

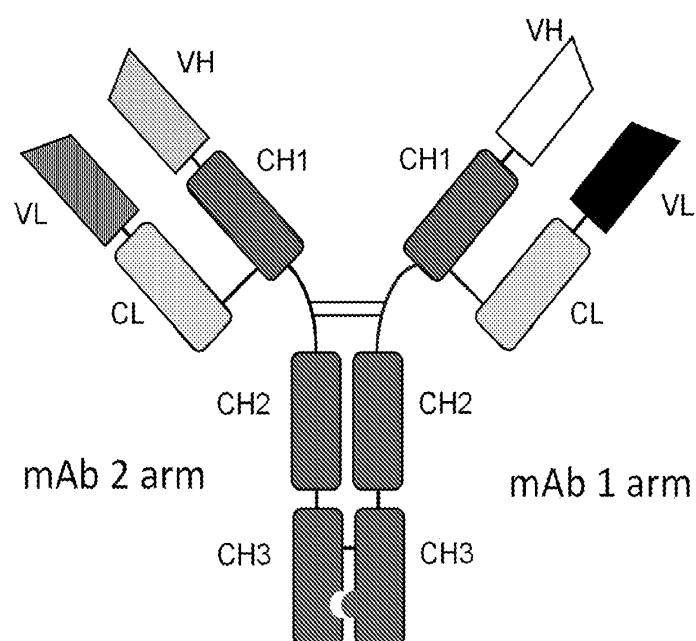


FIG. 1

Region		CHI																							
EU number		118	119	120	121	122	123	124	125	126	127	128	129	130	131	132	133	134	135	136	137	138	139	140	141
IgG1	A	S	T	K	G	P	S	V	F	P	L	A	P	S	S	S	K	S	T	S	G	T	A	A	A
IgG2	A	S	T	K	G	P	S	V	F	P	L	A	P	C	S	R	S	T	S	E	S	T	A	A	A
IgG3	A	S	T	K	G	P	S	V	F	P	L	A	P	C	S	R	S	T	S	G	T	A	A	A	A
IgG4	A	S	T	K	G	P	S	V	F	P	L	A	P	C	S	R	S	T	S	E	S	T	A	A	A

Region		CHI																							
EU number		142	143	144	145	146	147	148	149	150	151	152	153	154	155	156	157	158	159	160	161	162	163	164	165
IgG1	L	G	C	L	V	K	D	Y	F	P	E	P	V	T	V	S	W	N	S	G	A	L	T	S	S
IgG2	L	G	C	L	V	K	D	Y	F	P	E	P	V	T	V	S	W	N	S	G	A	L	T	S	S
IgG3	L	G	C	L	V	K	D	Y	F	P	E	P	V	T	V	S	W	N	S	G	A	L	T	S	S
IgG4	L	G	C	L	V	K	D	Y	F	P	E	P	V	T	V	S	W	N	S	G	A	L	T	S	S

Region		CHI																							
EU number		166	167	168	169	170	171	172	173	174	175	176	177	178	179	180	181	182	183	184	185	186	187	188	189
IgG1	G	V	H	T	T	F	P	A	V	L	Q	S	S	G	L	Y	S	L	S	S	V	V	T	V	P
IgG2	G	V	H	T	T	F	P	A	V	L	Q	S	S	G	L	Y	S	L	S	S	V	V	T	V	P*
IgG3	G	V	H	T	T	F	P	A	V	L	Q	S*	S	G	L	Y	S	L	S	S	V	V	T	V	P
IgG4	G	V	H	T	T	F	P	A	V	L	Q	S	S	G	L	Y	S	L	S	S	V	V	T	V	P

Region		CHI																							
EU number		190	191	192	193	194	195	196	197	198	199	200	201	202	203	204	205	206	207	208	209	210	211	212	213
IgG1	S	S	S	S	L	G	T	Q	T	Y	I	C	N	V	N	H	K	P	S	N	T	K	V	D	K
IgG2	S	S	S	N*	F*	G	T	Q	T	Y	T	C	N	V	D	H	K	P	S	N	T	K	V	D	K
IgG3	S	S	S	S*	L*	G	T	Q	T	Y	T	C	N	V	N	H	K	P	S	N	T	K	V	D	K
IgG4	S	S	S	S	L	G	T	K	T	Y	T	C	N	V	D	H	K	P	S	N	T	K	V	D	K

