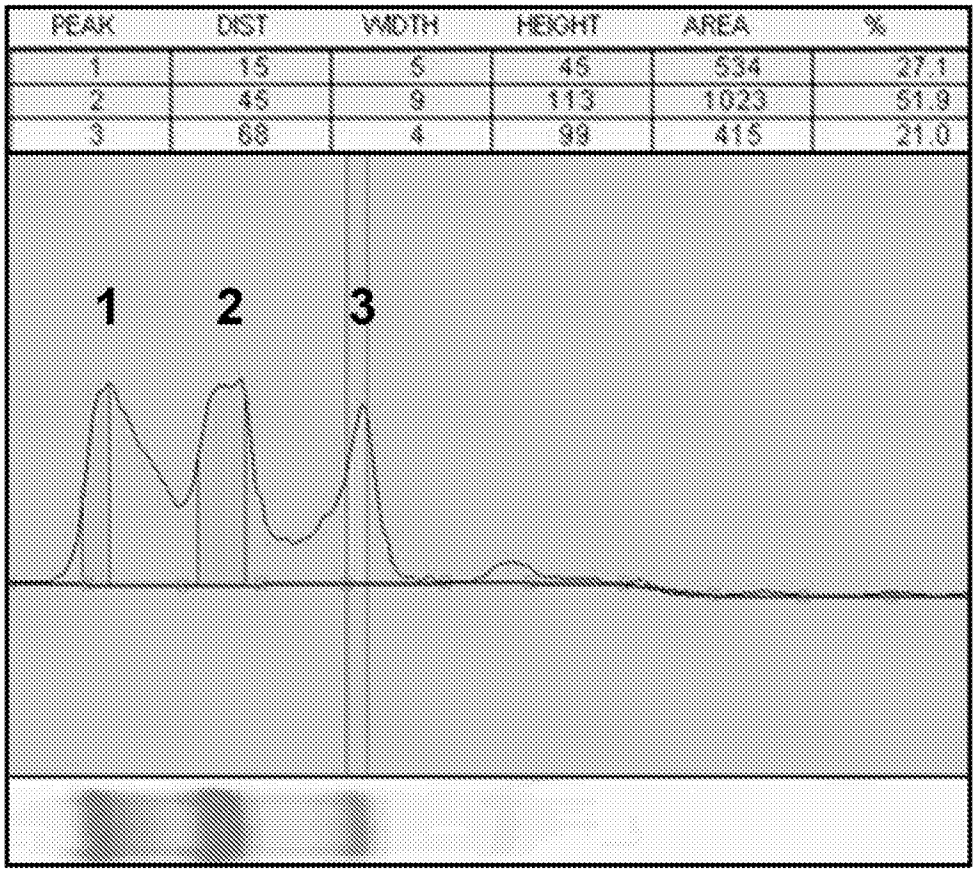

PEAK DIST WIDTH HEIGHT AREA %
1 15 5 45 534 27.1
2 45 9 113 1023 51.9
3 68 4 99 415 21.0
1
2
3

FIG. 19A

CH1																			
IMGT®	1.4	1.3	1.2	1.1	1	2	3	4	5	6	7	8	9	10	11	12	13	14	
EU	118	119	120	121	122	123	124	125	126	127	128	129	130	131	132	133	134	135	
IGHG1	A	S	T	K	G	P	S	V	F	P	L	A	P	S	S	K	S	T	
IGHG2														C		R			
IGHG3														C		R			
IGHG4														C		R			

IMGT®	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	35	36
EU	136	137	138	139	140	141	142	143	144	145	146	147	148	149	150	151	152	153
IGHG1	S	G	G	T	A	A	L	G	C	L	V	K	D	Y	F	P	E	P
IGHG2		E	S															
IGHG3		E	S															
IGHG4		E	S															

IMGT®	37	38	39	40	41	42	43	44	45	45.1	45.2	45.3	77	78	79	80	81	82
EU	154	155	156	157	158	159	160	161	162	163	164	165	166	167	168	169	170	171
IGHG1	V	T	V	S	W	N	S	G	A	L	T	S	G	V	H	T	F	P
IGHG2																		
IGHG3																		
IGHG4																		

FIG. 19A (cont.)

IMGT®	83	84	84.1	84.2	84.3	84.4	85.4	85.3	85.2	85.1	85	86	87	88	89	90	91	92
EU	172	173	174	175	176	177	178	179	180	181	182	183	184	185	186	187	188	189
IGHG1	A	V	L	Q	S	S	G	L	Y	S	L	S	S	V	V	T	V	P
IGHG2																		
IGHG3																		
IGHG4																		

IMGT®	93	94	95	96	98	99	100	101	102	103	104	105	106	107	108	109	110	113
EU	190	191	192	193	194	195	196	197	198	199	200	201	202	203	204	205	206	207
IGHG1	S	S	S	L	G	T	Q	T	Y	I	C	N	V	N	H	K	P	S
IGHG2			N	F						T				D				
IGHG3										T								
IGHG4							K			T				D				

IMGT®	114	115	116	117	118	119	120	121
EU	208	209	210	211	212	213	214	215
IGHG1	N	T	K	V	D	K	K	V
IGHG2							T	
IGHG3							R	
IGHG4							R	

FIG. 19B

IMGT®	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
EU	216	217	218	219	220	221	222	223	224	225	226	227	228	229	230
IGHG1-hinge	E	P	K	S	C	D	K	T	H	T	C	P	P	C	P

FIG. 19C

CH2																			
IMGT®	1.6	1.5	1.4	1.3	1.2	1.1	1	2	3	4	5	6	7	8	9	10	11	12	
EU	231	232	233	234	235	236	237	238	239	240	241	242	243	244	245	246	247	248	
IGHG1	A	P	E	L	L	G	G	P	S	V	F	L	F	P	P	K	P	K	
IGHG2			~	P	V	A													
IGHG3																			
IGHG4				F															
IMGT®	13	14	15	15.1	15.2	16	17	18	19	20	21	22	23	24	25	26	27	28	
EU	249	250	251	252	253	254	255	256	257	258	259	260	261	262	263	264	265	266	
IGHG1	D	T	L	M	I	S	R	T	P	E	V	T	C	V	V	V	D	V	
IGHG2																			
IGHG3																			
IGHG4																			
IMGT®	29	30	31	34	35	36	37	38	39	40	41	42	43	44	45	45.1	45.2	45.3	
EU	267	268	269	270	271	272	273	274	275	276	277	278	279	280	281	282	283	284	
IGHG1	S	H	E	D	P	E	V	K	F	N	W	Y	V	D	G	V	E	V	
IGHG2								Q											
IGHG3								Q		K									
IGHG4		Q						Q											

FIG. 19C (cont.)

IMGT®	45.4	77	78	79	80	81	82	83	84	84.1	84.2	84.3	84.4	85.4	85.3	85.2	85.1	85
EU	285	286	287	288	289	290	291	292	293	294	295	296	297	298	299	300	301	302
IGHG1	H	N	A	K	T	K	P	R	E	E	Q	Y	N	S	T	Y	R	V
IGHG2												F				F		
IGHG3																F		
IGHG4												F						

IMGT®	86	87.	88	89	90	91	92	93	94	95	96	97	98	99	100	101	102	103
EU	303	304	305	306	307	308	309	310	311	312	313	314	315	316	317	318	319	320
IGHG1	V	S	V	L	T	V	L	H	Q	D	W	L	N	G	K	E	Y	K
IGHG2							V											
IGHG3																		
IGHG4																		

IMGT®	104	105	106	107	108	109	110	113	114	115	116	117	118	119	120	121	122	123
EU	321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336	337	338
IGHG1	C	K	V	S	N	K	A	L	P	A	P	I	E	K	T	I	S	K
IGHG2							G											
IGHG3																		
IGHG4							G			S	S							

IMGT®	124	125
EU	339	340
IGHG1	A	K
IGHG2	T	
IGHG3	T	
IGHG4		

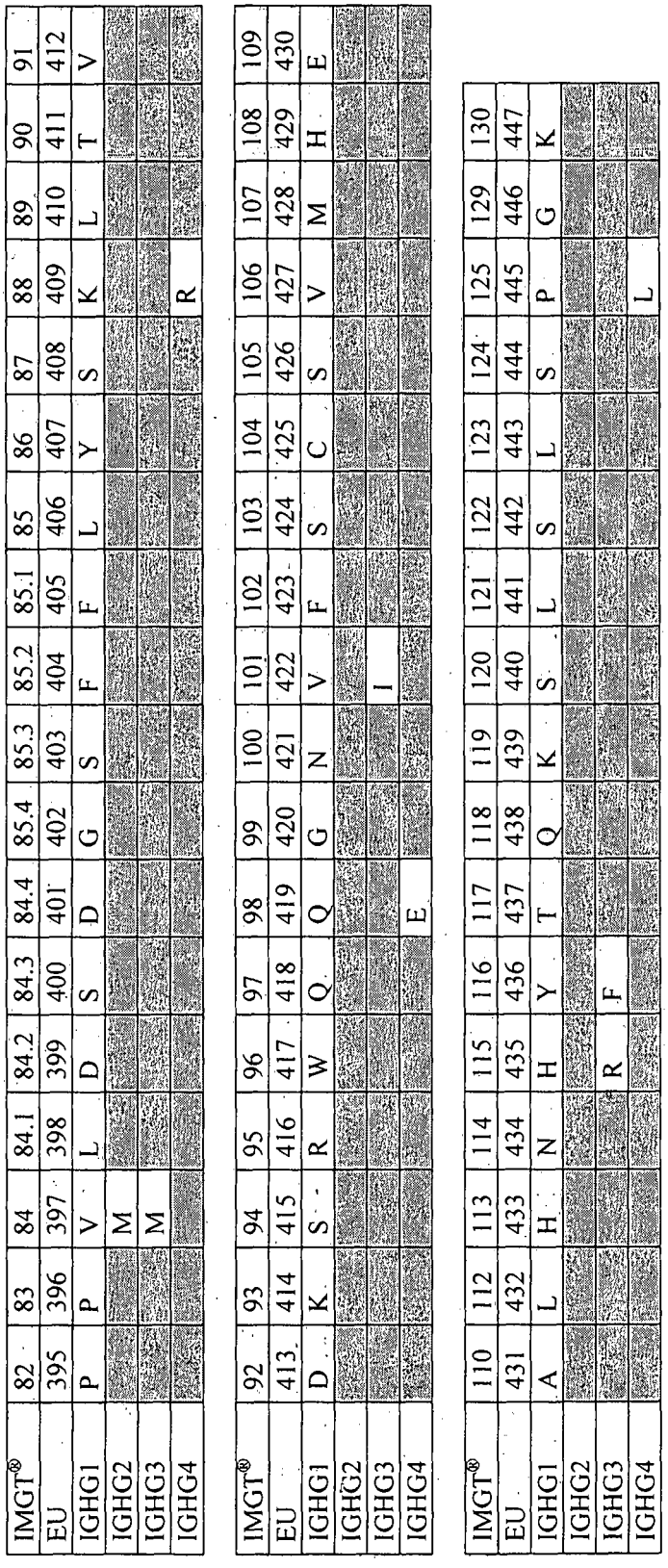
FIG. 19D

CH3																			
IMGT®	1.4	1.3	1.2	1.1	1	2	3	4	5	6	7	8	9	10	11	12	13	14
EU	341	342	343	344	345	346	347	348	349	350	351	352	353	354	355	356	357	358
IGHG1	G	Q	P	R	E	P	Q	V	Y	T	L	P	P	S	R	D	E	L
IGHG2																E		M
IGHG3																E		M
IGHG4															Q	E		M

IMGT®	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	35	36
EU	359	360	361	362	363	364	365	366	367	368	369	370	371	372	373	374	375	376
IGHG1	T	K	N	Q	V	S	L	T	C	L	V	K	G	F	Y	P	S	D
IGHG2																		
IGHG3																		
IGHG4																		

IMGT®	37	38	39	40	41	42	43	44	45	45.1	45.2	45.3	45.4	77	78	79	80	81
EU	377	378	379	380	381	382	383	384	385	386	387	388	389	390	391	392	393	394
IGHG1	I	A	V	E	W	E	S	N	G	Q	P	E	N	N	Y	K	T	T
IGHG2																		
IGHG3								S								N		
IGHG4																		

FIG. 19D (cont.)



IMGT®	82	83	84	84.1	84.2	84.3	84.4	85.4	85.3	85.2	85.1	85	86	87	88	89	90	91
EU	395	396	397	398	399	400	401	402	403	404	405	406	407	408	409	410	411	412
IGHG1	P	P	V	L	D	S	D	G	S	F	F	L	Y	S	K	L	T	V
IGHG2			M															
IGHG3			M															
IGHG4															R			

IMGT®	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107	108	109
EU	413	414	415	416	417	418	419	420	421	422	423	424	425	426	427	428	429	430
IGHG1	D	K	S	R	W	Q	Q	G	N	V	F	S	C	S	V	M	H	E
IGHG2																		
IGHG3										I								
IGHG4							E											

IMGT®	110	112	113	114	115	116	117	118	119	120	121	122	123	124	125	129	130
EU	431	432	433	434	435	436	437	438	439	440	441	442	443	444	445	446	447
IGHG1	A	L	H	N	H	Y	T	Q	K	S	L	S	L	S	P	G	K
IGHG2					R	F											
IGHG3															L		
IGHG4																	

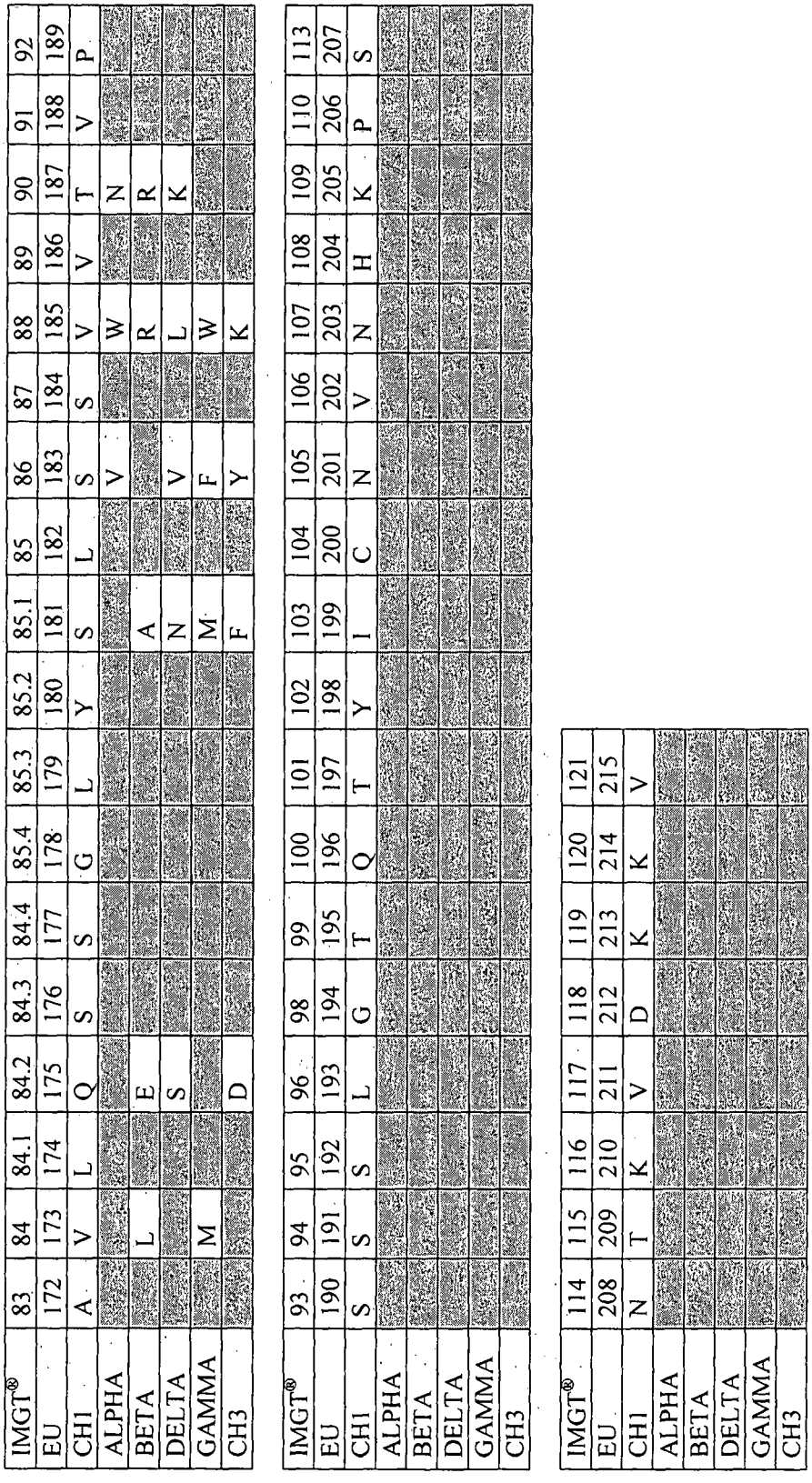
FIG. 20A

IMGT®	1.4	1.3	1.2	1.1	1	2	3	4	5	6	7	8	9	10	11	12	13	14
EU	118	119	120	121	122	123	124	125	126	127	128	129	130	131	132	133	134	135
CH1	A	S	T	K	G	P	S	V	F	P	L	A	P	S	S	K	S	T
ALPHA																		
BETA							E		A		F							
DELTA											M							
GAMMA							K		T									
CH3							Q		Y							D	E	

IMGT®	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	35	36
EU	136	137	138	139	140	141	142	143	144	145	146	147	148	149	150	151	152	153
CH1	S	G	G	T	A	A	L	G	C	L	V	K	D	Y	F	P	E	P
ALPHA						K		V				T						
BETA						T		V				T						
DELTA						N		A					E					
GAMMA						T		L				E	K					
CH3				Q		S		T										

IMGT®	37	38	39	40	41	42	43	44	45	45.1	45.2	45.3	77	78	79	80	81	82
EU	154	155	156	157	158	159	160	161	162	163	164	165	166	167	168	169	170	171
CH1	V	T	V	S	W	N	S	G	A	L	T	S	G	V	H	T	F	P
ALPHA															Y			
BETA																	D	
DELTA															F		A	
GAMMA																	G	
CH3															K		T	

FIG. 20A (cont.)