Region		CHI									
EU number		214	215	216	217	218		219	220		
IgG1	K*	V	E	P	K			S	C		
IgG2	T	V	E	R	K			C	C		
IgG3	R	V	E	L	K	T	P	L	G		
IgG4	R	V	E	S	K	Y	G				

FIG. 2A

Region		CL																							
EU number	108	109	110	111	112	113	114	115	116	117	118	119	120	121	122	123	124	125	126	127	128	129	130	131	
Kappa	R	T	V	A	A	P	S	V	F	I	F	P	P	S	D	E	Q	L	K	S	G	T	A	S	
Lambda	Q	P	K	A	A*	P	S*	V	T	L	F	P	P	S	S	E	E	L	Q	A	N	K	A	T	

Region		CL																							
EU number	132	133	134	135	136	137	138	139	140	141	142	143	144	145	146	147	148	149	150	151	152	153		154	
Kappa	V	V	C	L	L	N	N	F	Y	P	R	E	A	K	V	Q	W	K	V	D	N	A*		L	
Lambda	L	V	C	L	I	S	D	F	Y	P	G	A	V	T	V	A	W	K	A	D	S*	S	P	V	

Region		CL																							
EU number	155	156	157	158	159	160	161	162	163	164	165	166	167	168	169	170	171	172	173	174	175	176	177	178	
Kappa	Q	S	G	N	S	Q	E	S	V	T	E	Q	D	S	K	D	S	T	Y	S	L	S	S	T	
Lambda	K	A	G			V	E	T*	T	T	P	S	K	Q	S	N	N	K	Y	A	A	S	S	Y	

Region		CL																							
EU number	179	180	181	182	183	184	185	186	187	188	189	190	191	192	193	194	195	196	197	198	199	200	201	202	
Kappa	L	T	L	S	K	A	D	Y	E	K	H	K	V*	Y	A	C	E	V	T	H	Q	G	L	S	
Lambda	L	S	L	T	P	E	Q	W	K	S	H	R*	S	Y	S	C	Q	V	T	H	E	G			

Region		CL											
EU number	203	204	205	206	207	208	209	210	211	212	213	214	
Kappa	S	P	V	T	K	S	F	N	R	G	E	C	
Lambda	S	T	V	E	K	T	V	A	P	T	E	C	

FIG. 2B

Region		FR1																							
Kabat number	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	
Anti-CD47 VH	Q	V	Q	L	V	Q	S	G	A	E	V	K	K	P	G	A	S	V	K	V	S	C	K	A	
Anti-CLDN18.2 VH	E	V	Q	L	V	E	S	G	G	G	L	V	Q	P	G	G	S	L	R	L	S	C	A	A	

Region		CDR1												FR2											
Kabat number	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	
Anti-CD47 VH	S	G	Y	T	F	T	S	Y	W	M	H	W	V	R	Q	A	P	G	Q	G	L	E	W	I	
Anti-CLDN18.2 VH	S	G	F	I	F	S	S	F	G	M	H	W	V	R	Q	A	P	G	K	G	L	E	W	V	

Region		CDR2												FR3											
Kabat number	49	50	51	52	52a	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	
Anti-CD47 VH	G	N	I	D	P	S	D	S	E	T	H	Y	A	Q	K	F	Q	G	R	V	T	L	T	V	
Anti-CLDN18.2 VH	A	Y	I	S	S	G	R	S	T	M	Y	Y	A	D	S	V	K	G	R	F	T	I	S	R	

Region		CDR3												FR4											
Kabat number	72	73	74	75	76	77	78	79	80	81	82	82a	82b	82c	83	84	85	86	87	88	89	90	91	92	
Anti-CD47 VH	D	K	S	T	S	T	V	Y	M	E	L	S	S	L	R	S	E	D	T	A	V	Y	Y	C	
Anti-CLDN18.2 VH	D	N	S	K	N	T	L	Y	L	Q	M	N	S	L	T	A	E	D	T	A	V	Y	Y	C	