IMGT®	83	84	84.1	84.2	84.3	84.4	85.4	85.3	85.2	85.1	85	86	87	88	89	90	91	92
EU	172	173	174	175	176	177	178	179	180	181	182	183	184	185	186	187	188	189
CH1	A	V	L	Q	S	S	G	L	Y	S	L	S	S	V	V	T	V	P
ALPHA												V		W		N		
BETA		L		E						A				R		R		
DELTA				S						N		V		L		K		
GAMMA		M								M		F		W				
CH3				D						F		Y		K				

IMGT®	93	94	95	96	98	99	100	101	102	103	104	105	106	107	108	109	110	113
EU	190	191	192	193	194	195	196	197	198	199	200	201	202	203	204	205	206	207
CH1	S	S	S	L	G	T	Q	T	Y	I	C	N	V	N	H	K	P	S
ALPHA																		
BETA																		
DELTA																		
GAMMA																		
CH3																		

IMGT®	114	115	116	117	118	119	120	121
EU	208	209	210	211	212	213	214	215
CH1	N	T	K	V	D	K	K	V
ALPHA								
BETA								
DELTA								
GAMMA								
CH3								

FIG. 20B

IMGT®	1.6	1.5	1.4	1.3	1.2	1.1	1	2	3	4	5	6	7	8	9	10	11	12
EU	231	232	233	234	235	236	237	238	239	240	241	242	243	244	245	246	247	248
CH2	A	P	E	L	L	G	G	P	S	V	F	L	F	P	P	K	P	K
ALPHA																		
BETA									E		A							
DELTA													M					
GAMMA									K		T							
CH3											Y		L					

IMGT®	13	14	15	15.1	15.2	16	17	18	19	20	21	22	23	24	25	26	27	28
EU	249	250	251	252	253	254	255	256	257	258	259	260	261	262	263	264	265	266
CH2	D	T	L	M	I	S	R	T	P	E	V	T	C	V	V	V	D	V
ALPHA										K		V		V	V	T		
BETA										T		V				T		
DELTA										N		A					E	
GAMMA										T		L				E	K	
CH3										S								

IMGT®	29	30	31	34	35	36	37	38	39	40	41	42	43	44	45	45.1	45.2	45.3
EU	267	268	269	270	271	272	273	274	275	276	277	278	279	280	281	282	283	284
CH2	S	H	E	D	P	E	V	K	F	N	W	Y	V	D	G	V	E	V
ALPHA																		
BETA																		
DELTA																		
GAMMA																		
CH3																		

FIG. 20B (cont.)

IMGT®	45.4	77	78	79	80	81	82	83	84	84.1	84.2	84.3	84.4	85.4	85.3	85.2	85.1	85
EU	285	286	287	288	289	290	291	292	293	294	295	296	297	298	299	300	301	302
CH2	H	N	A	K	T	K	P	R	E	E	Q	Y	N	S	T	Y	R	V
ALPHA				Y													S	
BETA						D			L		E						A	
DELTA				F		A					S						N	
GAMMA						G		M									M	
CH3								T				D					F	

IMGT®	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100	101	102	103
EU	303	304	305	306	307	308	309	310	311	312	313	314	315	316	317	318	319	320
CH2	V	S	V	L	T	V	L	H	Q	D	W	L	N	G	K	E	Y	K
ALPHA			W		N													
BETA	S		R		R													
DELTA			L		K													
GAMMA	F		W															
CH3	Y		K															

FIG. 20B (cont.)

IMGT®	104	105	106	107	108	109	110	113	114	115	116	117	118	119	120	121	122	123
EU	321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336	337	338
CH2	C	K	V	S	N	K	A	L	P	A	P	I	E	K	T	I	S	K
ALPHA																		
BETA																		
DELTA																		
GAMMA																		
CH3																		

IMGT®	124	125
EU	339	340
CH2	A	K
ALPHA		
BETA		
DELTA		
GAMMA		
CH3		

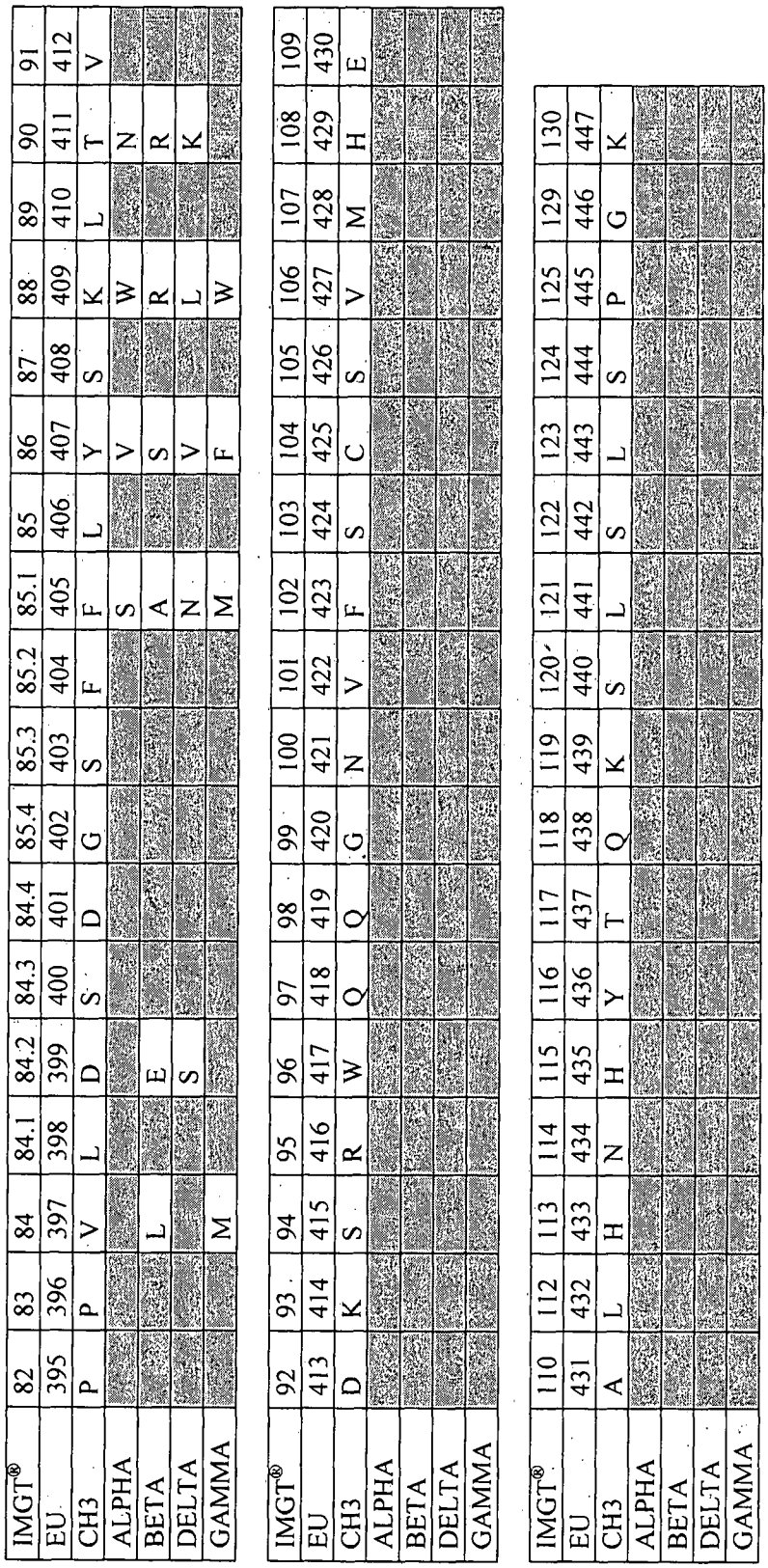
FIG. 20C

IMGT®	1.4	1.3	1.2	1.1	1	2	3	4	5	6	7	8	9	10	11	12	13	14
EU	341	342	343	344	345	346	347	348	349	350	351	352	353	354	355	356	357	358
CH3	G	Q	P	R	E	P	Q	V	Y	T	L	P	P	S	R	D	E	L
ALPHA																		
BETA							E		A		F							
DELTA											M							
GAMMA							K		T									

IMGT®	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	35	36
EU	359	360	361	362	363	364	365	366	367	368	369	370	371	372	373	374	375	376
CH3	T	K	N	Q	V	S	L	T	C	L	V	K	G	F	Y	P	S	D
ALPHA						K		V				T						
BETA						T		V				T						
DELTA						N		A					E					
GAMMA						T		L				E	K					

IMGT®	37	38	39	40	41	42	43	44	45	45.1	45.2	45.3	45.4	77	78	79	80	81
EU	377	378	379	380	381	382	383	384	385	386	387	388	389	390	391	392	393	394
CH3	I	A	V	E	W	E	S	N	G	Q	P	E	N	N	Y	K	T	T
ALPHA																Y		
BETA																		D
DELTA																F		A
GAMMA																		G

FIG. 20C (cont.)

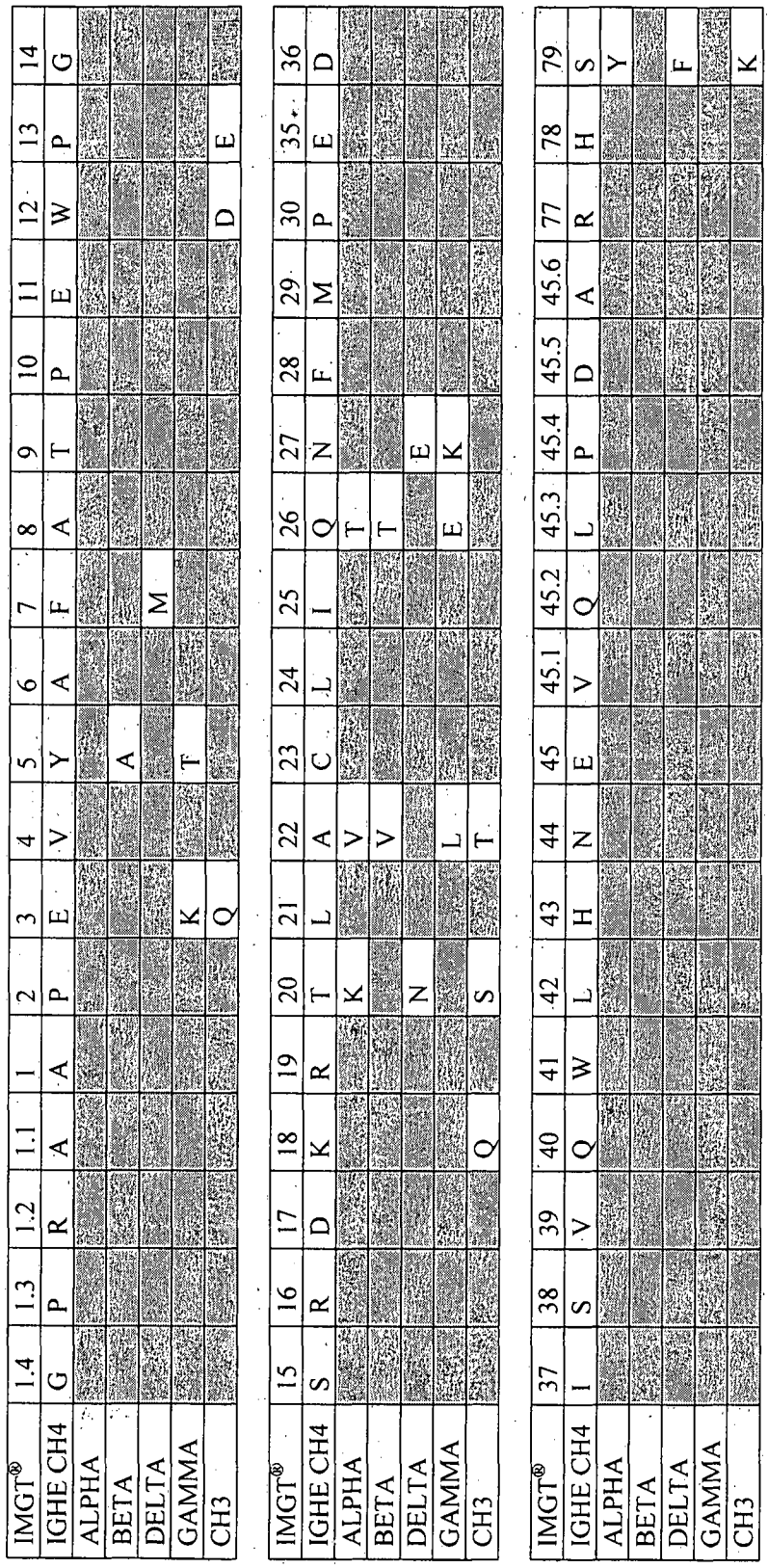


IMGT®	82	83	84	84.1	84.2	84.3	84.4	85.4	85.3	85.2	85.1	85	86	87	88	89	90	91
EU	395	396	397	398	399	400	401	402	403	404	405	406	407	408	409	410	411	412
CH3	P	P	V	L	D	S	D	G	S	F	F	L	Y	S	K	L	T	V
ALPHA											S		V		W		N	
BETA			L		E						A		S		R		R	
DELTA					S						N		V		L		K	
GAMMA			M								M		F		W			

IMGT®	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107	108	109
EU	413	414	415	416	417	418	419	420	421	422	423	424	425	426	427	428	429	430
CH3	D	K	S	R	W	Q	Q	G	N	V	F	S	C	S	V	M	H	E
ALPHA																		
BETA																		
DELTA																		
GAMMA																		

IMGT®	110	112	113	114	115	116	117	118	119	120'	121	122	123	124	125	129	130
EU	431	432	433	434	435	436	437	438	439	440	441	442	443	444	445	446	447
CH3	A	L	H	N	H	Y	T	Q	K	S	L	S	L	S	P	G	K
ALPHA																	
BETA																	
DELTA																	
GAMMA																	