Region		CDR3												FR4											
Kabat number	93	94	95	96	97	98	99	100	100a	100b	101	102	103	104	105	106	107	108	109	110	111	112	113		
Anti-CD47 VH	A	G	T	D	L						A	Y	W	G	Q	G	T	L	V	T	V	S	S		
Anti-CLDN18.2 VH	A	R	G	G	F	Y	G	N	S	L	D	Y	W	G	Q	G	T	L	V	T	V	S	S		

FIG. 2C

Region		FR1																							
Kabat number		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24
Anti-CD47 VL	E	I	V	L	T	Q	S	P	G	T	L	S	L	S	P	G	E	R	A	T	L	S	C	H	
Anti-CLDN18.2 VL	D	I	Q	M	T	Q	S	P	S	S	L	S	A	S	V	G	D	R	V	T	I	T	C	K	

Region		CDR1												FR2											
Kabat number		25	26	27	27A	27B	27C	27D	27E	27F	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42
Anti-CD47 VL	A	S	Q								N	I	N	V	W	L	S	W	Y	Q	Q	K	P	G	Q
Anti-CLDN18.2 VL	S	S	L	S	L	L	L	N	S	G	N	Q	K	N	Y	L	T	W	Y	Q	Q	K	P	G	K

Region		FR2												FR3											
Kabat number		43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66
Anti-CD47 VL	A	P	R	L	L	L	I	Y	K	A	S	N	L	H	T	G	I	P	D	R	F	S	G	S	G
Anti-CLDN18.2 VL	A	P	K	L	L	L	I	Y	W	A	S	T	R	E	S	G	V	P	S	R	F	S	G	S	G

Region		FR3																							
Kabat number		67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90
Anti-CD47 VL	S	G	T	D	D	F	T	L	T	I	S	R	L	E	P	E	D	F	A	V	Y	Y	C	Q	Q
Anti-CLDN18.2 VL	S	G	T	D	D	F	T	L	T	I	S	S	L	Q	P	E	D	F	A	T	Y	S	C	Q	N

Region		CDR3												FR4											
Kabat number		91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107							
Anti-CD47 VL	G	Q	S	Y	P	P	F	T	F	G	Q	G	T	K	V	E	I	K							
Anti-CLDN18.2 VL	A	Y	S	Y	P	P	L	T	F	G	Q	G	T	K	V	E	I	K							

FIG. 2D

Region		FR1																							
Kabat number	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	
Anti-CD3 VH	D	I	K	L	Q	Q	S	G	A	E	L	A	R	P	G	A	S	V	K	M	S	C	K	T	
Anti-DLL3 VH	E	V	R	L	S	Q	S	G	G	Q	M	K	K	P	G	E	S	M	R	L	S	C	R	A	

Region		CDR1												FR2											
Kabat number	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	
Anti-CD3 VH	S	G	Y	T	F	T	R	Y	T	M	H	W	V	K	Q	R	P	G	Q	G	L	E	W	I	
Anti-DLL3 VH	S	G	Y	T	F	T	S	Y	V	M	H	W	V	R	Q	A	P	G	R	R	P	E	W	I	

Region		CDR2												FR3											
Kabat number	49	50	51	52	52a	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	
Anti-CD3 VH	G	Y	I	N	P	S	R	G	Y	T	N	Y	N	Q	K	F	K	D	K	A	T	L	T	T	
Anti-DLL3 VH	G	Y	I	N	P	Y	N	D	A	T	K	Y	A	R	K	F	Q	G	R	A	T	L	T	S	

Region		FR3																							
Kabat number	72	73	74	75	76	77	78	79	80	81	82	82a	82b	82c	83	84	85	86	87	88	89	90	91	92	
Anti-CD3 VH	D	K	S	S	S	T	A	Y	M	Q	L	S	S	L	T	S	E	D	S	A	V	Y	Y	C	
Anti-DLL3 VH	D	K	Y	S	D	T	A	F	L	E	L	R	S	L	T	S	D	D	T	A	V	Y	Y	C	

Region		CDR3												FR4											
Kabat number	93	94	95	96	97	98	99	100	100a	100b	101	102	103	104	105	106	107	108	109	110	111	112	113		
Anti-CD3 VH	A	R	Y	Y	D	D	H	Y	S	L	D	Y	W	G	Q	G	T	T	L	T	V	S	S		
Anti-DLL3 VH	A	R	G	G	Y	D	Y	D	G		D	Y	W	G	R	G	A	P	V	T	V	S	S		