FIG. 21A

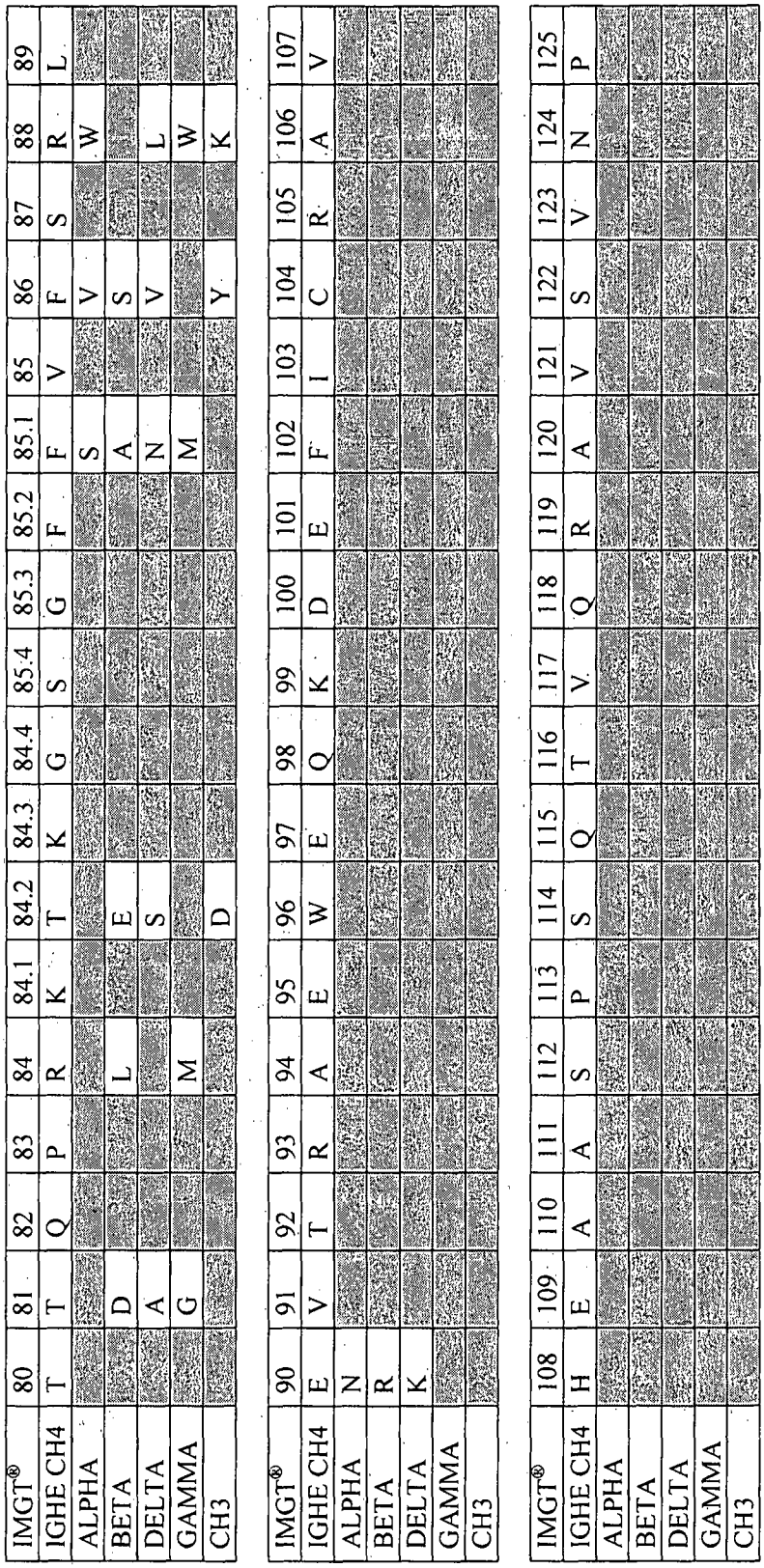


IMGT®	1.4	1.3	1.2	1.1	1	2	3	4	5	6	7	8	9	10	11	12	13	14
IGHE CH4	G	P	R	A	A	P	E	V	Y	A	F	A	T	P	E	W	P	G
ALPHA																		
BETA									A									
DELTA									A		M							
GAMMA							K		T									
CH3							Q								D	E		

IMGT®	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	35	36
IGHE CH4	S	R	D	K	R	T	L	A	C	L	I	Q	N	F	M	P	E	D
ALPHA						K		V				T						
BETA								V				T						
DELTA						N							E					
GAMMA								L				E	K					
CH3				Q		S		T										

IMGT®	37	38	39	40	41	42	43	44	45	45.1	45.2	45.3	45.4	45.5	45.6	77	78	79
IGHE CH4	I	S	V	Q	W	L	H	N	E	V	Q	L	P	D	A	R	H	S
ALPHA																		Y
BETA																		
DELTA																		F
GAMMA																		
CH3																		K

FIG. 21A (cont.)

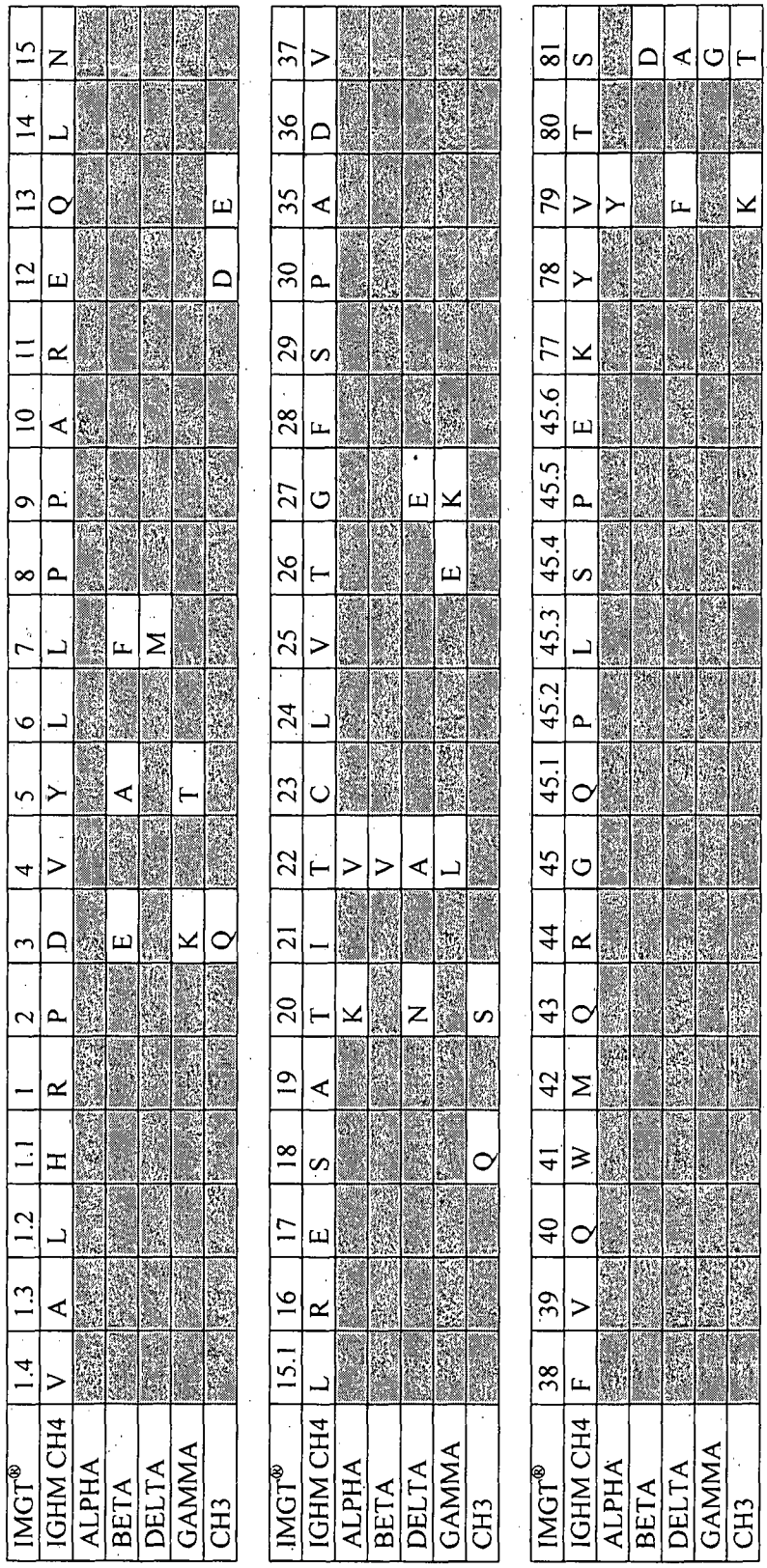


IMGT®	80	81	82	83	84	84.1	84.2	84.3	84.4	85.4	85.3	85.2	85.1	85	86	87	88	89
IGHE CH4	T	T	Q	P	R	K	T	K	G	S	G	F	F	V	F	S	R	L
ALPHA													S		V		W	
BETA		D			L		E						A		S			
DELTA		A					S						N		V		L	
GAMMA		G			M								M				W	
CH3							D								Y		K	

IMGT®	90	91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107
IGHE CH4	E	V	T	R	A	E	W	E	Q	K	D	E	F	I	C	R	A	V
ALPHA	N																	
BETA	R																	
DELTA	K																	
GAMMA																		
CH3																		

IMGT®	108	109	110	111	112	113	114	115	116	117	118	119	120	121	122	123	124	125
IGHE CH4	H	E	A	A	S	P	S	Q	T	V	Q	R	A	V	S	V	N	P
ALPHA																		
BETA																		
DELTA																		
GAMMA																		
CH3																		

FIG. 21B



IMGT®	1.4	1.3	1.2	1.1	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
IGHM CH4	V	A	L	H	R	P	D	V	Y	L	L	P	P	A	R	E	Q	L	N
ALPHA																			
BETA							E		A		F								
DELTA											M								
GAMMA							K		T										
CH3							Q									D	E		

IMGT®	15.1	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	35	36	37
IGHM CH4	L	R	E	S	A	T	I	T	C	L	V	T	G	F	S	P	A	D	V
ALPHA						K		V											
BETA								V											
DELTA						N		A					E						
GAMMA								L				E	K						
CH3				Q		S													

IMGT®	38	39	40	41	42	43	44	45	45.1	45.2	45.3	45.4	45.5	45.6	77	78	79	80	81
IGHM CH4	F	V	Q	W	M	Q	R	G	Q	P	L	S	P	E	K	Y	V	T	S
ALPHA																	Y		D
BETA																			
DELTA																	F		A
GAMMA																			G
CH3																	K		T

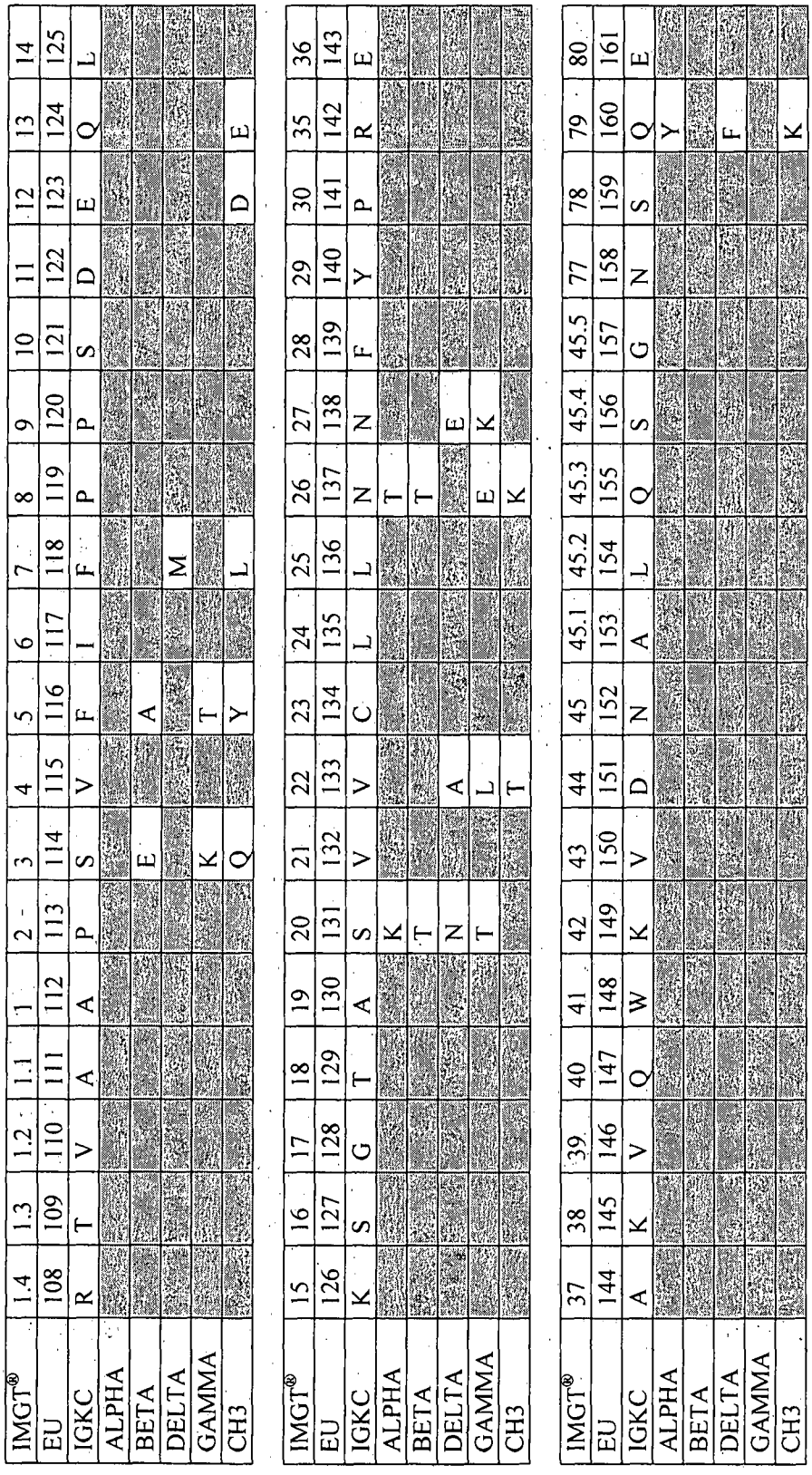
FIG. 21B (cont.)

IMGT®	82	83	84	84.1	84.2	84.3	84.4	84.5	85.5	85.4	85.3	85.2	85.1	85	86	87	88	89	90
IGHM CH4	A	P	M	P	E	P	Q	A	P	G	R	Y	F	A	H	S	I	L	T
ALPHA													S		V		W		N
BETA			L										A		S		R		R
DELTA					S								N		V		L		K
GAMMA													M		F		W		
CH3					D										Y		K		

IMGT®	91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107	108	109
IGHM CH4	V	S	E	E	E	W	N	T	G	E	T	Y	T	C	V	V	A	H	E
ALPHA																			
BETA																			
DELTA																			
GAMMA																			
CH3																			

IMGT®	110	112	113	114	115	116	117	118	119	120	121	122	123	124	125
IGHM CH4	A	L	P	N	R	V	T	E	R	T	V	D	K	S	T
ALPHA															
BETA															
DELTA															
GAMMA															
CH3															

FIG. 22



IMGT®	1.4	1.3	1.2	1.1	1	2	3	4	5	6	7	8	9	10	11	12	13	14
EU	108	109	110	111	112	113	114	115	116	117	118	119	120	121	122	123	124	125
IGKC	R	T	V	A	A	P	S	V	F	I	F	P	P	S	D	E	Q	L
ALPHA																		
BETA							E		A									
DELTA											M							
GAMMA							K		T									
CH3							Q		Y		L				D	E		

IMGT®	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	35	36
EU	126	127	128	129	130	131	132	133	134	135	136	137	138	139	140	141	142	143
IGKC	K	S	G	T	A	S	V	V	C	L	L	N	N	F	Y	P	R	E
ALPHA						K						T						
BETA						T						T						
DELTA						N		A					E					
GAMMA						T		L				E	K					
CH3								T				K						

IMGT®	37	38	39	40	41	42	43	44	45	45.1	45.2	45.3	45.4	45.5	77	78	79	80
EU	144	145	146	147	148	149	150	151	152	153	154	155	156	157	158	159	160	161
IGKC	A	K	V	Q	W	K	V	D	N	A	L	Q	S	G	N	S	Q	E
ALPHA																	Y	
BETA																		
DELTA																	F	
GAMMA																		
CH3																	K	

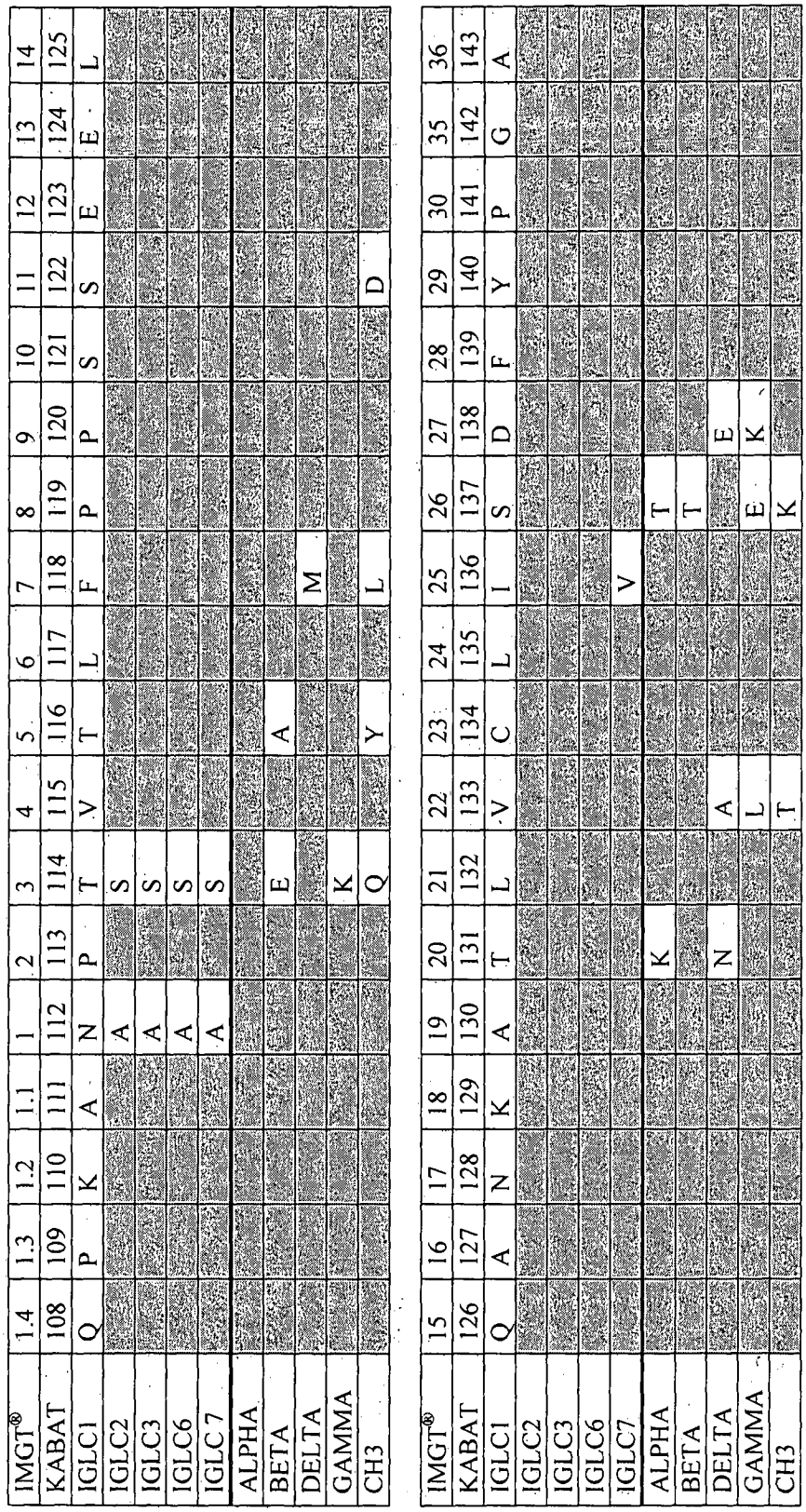
FIG. 22 (cont.)

IMGT®	81	82	83	84	84.1	84.2	84.3	84.4	84.5	85.4	85.3	85.2	85.1	85	86	87	88	89
EU	162	163	164	165	166	167	168	169	170	171	172	173	174	175	176	177	178	179
IGKC	S	V	T	E	Q	D	S	K	D	S	T	Y	S	L	S	S	T	L
ALPHA															V		W	
BETA	D		L			E							A				R	
DELTA	A					S							N		V		L	
GAMMA	G		M										M		F		W	
CH3	T		V			S							F		Y		K	

IMGT®	90	91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107
EU	180	181	182	183	184	185	186	187	188	189	190	191	192	193	194	195	196	197
IGKC	T	L	S	K	A	D	Y	E	K	H	K	V	Y	A	C	E	V	T
ALPHA	N																	
BETA	R																	
DELTA	K																	
GAMMA																		
CH3																		

IMGT®	108	109	110	113	114	115	116	117	118	119	120	121	122	123	124	125	126
EU	198	199	200	201	202	203	204	205	206	207	208	209	210	211	212	213	214
IGKC	H	Q	G	L	S	S	P	V	T	K	S	F	N	R	G	E	C
ALPHA																	
BETA																	
DELTA																	
GAMMA																	
CH3																	

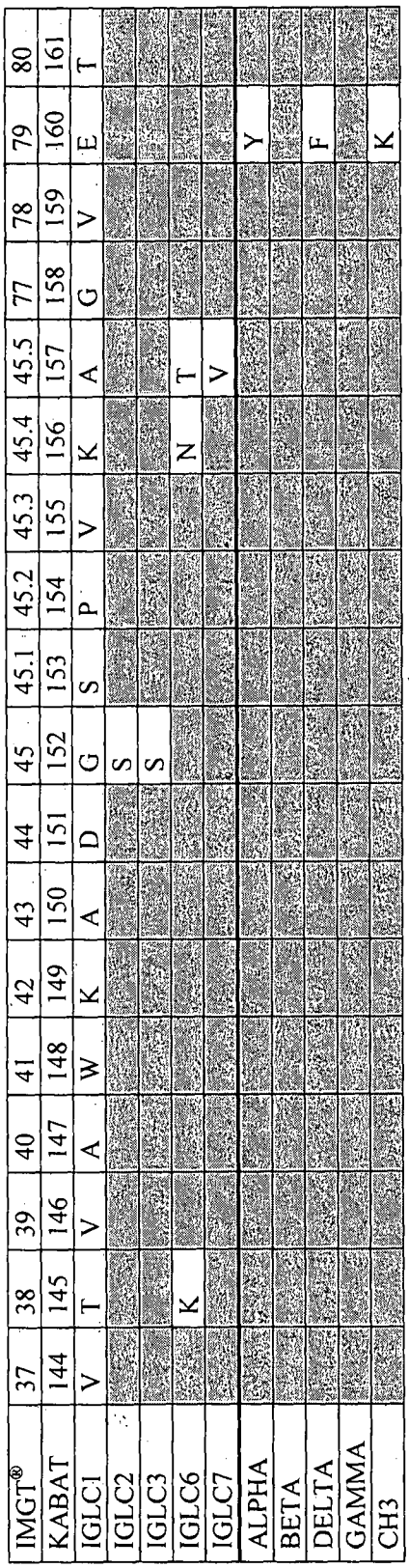
FIG. 23A



IMGT®	1.4	1.3	1.2	1.1	1	2	3	4	5	6	7	8	9	10	11	12	13	14
KABAT	108	109	110	111	112	113	114	115	116	117	118	119	120	121	122	123	124	125
IGLC1	Q	P	K	A	N	P	T	V	T	L	F	P	P	S	S	E	E	L
IGLC2					A		S											
IGLC3					A		S											
IGLC6					A		S											
IGLC7					A		S											
ALPHA																		
BETA							E		A									
DELTA											M							
GAMMA							K											
CH3							Q		Y		L				D			

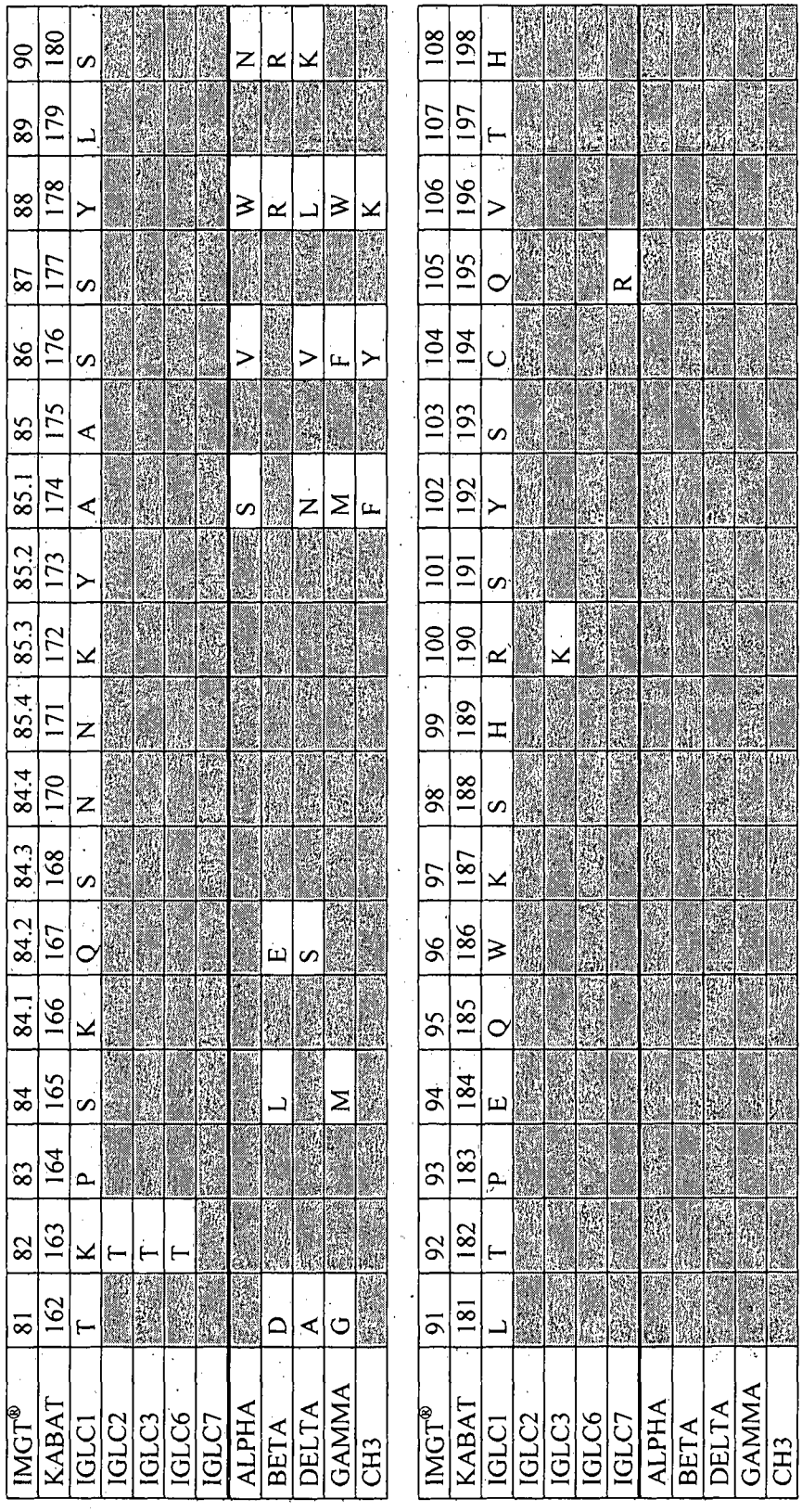
IMGT®	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	35	36
KABAT	126	127	128	129	130	131	132	133	134	135	136	137	138	139	140	141	142	143
IGLC1	Q	A	N	K	A	T	L	V	C	L	I	S	D	F	Y	P	G	A
IGLC2																		
IGLC3																		
IGLC6																		
IGLC7											V							
ALPHA						K						T						
BETA												T						
DELTA						N		A					E					
GAMMA								L				E	K					
CH3								T				K						

FIG. 23A (cont.)



IMGT®	37	38	39	40	41	42	43	44	45	45.1	45.2	45.3	45.4	45.5	77	78	79	80
KABAT	144	145	146	147	148	149	150	151	152	153	154	155	156	157	158	159	160	161
IGLC1	V	T	V	A	W	K	A	D	G	S	P	V	K	A	G	V	E	T
IGLC2									S									
IGLC3									S									
IGLC6		K											N	T				
IGLC7														V				
ALPHA																	Y	
BETA																		
DELTA																	F	
GAMMA																		
CH3																	K	

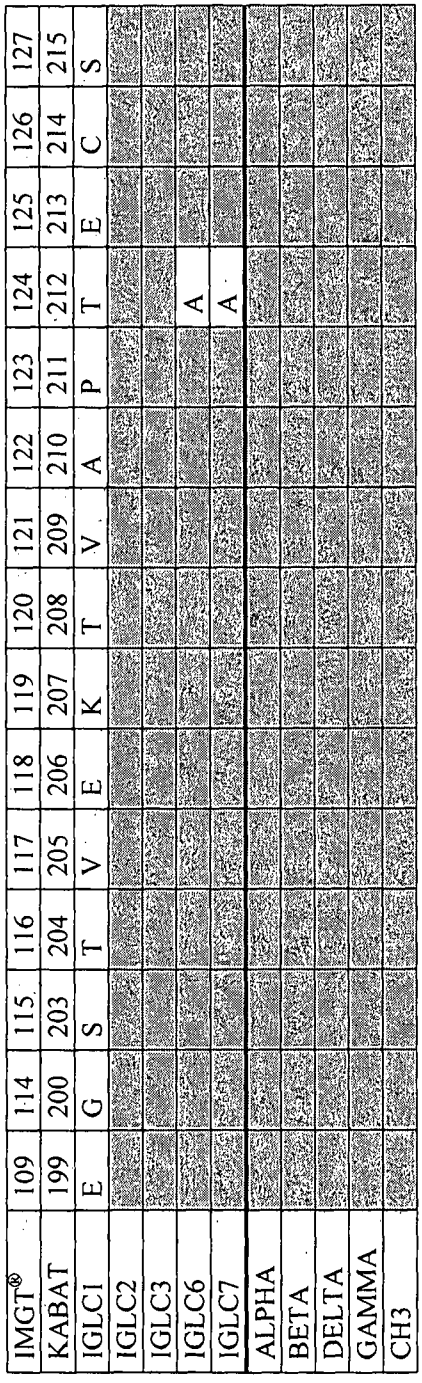
FIG. 23B



IMGT®	81	82	83	84	84.1	84.2	84.3	84.4	85.4	85.3	85.2	85.1	85	86	87	88	89	90
KABAT	162	163	164	165	166	167	168	170	171	172	173	174	175	176	177	178	179	180
IGLC1	T	K	P	S	K	Q	S	N	N	K	Y	A	A	S	S	Y	L	S
IGLC2		T																
IGLC3		T																
IGLC6		T																
IGLC7																		
ALPHA												S		V		W		N
BETA	D			L		E										R		R
DELTA	A					S						N		V		L		K
GAMMA	G			M								M		F		W		
CH3												F		Y		K		

IMGT®	91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107	108
KABAT	181	182	183	184	185	186	187	188	189	190	191	192	193	194	195	196	197	198
IGLC1	L	T	P	E	Q	W	K	S	H	R	S	Y	S	C	Q	V	T	H
IGLC2										K								
IGLC3																		
IGLC6																		
IGLC7															R			
ALPHA																		
BETA																		
DELTA																		
GAMMA																		
CH3																		

FIG. 23B (cont.)



IMGT®	109	114	115	116	117	118	119	120	121	122	123	124	125	126	127
KABAT	199	200	203	204	205	206	207	208	209	210	211	212	213	214	215
IGLC1	E	G	S	T	V	E	K	T	V	A	P	T	E	C	S
IGLC2															
IGLC3															
IGLC6												A			
IGLC7												A			
ALPHA															
BETA															
DELTA															
GAMMA															
CH3															

FIG. 23B

IMGT®	81	82	83	84	84.1	84.2	84.3	84.4	85.4	85.3	85.2	85.1	85	86	87	88	89	90
KABAT	162	163	164	165	166	167	168	170	171	172	173	174	175	176	177	178	179	180
IGLC1	T	K	P	S	K	Q	S	N	N	K	Y	A	A	S	S	Y	L	S
IGLC2		T																
IGLC3		T																
IGLC6		T																
IGLC7																		
ALPHA												S		V		W		N
BETA	D			L		E										R		R
DELTA	A					S						N		V		L		K
GAMMA	G			M								M		F		W		
CH3												F		Y		K		

IMGT®	91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107	108
KABAT	181	182	183	184	185	186	187	188	189	190	191	192	193	194	195	196	197	198
IGLC1	L	T	P	E	Q	W	K	S	H	R	S	Y	S	C	Q	V	T	H
IGLC2																		
IGLC3										K								
IGLC6																		
IGLC7															R			
ALPHA																		
BETA																		
DELTA																		
GAMMA																		
CH3																		

IMGT®	109	114	115	116	117	118	119	120	121	122	123	124	125	126	127
KABAT	199	200	203	204	205	206	207	208	209	210	211	212	213	214	215
IGLC1	E	G	S	T	V	E	K	T	V	A	P	T	E	C	S
IGLC2															
IGLC3															
IGLC6												A			
IGLC7												A			
ALPHA															
BETA															
DELTA															
GAMMA															
CH3															

FIG. 24

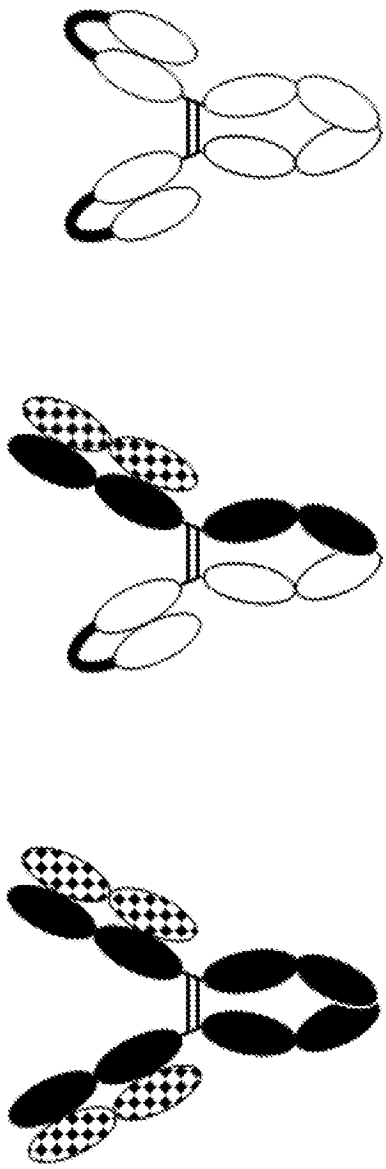

scFv(anti-HER2)-BT beta chain
VH(anti-EGFR)-CH1-BT alpha chain
VL(anti-EGFR)-CK chain
(Gly4Ser)3 linker