FIG. 2E

Region		FR1																							
Kabat number		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24
Anti-CD3 VL	D	I	Q	L	T	Q	Q	S	P	A	I	M	S	A	S	P	G	E	K	V	T	M	T	C	R
Anti-DLL3 VL	E	I	V	L	T	Q	Q	S	P	G	T	L	S	L	S	P	G	E	R	A	T	L	S	C	H

Region		CDR1												FR2											
Kabat number		25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48
Anti-CD3 VL	A	S	S	S	S	S	V	S	Y	M	N	W	Y	Q	Q	K	S	G	T	S	P	K	R	W	I
Anti-DLL3 VL	A	S	Q	N	I	N	I	N	V	W	L	S	W	Y	Q	K	P	G	Q	A	P	R	L	L	I

Region		CDR2												FR3											
Kabat number		49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72
Anti-CD3 VL	Y	D	T	S	K	K	V	A	S	G	V	P	Y	R	F	S	G	S	G	S	G	T	S	Y	S
Anti-DLL3 VL	Y	K	A	S	S	N	L	H	T	G	I	P	D	R	F	S	G	S	G	S	G	T	D	F	T

Region		FR3												CDR3											
Kabat number		73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96
Anti-CD3 VL	L	T	I	I	S	S	M	E	A	E	D	A	A	T	Y	Y	C	Q	Q	W	S	S	N	P	L
Anti-DLL3 VL	L	T	I	I	S	R	L	E	P	E	D	F	A	V	Y	Y	C	Q	Q	G	Q	S	Y	P	F

Region		FR4											
Kabat number		97	98	99	100	101	102	103	104	105	106	107	
Anti-CD3 VL	T	F	G	A	G	T	K	L	E	L	K		
Anti-DLL3 VL	T	F	G	Q	G	G	T	K	V	E	I	K	

FIG. 2F

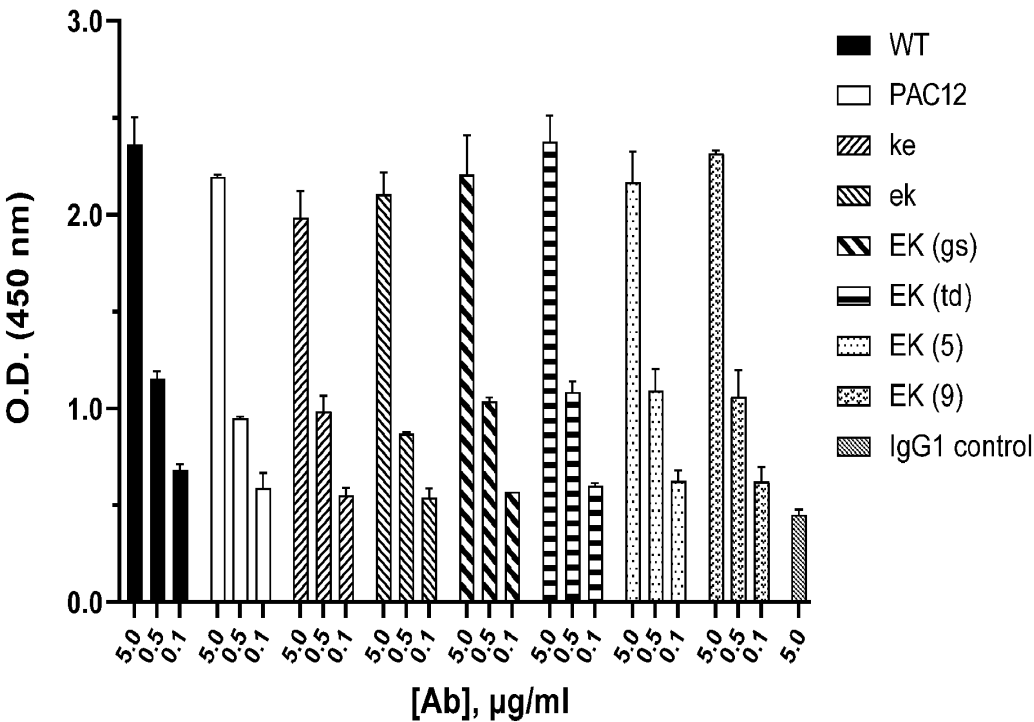


FIG. 3A

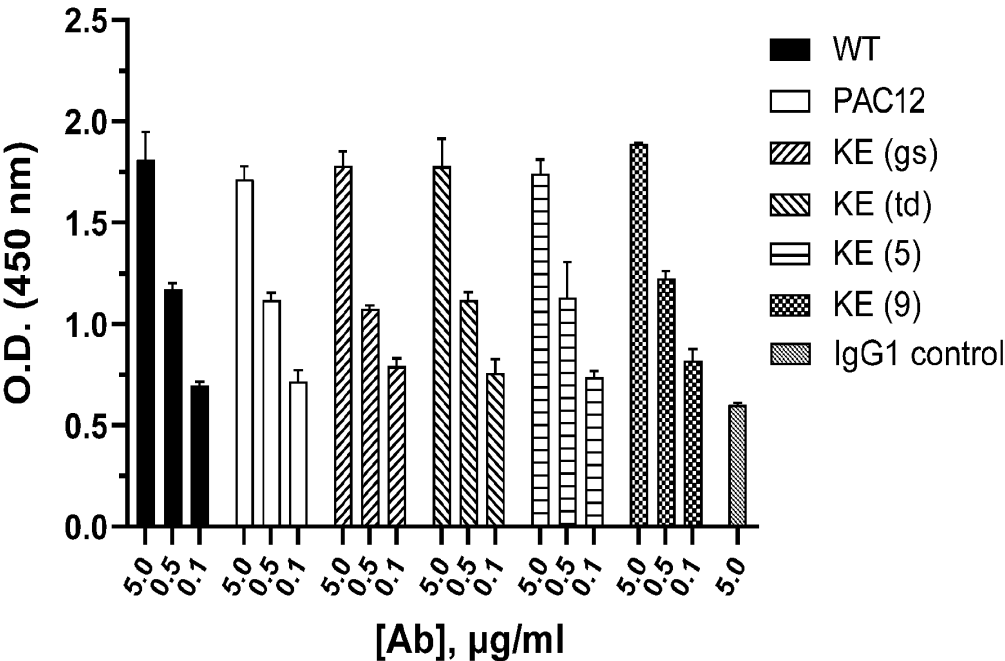