FIG. 25

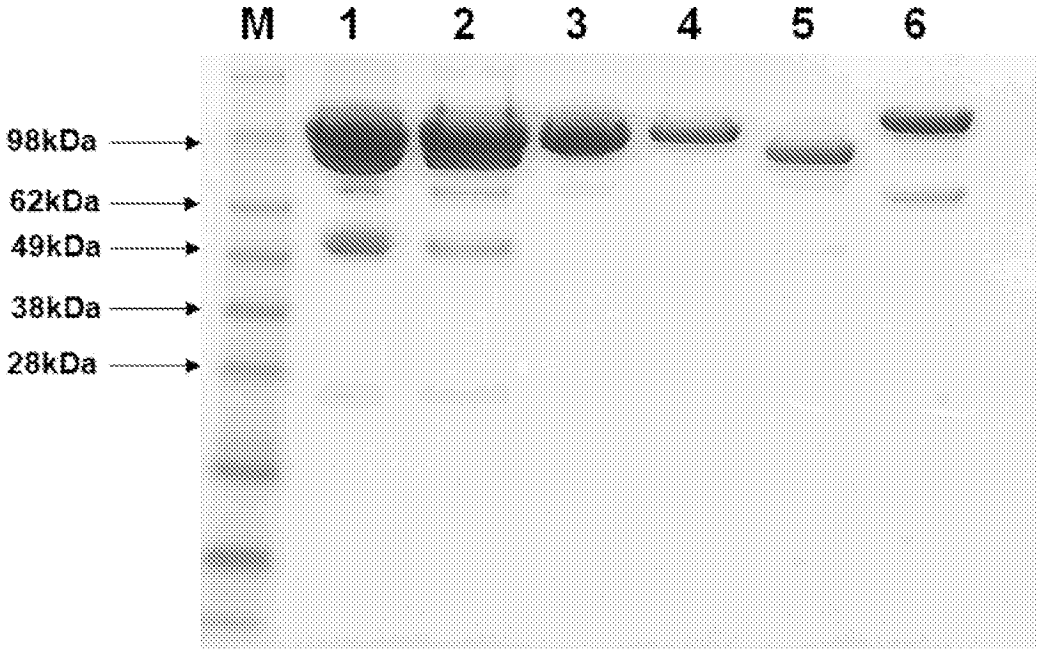


FIG. 26

PEAK	DST	WIDTH	HEIGHT	AREA	%
1	19	4	43	193	7.1
2	31	15	95	1344	49.5
3	47	12	91	1176	43.3

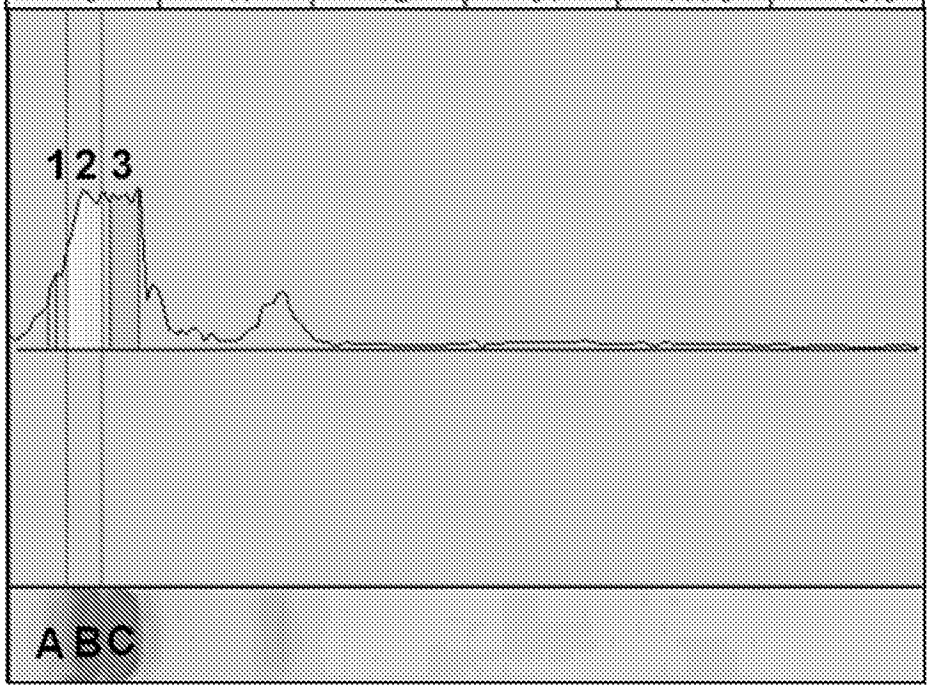


FIG. 27

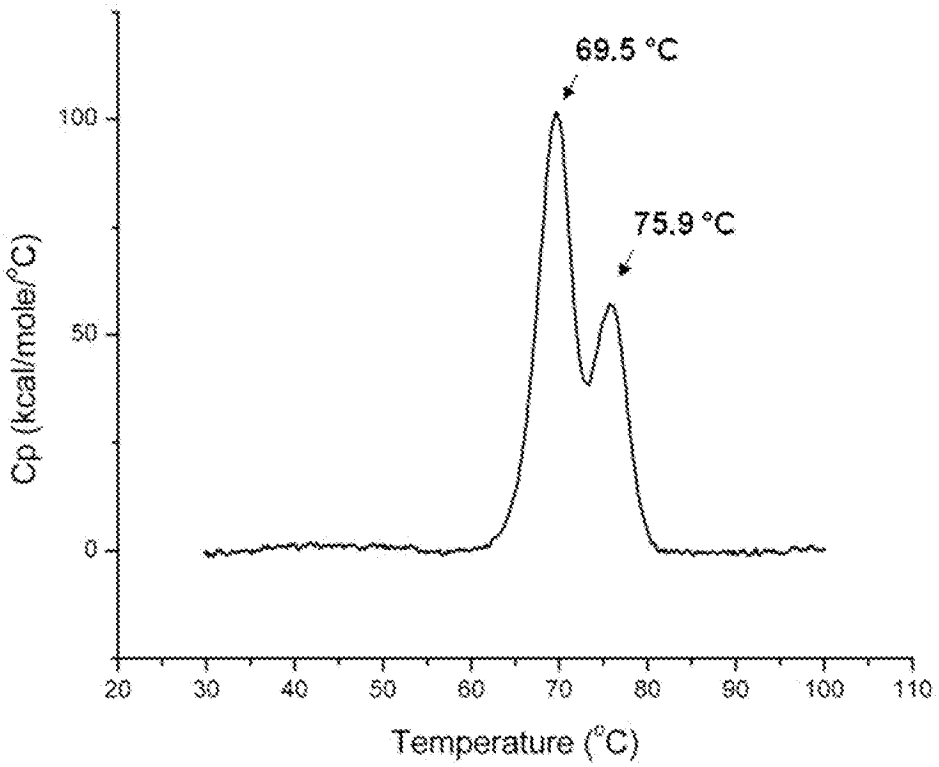

69.5 °C
75.9 °C
Cp (kcal/mole/°C)
100
50
0
20
30
40
50
60
70
80
90
100
110
Temperature (°C)

FIG. 28A

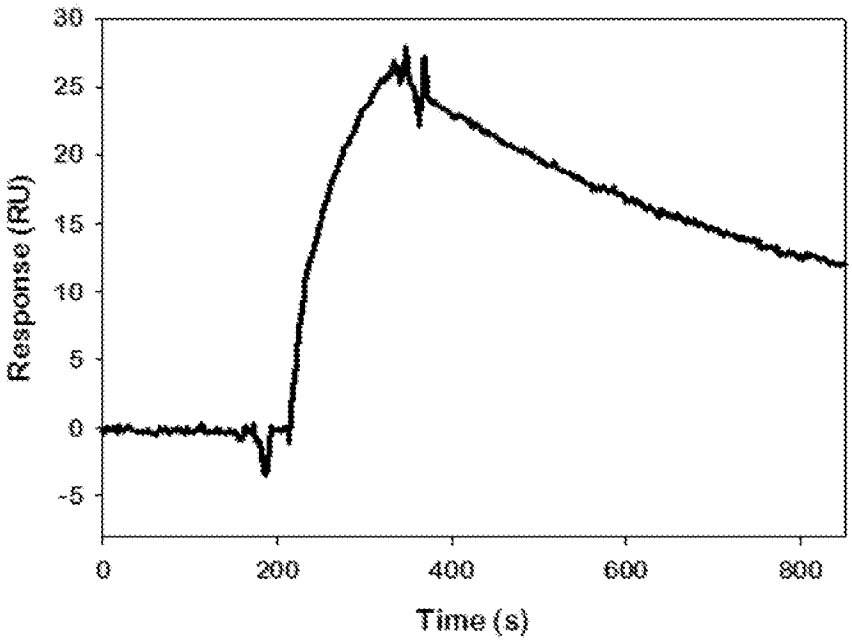
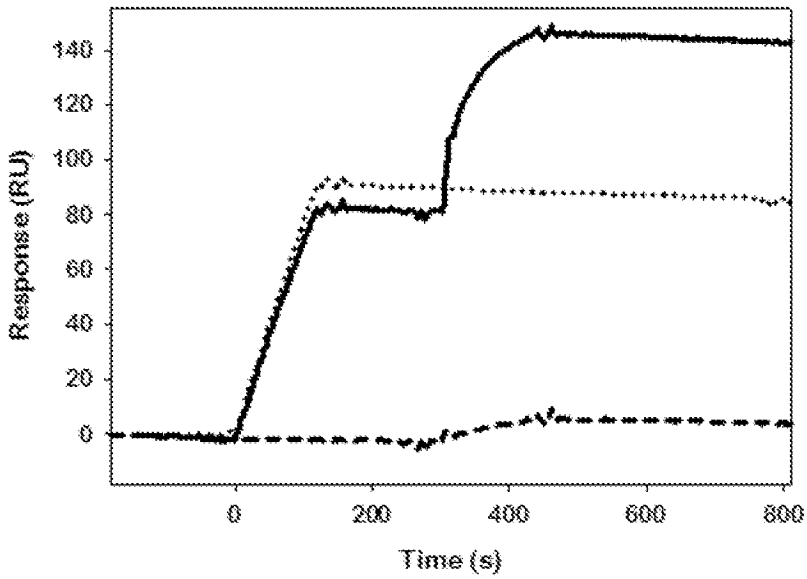

30
25
20
15
10
5
0
-5
Response (RU)
0
200
400
600
800
Time (s)
Immobilized human EGFR-Fc + (Erbitux-BT alpha_Herceptin scFv-BT beta)

FIG. 28B



Immobilized human HER2-Fc
+ human EGFR-Fc
+ (Erbitux-BT alpha_Herceptin scFv-BT beta)

Immobilized human HER2-Fc
+ (Erbitux-BT alpha_Herceptin scFv-BT beta)

Immobilized human HER2-Fc
+ human EGFR-Fc

FIG. 29

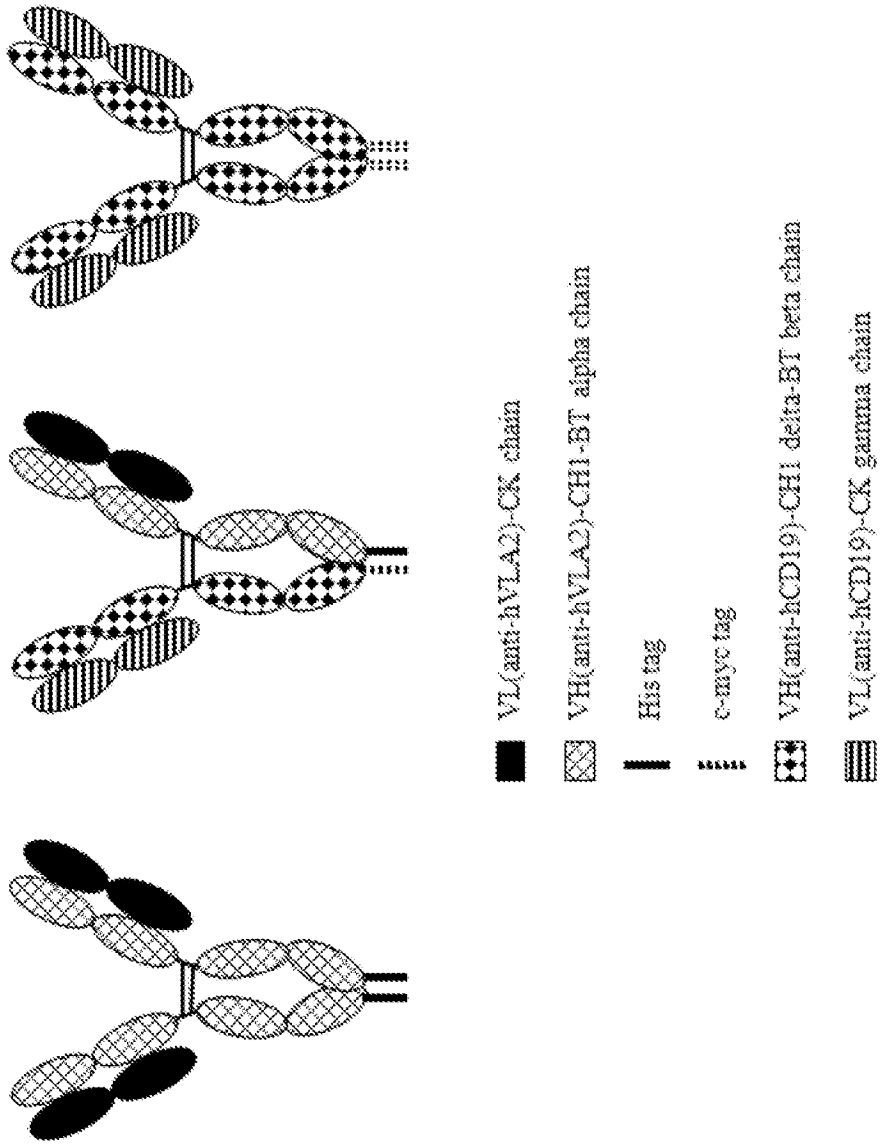

VL(anti-hVLA2)-CK chain
VH(anti-hVLA2)-CH1-BT alpha chain
His tag
c-myc tag
VH(anti-hCD19)-CH1 delta-BT beta chain
VL(anti-hCD19)-CK gamma chain