FIG. 3B

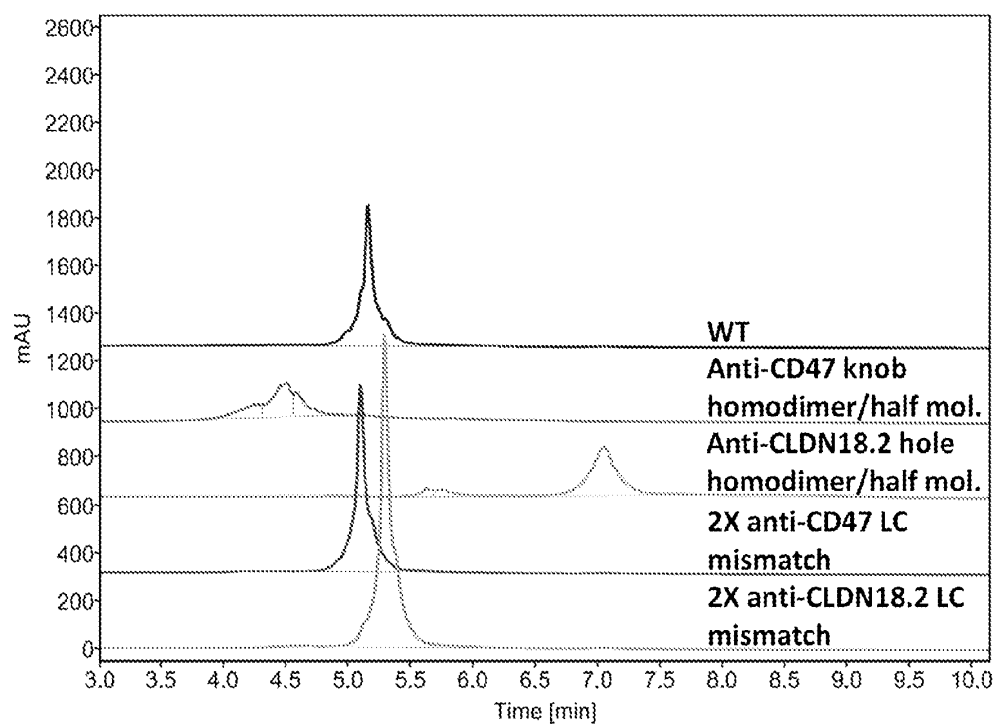


FIG. 3C

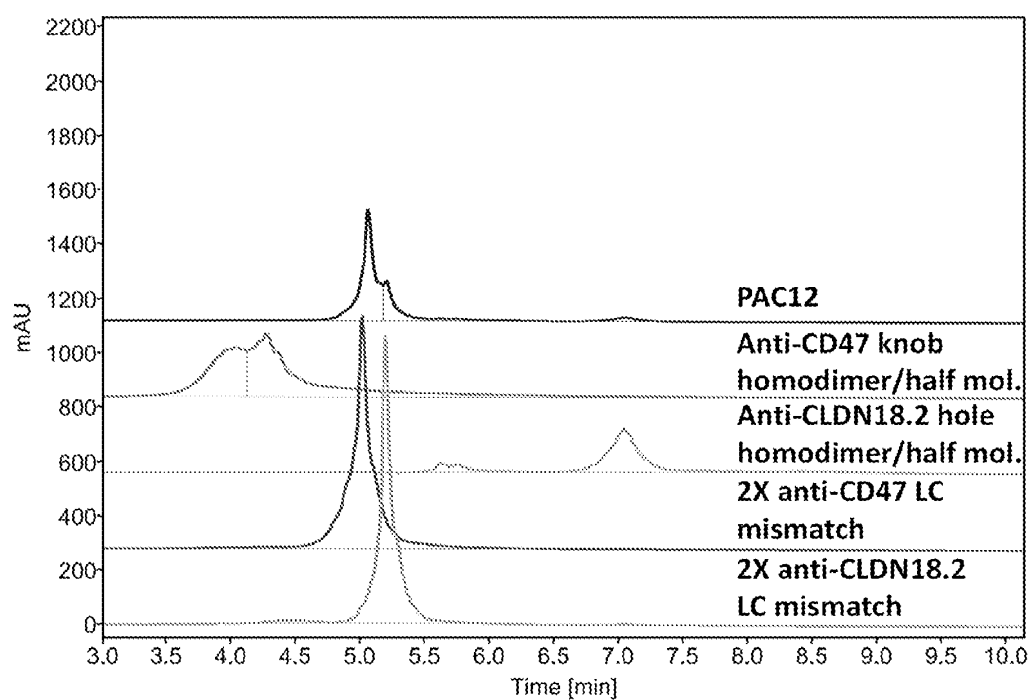
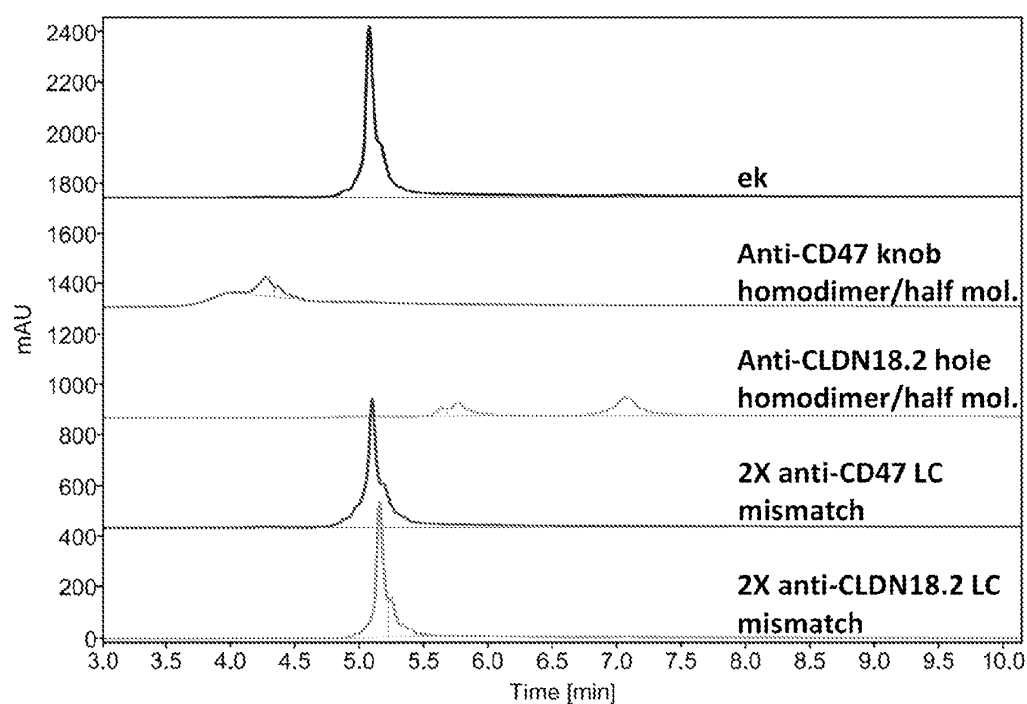
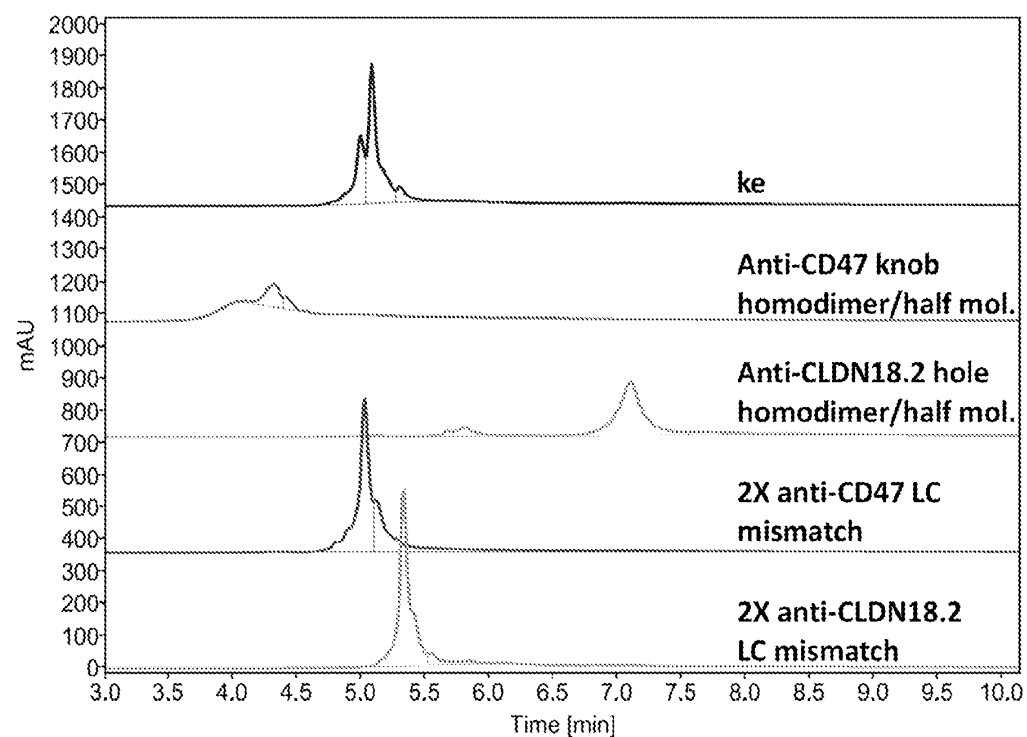