FIG. 30

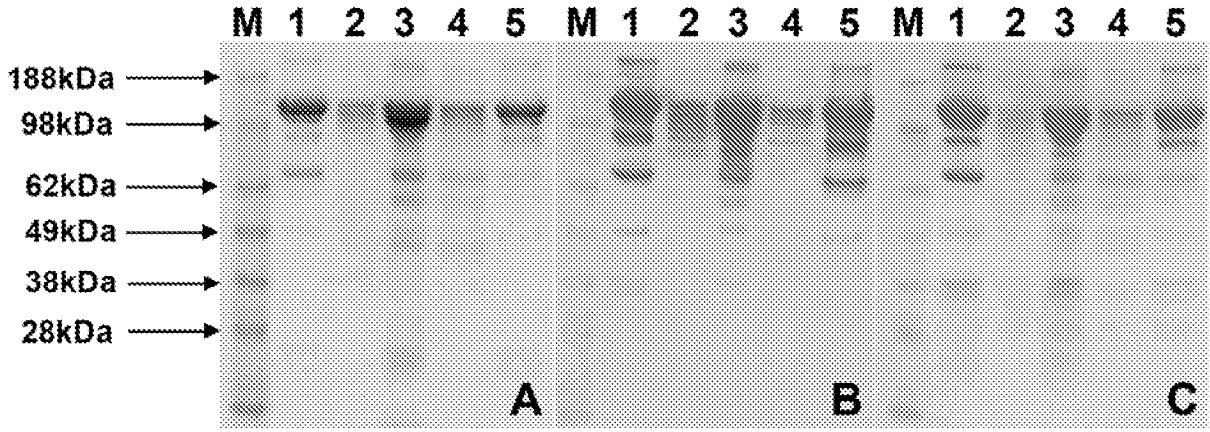

M 1 2 3 4 5 M 1 2 3 4 5 M 1 2 3 4 5
188kDa
98kDa
62kDa
49kDa
38kDa
28kDa
A
B
C

FIG. 31

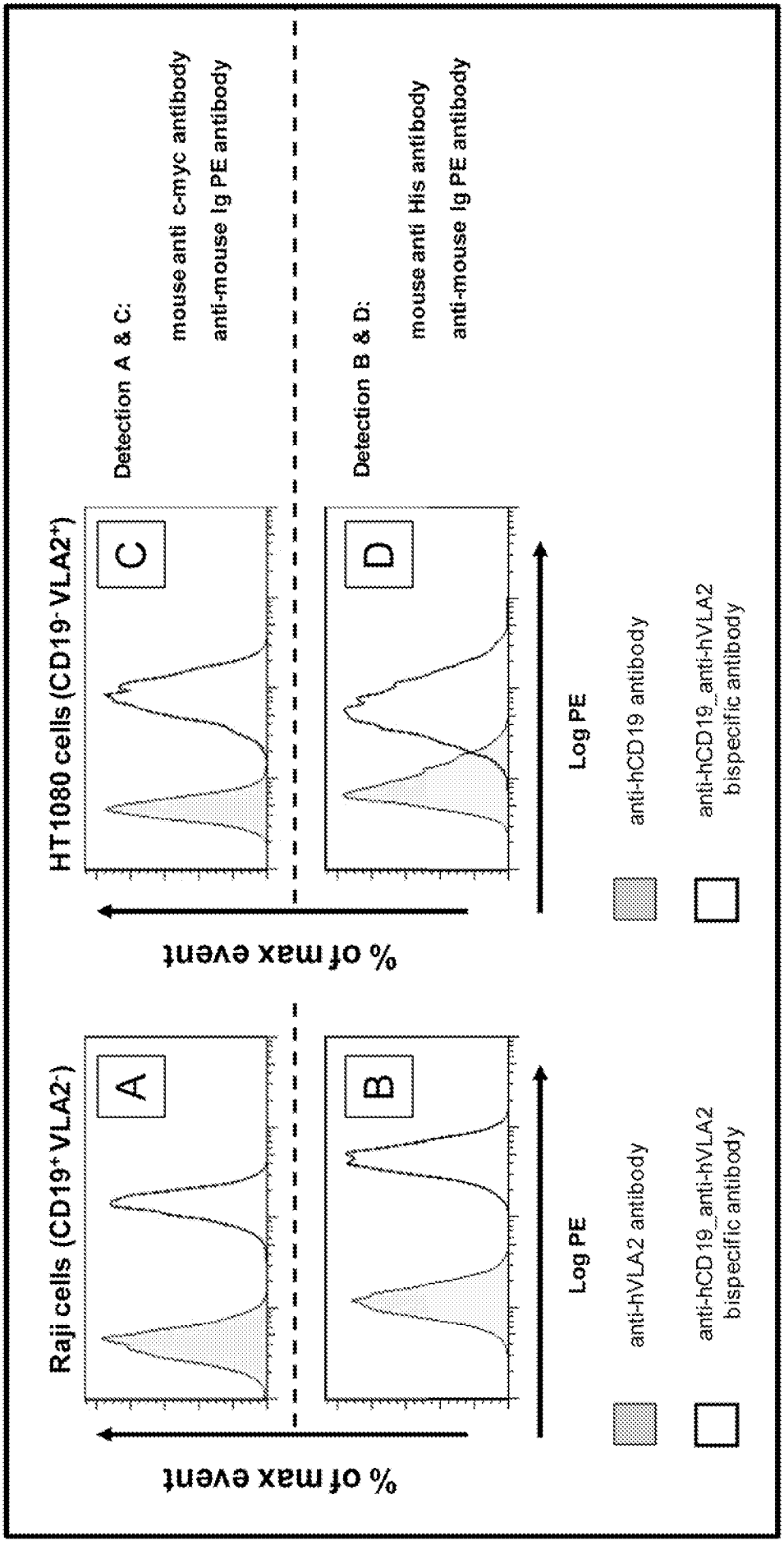

Raji cells (CD19+VLA2-)
A
B
% of max event
Log PE
anti-hVLA2 antibody
anti-hCD19_anti-hVLA2 bispecific antibody
HT1080 cells (CD19-VLA2+)
C
D
Detection A & C:
mouse anti c-myc antibody
anti-mouse Ig PE antibody
Detection B & D:
mouse anti His antibody
anti-mouse Ig PE antibody
anti-hCD19 antibody
anti-hCD19_anti-hVLA2 bispecific antibody

FIG. 32

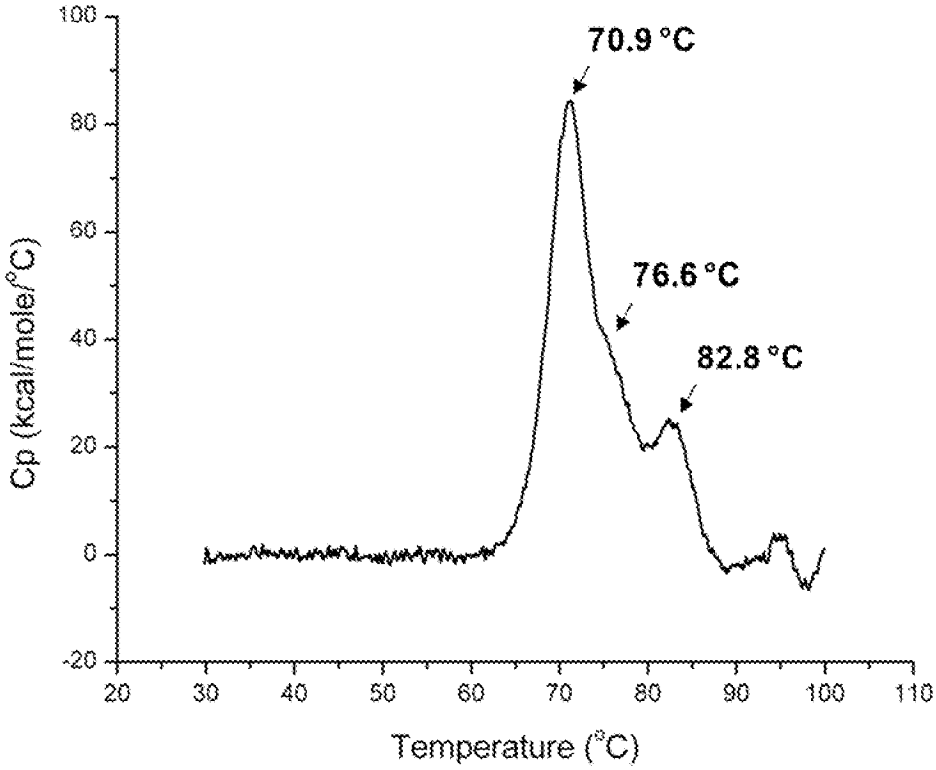

70.9 °C
76.6 °C
82.8 °C
Cp (kcal/mole/°C)
Temperature (°C)
100
80
60
40
20
0
-20
20
30
40
50
60
70
80
90
100
110

FIG. 33

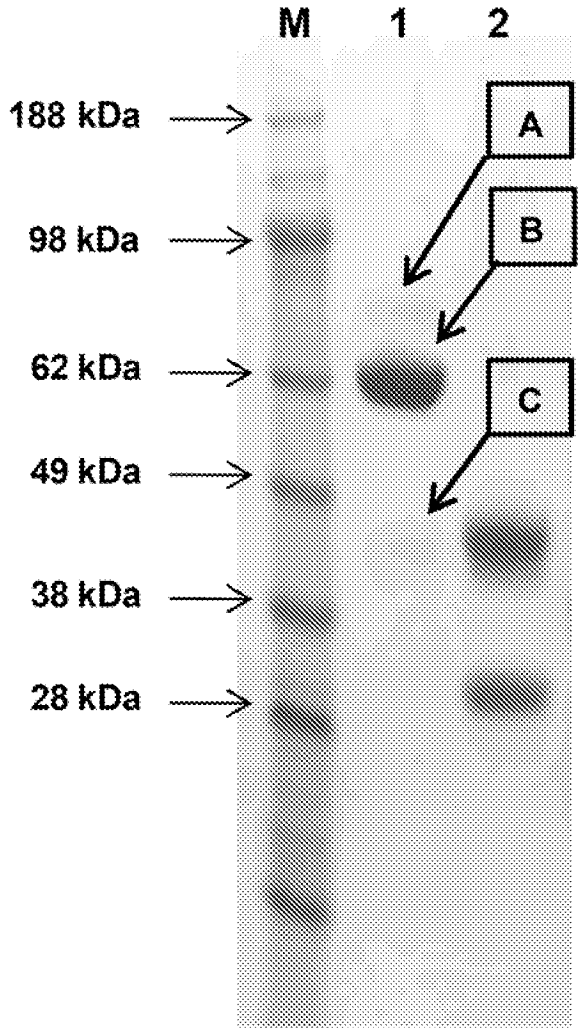
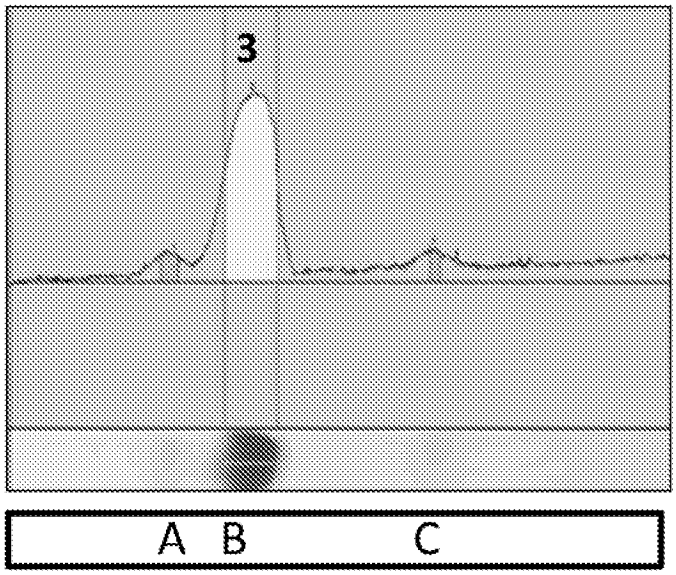


FIG. 34



LANE	PEAK	DIST	WIDTH	HEIGHT	AREA	%
1	1	210	22	23	550	5.4
1	2	309	67	153	9206	91.0
1	3	537	13	28	360	3.6

FIG. 35

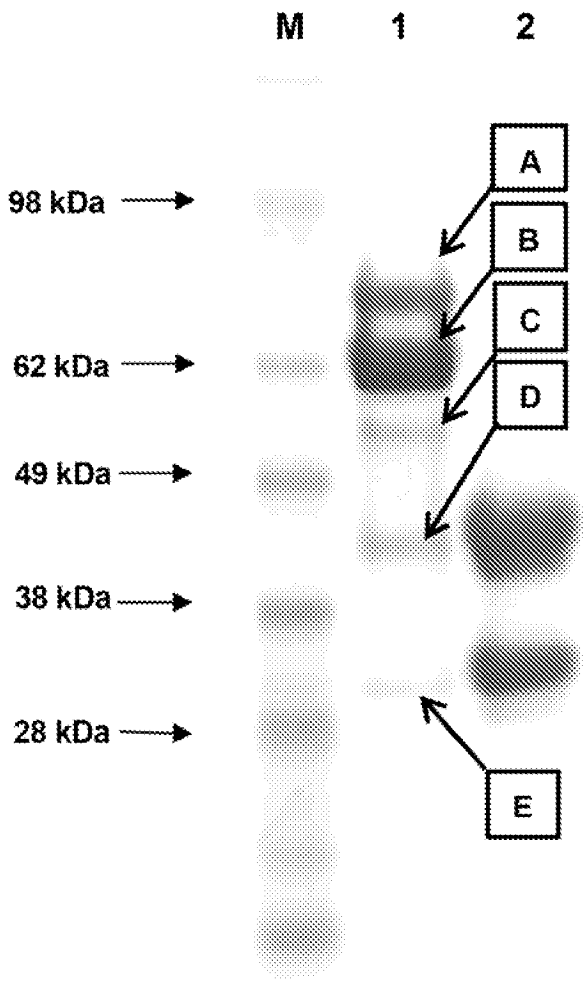
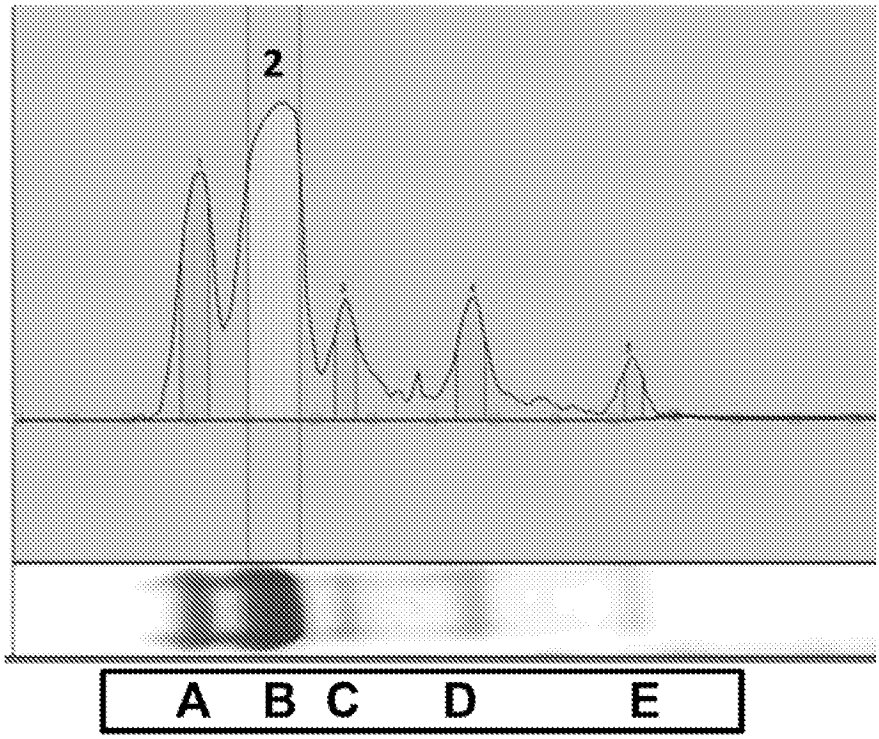


FIG. 36



LANE	PEAK	DIST	WIDTH	HEIGHT	AREA	%
1	1	54	8	150	1210	23.0
1	2	78	15	191	2850	54.2
1	3	95	6	74	436	8.3
1	4	132	8	74	560	10.7
1	5	177	5	38	197	3.7

FIG. 37

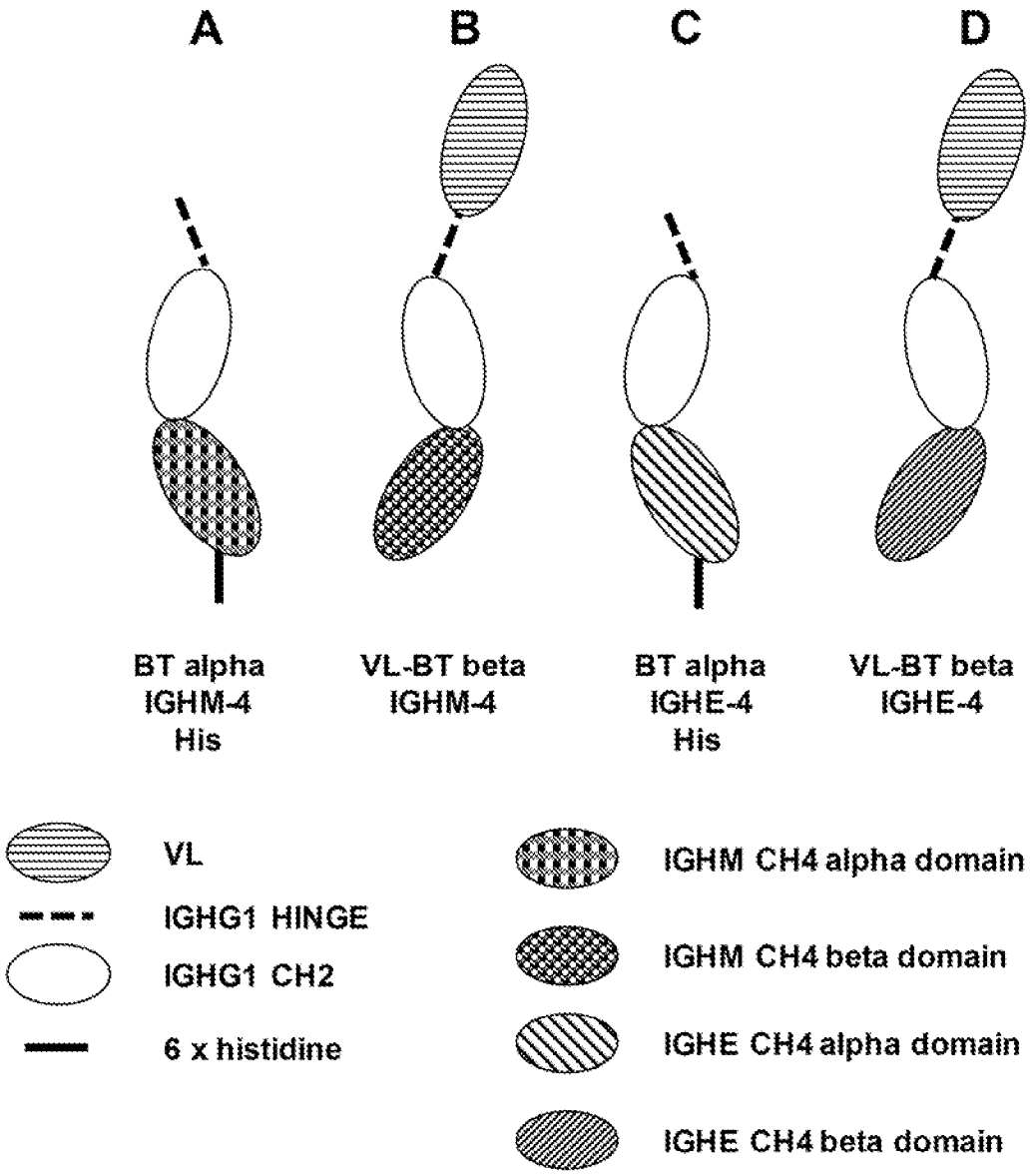

A
B
C
D
BT alpha IGHM-4 His
VL-BT beta IGHM-4
BT alpha IGHE-4 His
VL-BT beta IGHE-4
VL
IGHG1 HINGE
IGHG1 CH2
6 x histidine
IGHM CH4 alpha domain
IGHM CH4 beta domain
IGHE CH4 alpha domain
IGHE CH4 beta domain

FIG. 38

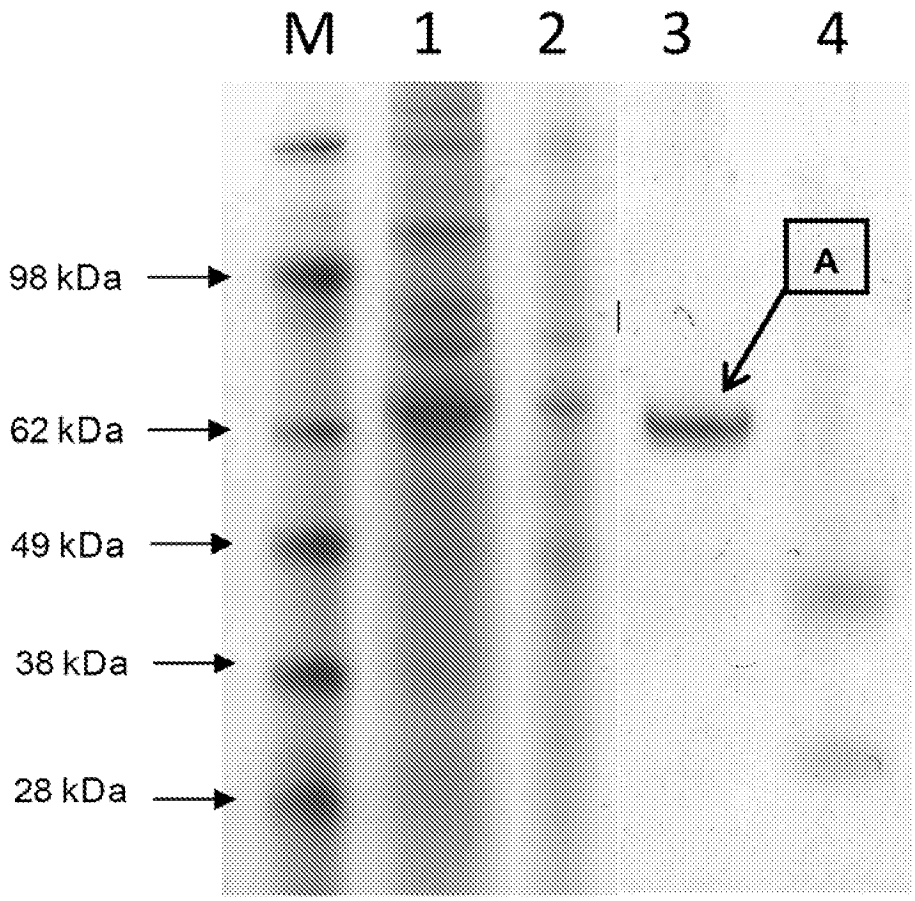


FIG. 39

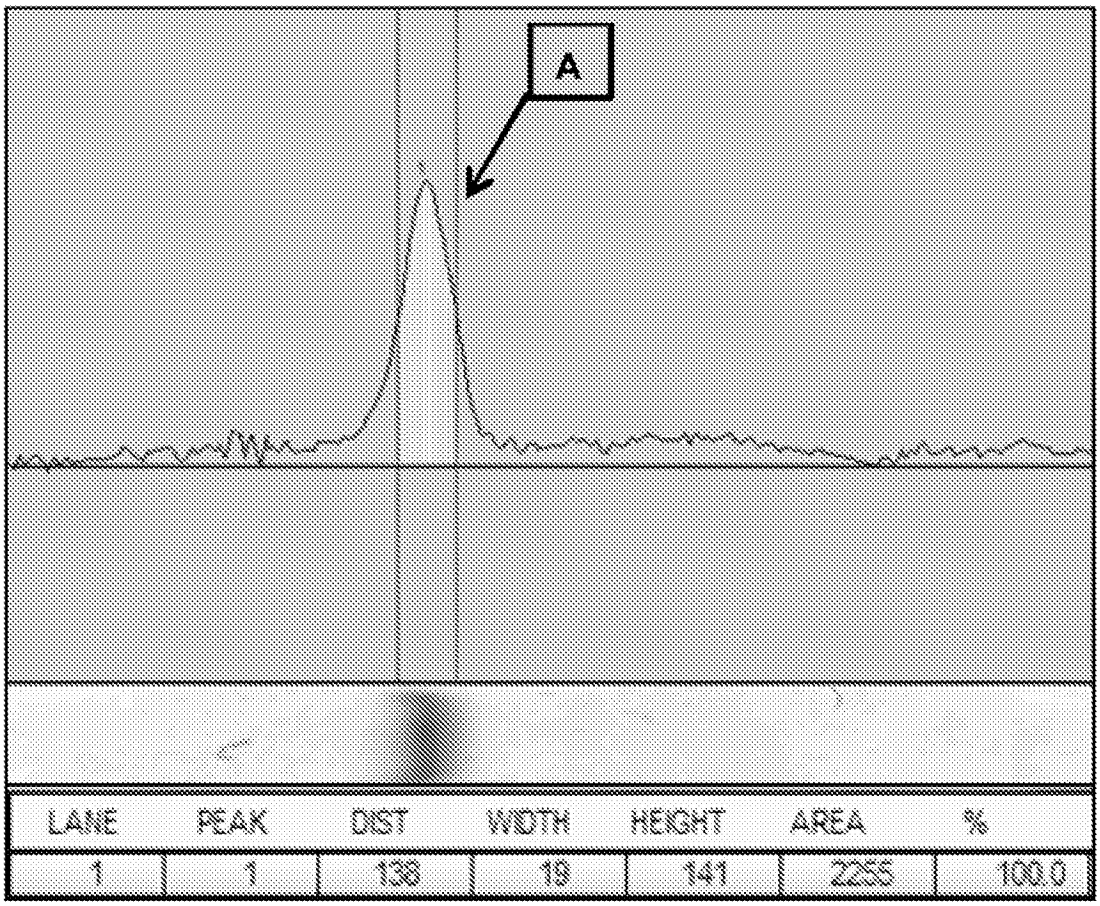


FIG. 40

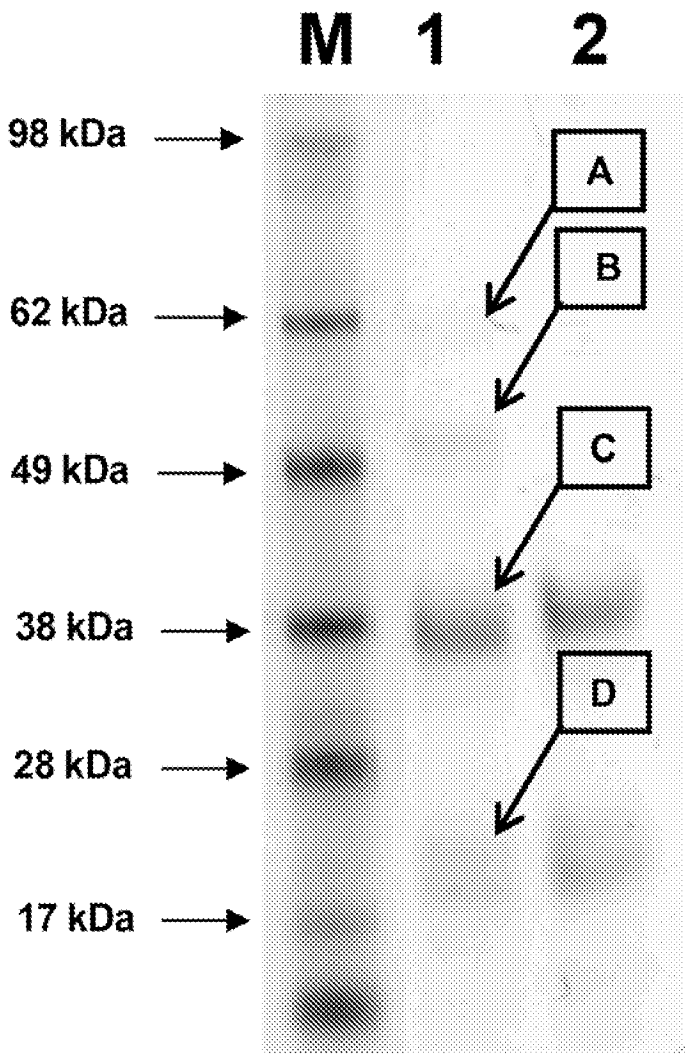
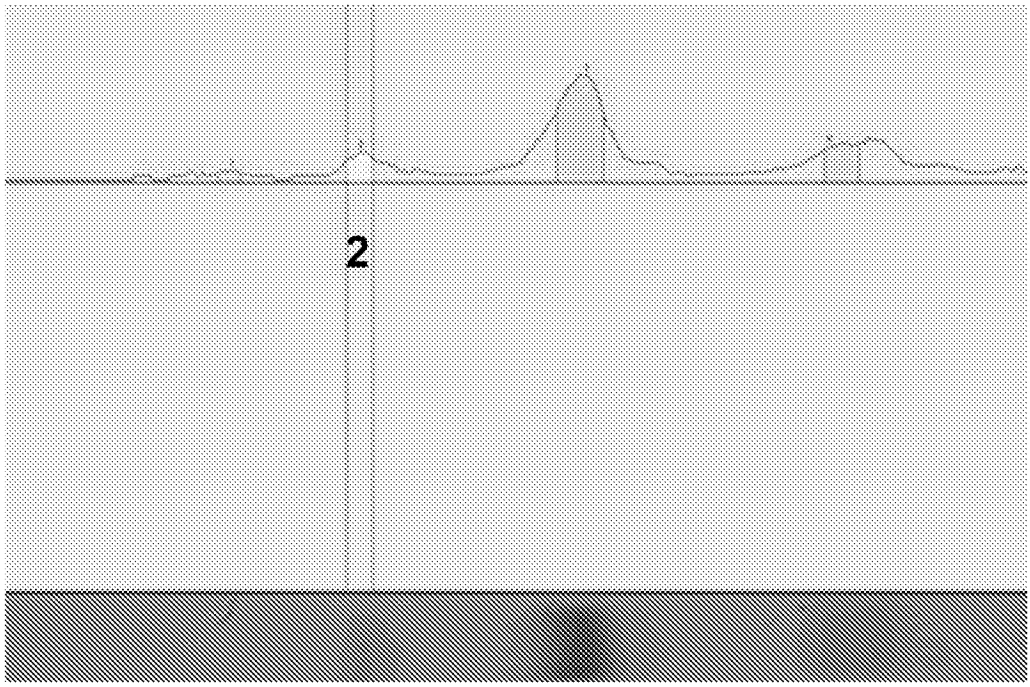


FIG. 41



LANE	PEAK	DIST	WIDTH	HEIGHT	AREA	%
1	1	108	10	7	73	3.9
1	2	167	12	18	203	10.8
1	3	271	22	62	1236	65.7
1	4	382	16	21	368	19.6

FIG. 42

```
                    A          AB      B          BC          C       CD
                  (1-15)               (16-26)    (27-38)     (39-45)
                  ---------------->    ---------->           ------>
                    1        10  15    16   23 26 27      38 39 41  45
            87654321|........|....|123|......|..|.|........| |.|...|1234567
CH1
J00220 IGHA1        ASPTSPKVFPLSLCSTQP....DGNVVIACLVQ GFFPQ.EPLS VTWSESGQGV....
J00221 IGHA2        ASPTSPKVFPLSLDSTPQ....DGNVVVACLVQ GFFPQ.EPLS VTWSESGQNV....
K02875 IGHD         APTKAPDVFPIISGCRHPKD..NSPVVLACLIT GYHP..TSVT VTWYMGTQSQ....
J00222 IGHE         ASTQSPSVFPLTRCCKNIPSN.ATSVTLGCLAT GYFP..EPVM VTCDTGSLNG....
J00228 IGHG1        ASTKGPSVFPLAPSSKSTS...GGTAALGCLVK DYFP..EPVT VSWNSGALTS....
J00230 IGHG2        ASTKGPSVFPLAPCSRSTS...ESTAALGCLVK DYFP..EPVT VSWNSGALTS....
X03604 IGHG3        ASTKGPSVFPLAPCSRSTS...GGTAALGCLVK DYFP..EPVT VSWNSGALTS....
K01316 IGHG4        ASTKGPSVFPLAPCSRSTS...ESTAALGCLVK DYFP..EPVT VSWNSGALTS....
X06766 IGHGP        ASTKGPSVFPLVPSSRSVS...EGTAALGCLVK DYFP..EPVT VSWNSGALTR....
X14940 IGHM         GSASAPTLFPLVSCENSPSD..TSSVAVGCLAQ DFLP..DSIT LSWKYKNNSDIS..
CH2
J00220 IGHA1           CCHPRLSLHRPALEDLLL..GSEANLTCTLT GLRDA.SGVT FTWTPSSGKS....
J00221 IGHA2           CCHPRLSLHRPALEDLLL..GSEANLTCTLT GLRDA.SGAT FTWTPSSGKS....
K02878 IGHD         ECPSHTQPLGVYLLTPAVQDLWL..RDKATFTCFVV GSDL..KDAH LTWEVAGKVPTG..
J00222 IGHE         VCSRDFTPPTVKILQSSCDGGGHF.PPTIQLLCLVS GYTP..GTIN ITWLEDGQVMD...
J00223 IGHEP1 .....NPRGVSAYLSRPSPFDLFI..RKSPTITCLVV DLAPSKWTVN LTWSRASGKPV...
J00228 IGHG1        APELLGGPSVFLFPPKPKDTLMI.SRTPEVTCVVV DVSHEDPEVK FNWYVDGVEVH...
J00230 IGHG2        AP.PVAGPSVFLFPPKPKDTLMI.SRTPEVTCVVV DVSHEDPEVQ FNWYVDGVEVH...
X03604 IGHG3        APELLGGPSVFLFPPKPKDTLMI.SRTPEVTCVVV DVSHEDPEVQ FKWYVDGVEVH...
K01316 IGHG4        APEFLGGPSVFLFPPKPKDTLMI.SRTPEVTCVVV DVSQEDPEVQ FNWYVDGVEVH...
X06766 IGHGP        TTEPLGGPSVFLFPPKPKDTLMI.SRTPEVTCVVV DVSHEDPEVK FNWYVDGVEVH...
X14940 IGHM            VIAELPPKVSVFVPPRDGFFGN.PRKSKLICQAT GFSP..RQIQ VSWLREGKQVGS..
```

FIG. 42 (cont.)

```
                    A          AB      B          BC          C       CD
                  (1-15)               (16-26)    (27-38)     (39-45)
                  ---------->  ------------>      ---------   ------->
                        1       10  15   16   23 26  27      38 39 41  45
                 87654321|.....|....|123|.....|..| |.....|    |.|...|1234567
CH3
J00220 IGHA1      GNTFRPEVHLLPPPSEELAL..NELVTLTCLAR GFSP..KDVL  VRWLQGSQELPRE.
J00221 IGHA2      GNTFRPEVHLLPPPSEELAL..NELVTLTCLAR GFSP..KDVL  VRWLQGSQELPRE.
K02879 IGHD         AAQAPVKLSLNLLASSDPP..EAASWLLCEVS GFSP..PNIL  LMWLEDQREVNTS.
J00222 IGHE       DSNPRGVSAYLSRPSPFDLFI..RKSPTITCLVV DLAPSKGTVN  LTWSRASGKPV...
J00223 IGHEP1      GPRAAPEVYAFATPEWLGS...RDKRTLTCLIQ NFMP..EDIS  VQWLHNEVQLPDA.
J00228 IGHG1      GQPREPQVYTLPPSRDELT...KNQVSLTCLVK GFYP..SDIA  VEWESNGQPEN...
J00230 IGHG2      GQPREPQVYTLPPSREEMT...KNQVSLTCLVK GFYP..SDIA  VEWESNGQPEN...
X03604 IGHG3      GQPREPQVYTLPPSREEMT...KNQVSLTCLVK GFYP..SDIA  VEWESSGQPEN...
K01316 IGHG4      GQPREPQVYTLPPSQEEMT...KNQVSLTCLVK GFYP..SDIA  VEWESNGQPEN...
X06766 IGHGP      GQPREPQVYTLPPSQK.MT...KNQVTLTCLVK GFYP..SDIT  VEWESNGQPEN...
X14940 IGHM       DQDTAIRVFAIPPSFASIFL..TKSTKLTCLVT DLTTY.DSVT  ISWTRQNGEAV...
CH4
J00222 IGHE        GPRAAPEVYAFATPEWPGS...RDKRTLACLIQ NFMP..EDIS  VQWLHNEVQLPDA.
X14940 IGHM       GVALHRPDVYLLPPAREQLNL..RESATITCLVT GFSP..ADVF  VQWMQRGQPLSPE.
```

FIG. 42 (cont.)

```
                                D          DE          E      EF   F          FG          G
                               (77-84)                (85-96)     (97-104)   (105-117)   (118-128)
                               ------->             ---------->  ------->               ---------->
                              77      84           85   89    96  97   104 105       117 118     128
                               |      |12345677654321|...|.....|12|.....| |...........| |.........|
CH1
J00220  IGHA1                  TARNFPPSQDASG.....DLYTTSSQLTLPATQC..LAGKSVTC HVKHY...TNPSQ DVTVPCP
J00221  IGHA2                  TARNFPPSQDASG.....DLYTTSSQLTLPATQC..PDGKSVTC HVKHY...TNPSQ DVTVPCP
K02875  IGHD                   PQRTFPEIQRRD......SYYMTSSQLSTPLQQW...RQGEYKC VVQHT...ASKSK KEIFRWP
J00222  IGHE                   TTMTLPATTLTLS.....GHYATISLLTVSGAW....AKQMFTC RVAHTPSSTDWVD NKTFS
J00228  IGHG1                  GVHTFPAVLQSS......GLYSLSSVVTVPSSSL...GTQTYIC NVNHKP..SNTKV DKKV
J00230  IGHG2                  GVHTFPAVLQSS......GLYSLSSVVTVPSSNF...GTQTYTC NVDHKP..SNTKV DKTV
X03604  IGHG3                  GVHTFPAVLQSS......GLYSLSSVVTVPSSSL...GTQTYTC NVNHKP..SNTKV DKRV
K01316  IGHG4                  GVHTFPAVLQSS......GLYSLSSVVTVPSSSL...GTKTYTC NVDHKP..SNTKV DKRV
X06766  IGHGP                  SVHTFPAVLQSS......GLYSLSSVVTVPSSSL...GTQTYTC NVDHKP..SNTKV DKTV
X14940  IGHM                   STRGFPSVLRGG.......KYAATSQVLLPSKDVMQGTDEHVVC KVQHP....NGNK EKNVPLP
CH2
J00220  IGHA1                  .AVQGPPERDLCG......CYSVSSVLPGCAEPW..NHGKTFTC TAAYPE..SKTPL TATLSKS
J00221  IGHA2                  .AVQGPPERDLCG......CYSVSSVLPGCAQPW..NHGETFTC TAAHPE..LKTPL TANITKS
K02878  IGHD                   GVEEGLLERHSN......GSQSQHSRLTLPRSLW..NAGTSVTC TLNHPS..LPPQR LMALREP
J00222  IGHE                   VDLSTASTTQEG......ELASTQSELTLSQKHW..LSDRTYTC QVTYQ....GHTF EDSTKKCA
J00223  IGHEP1                 NHSTRKEEKQRN......GTLTVTSTVPVGTRDW..IEGETYQC RVTHPQ..LPRAL VRSTTKTS
J00228  IGHG1                  NAKTKPREEQYN......STYRVVSVLTVLHQDW..LNGKEYKC KVSNKA..LPAPI EKTISKAK
J00230  IGHG2                  NAKTKPREEQFN......STFRVVSVLTVVHQDW..LNGKEYKC KVSNKG..LPAPI EKTISKTK
X03604  IGHG3                  NAKTKPREEQYN......STFRVVSVLTVLHQDW..LNGKEYKC KVSNKA..LPAPI EKTISKTK
K01316  IGHG4                  NAKTKPREEQFN......STYRVVSVLTVLHQDW..LNGKEYKC KVSNKG..LPSSI EKTISKAK
X06766  IGHGP                  NAKTKPWEEQYN......STYHVVSVLTVVHQNW..LNGREYKC KVSNKG..LPAPI EKTISKTK
X14940  IGHM                   GVTTDQVQAEAKES..GPTTYKVTSTLTIKESDW..LGQSMFTC RVDHR....GLTF QQNASSMCVP
```

SUBSTITUTE SHEET (RULE 26)

FIG. 42 (cont.)

```
                  D        DE          E      EF   F          FG          G
                (77-84)               (85-96)      (97-104)   (105-117)   (118-128)
                -------->             ----------->  -------->           ----------->
             77     84                 85   89   96  97   104 105         117 118     128
              |      |12345677654321|...|.....|12|.....| |.............| |..........|
CH3
J00220 IGHA1    KYLTWASRQEPSQG...TTTFAVTSILRVAAEDW..KKGDTFSC MVGHEA.LPLAFT QKTIDRLA
J00221 IGHA2    KYLTWASRQEPSQG...TTTFAVTSILRVAAEDW..KKGDTFSC MVGHEA.LPLAFT QKTIDRLA
K02879 IGHD     GFAPARPPPQPRS.....TTFWAWSVLRVPAPPS..PQPATYTC VVSHEDSRTLLNA SRSLEVS
J00222 IGHE     NHSTRKEEKQRN.....GTLTVTSTLPVGTRDW..IEGETYQC RVTHPH..LPRAL MRSTTKTS
J00223 IGHEP1   RHSTTQPRKTKG.....SGFFIFSRLEVTRAEW..EQKDEFIC RAVHEAAIPSQTV QRAVSVNP
J00228 IGHG1    NYKTTPPVLDSD.....GSFFLYSKLTVDKSRW..QQGNVFSC SVMHEA.LHNHYT QKSLSLSP
J00230 IGHG2    NYKTTPPMLDSD.....GSFFLYSKLTVDKSRW..QQGNVFSC SVMHEA.LHNHYT QKSLSLSP
X03604 IGHG3    NYNTTPPMLDSD.....GSFFLYSKLTVDKSRW..QQGNIFSC SVMHEA.LHNRFT QKSLSLSP
K01316 IGHG4    NYKTTPPVLDSD.....GSFFLYSRLTVDKSRW..QEGNVFSC SVMHEA.LHNHYT QKSLSLSL
X06766 IGHGP    NYKTTPPMLDSN.....GSFFLYSKLTVDKSRW..QQGNVFSC SVMHEG.LHNHYT QKSLSLSP
X14940 IGHM     KTHTNISESHPN.....ATFSAVGEASICEDDW..NSGERFTC TVTHTD..LPSPL KQTISRPK
CH4
J00222 IGHE     RHSTTQPRKTKG.....SGFFVFSRLEVTRAEW..EQKDEFIC RAVHEAASPSQTV QRAVSVNP
X14940 IGHM     KYVTSAPMPEPQA....PGRYFAHSILTVSEEEW..NTGETYTC VVAHEA.LPNRVT ERTVDKST
```

FIG. 43

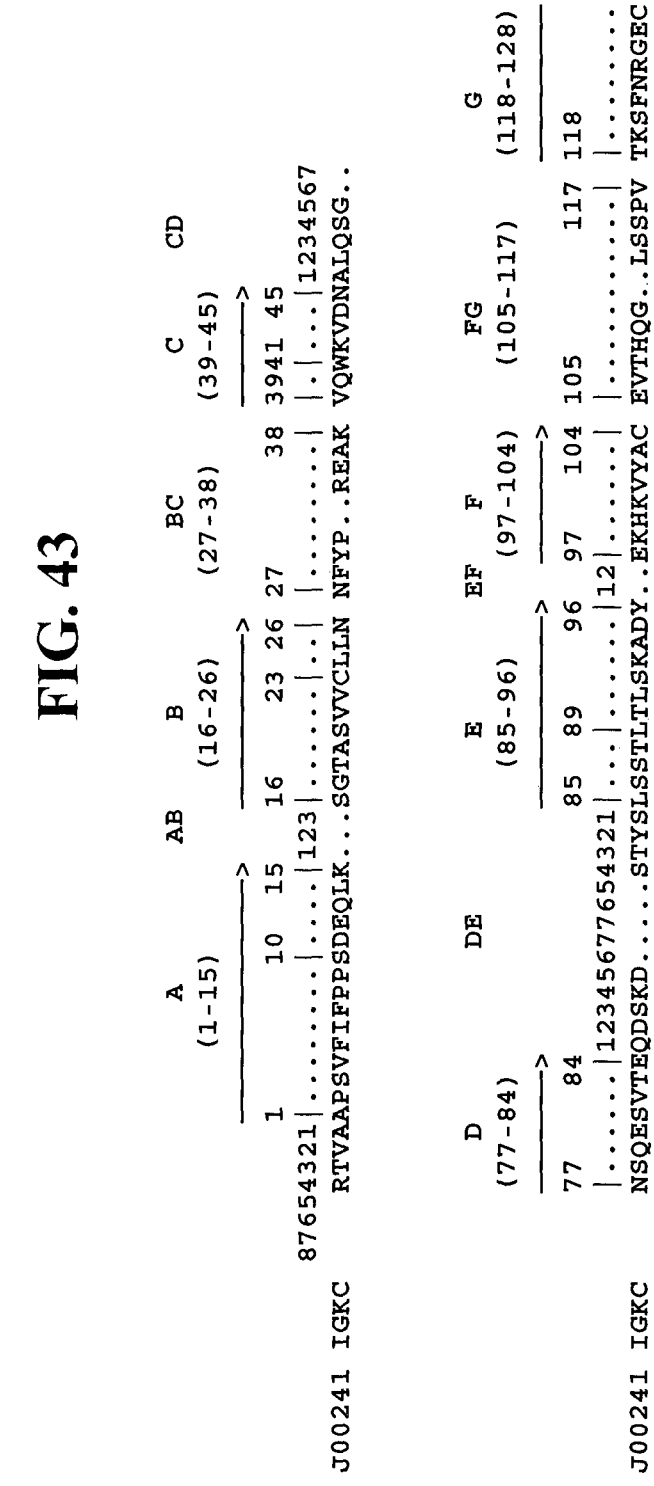

A AB B BC C CD
(1-15) (16-26) (27-38) (39-45)
1 10 15 16 23 26 27 38 3941 45
87654321 123 12 1234567
J00241 IGKC RTVAAPSVFIFPPSDEQLK...SGTASVVCLLN NFYP..REAK VQWKVDNALQSG..
D DE E EF F FG G
(77-84) (85-96) (97-104) (105-117) (118-128)
77 84 85 89 96 97 104 105 117 118
12
J00241 IGKC NSQESVTEQDSKD.....STYSLSSTLTLSKADY..EKHKVYAC EVTHQG..LSSPV TKSFNRGEC

FIG. 44

```
                                A          AB      B          BC          C      CD
                            (1-15)                (16-26)    (27-38)    (39-45)
                            ------------>       ---------->           ---------->
                                     1        10  15  16    23 26 27        38 3941 45
                            87654321|........|....|123|.....|..| |.........| |.|...|1234567
J00252 IGLC1                  PKANPTVTLFPPSSEELQ...ANKATLVCLIS DFYP..GAVT VAWKADGSPVKA..
J00253 IGLC2                GQPKAAPSVTLFPPSSEELQ...ANKATLVCLIS DFYP..GAVT VAWKADSSPVKA..
J00254 IGLC3                  PKAAPSVTLFPPSSEELQ...ANKATLVCLIS DFYP..GAVT VAWKADSSPVKA..
J03011 IGLC6                GQPKAAPSVTLFPPSSEELQ...ANKATLVCLIS DFYP..GAVK VAWKADGSPVNT..
X51755 IGLC7                GQPKAAPSVTLFPPSSEELQ...ANKATLVCLVS DFYP..GAVT VAWKADGSPVKV..

                                D          DE          E      EF  F          FG          G
                            (77-84)                  (85-96)    (97-104)   (105-117)   (118-128)
                            ------------>            ---------->           ---------- ----------
                                     77  84          85  89    96  97  104 105     117 118
                            |.......|12345677654321|...|.......|12|.....| |.............| |..........
J00252 IGLC1                GVETTKPSKQSN.......NKYAASSYLSLTPEQW..KSHRSYSC QVTHE....GSTV EKTVAPTECS
J00253 IGLC2                GVETTTPSKQSN.......NKYAASSYLSLTPEQW..KSHRSYSC QVTHE....GSTV EKTVAPTECS
J00254 IGLC3                GVETTTPSKQSN.......NKYAASSYLSLTPEQW..KSHKSYSC QVTHE....GSTV EKTVAPTECS
J03011 IGLC6                GVETTTPSKQSN.......NKYAASSYLSLTPEQW..KSHRSYSC QVTHE....GSTV EKTVAPAECS
X51755 IGLC7                GVETTKPSKQSN.......NKYAASSYLSLTPEQW..KSHRSYSC RVTHE....GSTV EKTVAPAECS
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FIG. 45

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                                A          AB     B          BC          C     CD
                              (1-15)              (16-26)    (27-38)    (39-45)
                            ---------->        ---------->            ------->
                             1        10  15    16     23 26 27      38  3941 45
                           87654321|.....|....|123|.....|..| |.....|   |.|...|1234567
X02883 TRAC                   XIQNPDPAVYQLRDSK.....SSDKSVCLFT DFDS...QTN  VSQSKDS.....
M12887 TRBC1                 EDLNKVFPPEVAVFEPSEAEISH..TQKATLVCLAT GFFP..DHVE  LSWWVNGKEVHS..
M12888 TRBC2                 EDLKNVFPPEVAVFEPSEAEISH..TQKATLVCLAT GFYP..DHVE  LSWWVNGKEVHS..
M22148 TRDC                   XSQPHTKPSVFVMKNG.....TNVACLVK EFYP..KDIR  INLVSSKKI.....
M14996 TRGC1               DKQLDADVSPKPTIFLPSIAETKL..QKAGTYLCLLE KFFP..DVIK  IHWQEKKSNTIL..
M15002 TRGC2(2x)           DKQLDADVSPKPTIFLPSIAETKL..QKAGTYLCLLE KFFP..DIIK  IHWQEKKSNTIL..
M17323 TRGC2(3x)           DKQLDADVSPKPTIFLPSIAETKL..QKAGTYLCLLE KFFP..DIIK  IHWQEKKSNTIL..

                              D          DE          E      EF   F              FG                 G
                            (77-84)                  (85-96)     (97-104)       (105-117)          (118-128)
                            ------->                 ---------->  ------->                         ----------
                            77    84                 85  89    96 97   104 105                 117 118
                            |......|12345677654321|...|......|12|......| |......123456654321.....| |........
X02883 TRAC                DVYITDKTVLDMRSM.DFKSNSAVAWSNKS.....DFAC ANAFNN.............SIIPE DTFFPSP
M12887 TRBC1               GVSTDPQPLKEQPAL.NDSRYCLSSRLRVSATFWQ.NPRNHFRC QVQFYGLSENDEWTQDRAKPVTQIV SAEAWGRA
M12888 TRBC2               GVSTDPQPLKEQPAL.NDSRYCLSSRLRVSATFWQ.NPRNHFRC QVQFYGLSENDEWTQDRAKPVTQIV SAEAWGRA
M22148 TRDC                TEFDPAIVISPS......GKYNAVKLGKYED.....SNSVTC SVQHDNK.............TVHST DFEVKTDST
M14996 TRGC1               GSQEGNTMKTND.....TYMKFSWLTVPEK....SLDKEHRC IVRHENN............KNGVDQ EIIFPPIKT
M15002 TRGC2(2x)           GSQEGNTMKTND.....TYMKFSWLTVPEE....SLDKEHRC IVRHENN............KNGIDQ EIIFPPIKT
M17323 TRGC2(3x)           GSQEGNTMKTND.....TYMKFSWLTVPEE....SLDKEHRC IVRHENN............KNGIDQ EIIFPPIKT
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