FIG. 3D

**FIG. 3E****FIG. 3F**

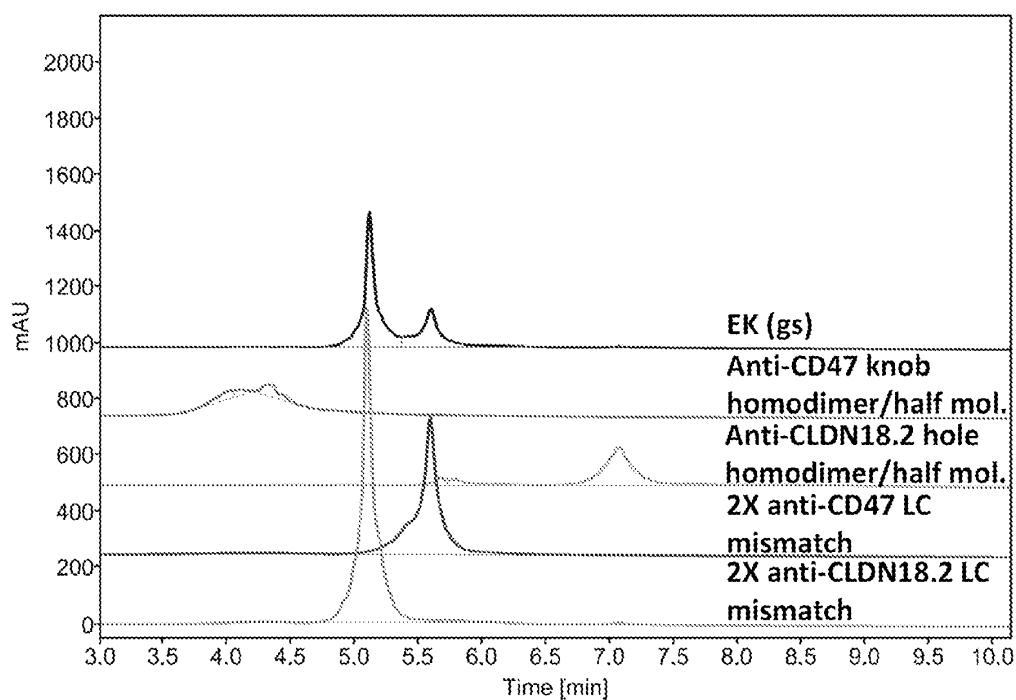


FIG. 3G

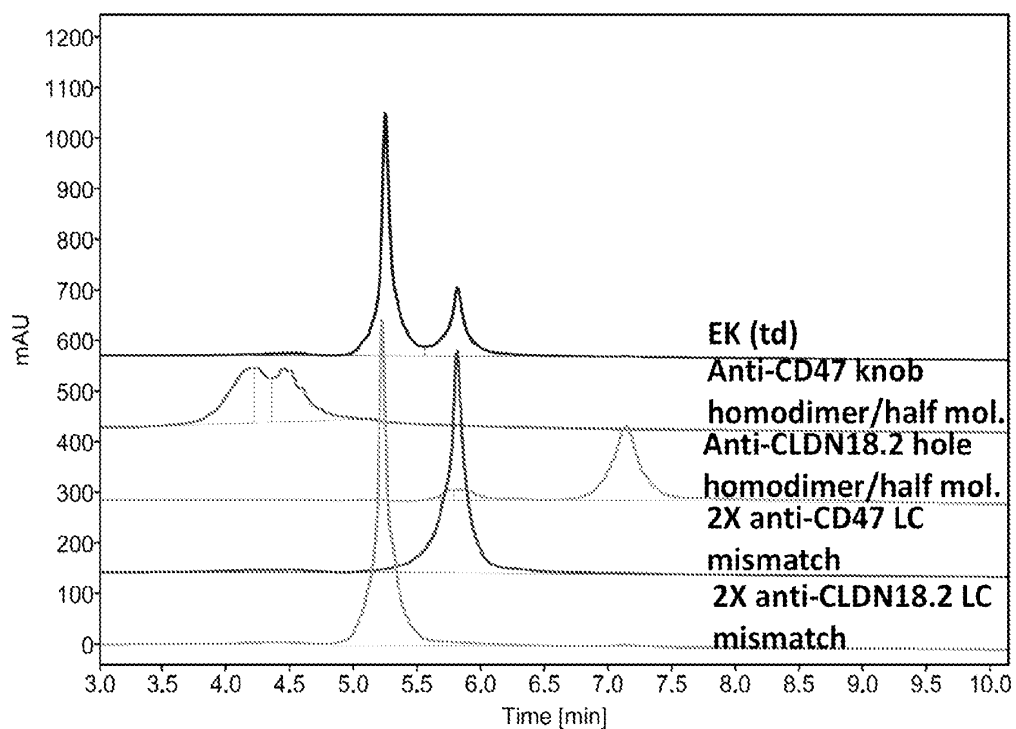
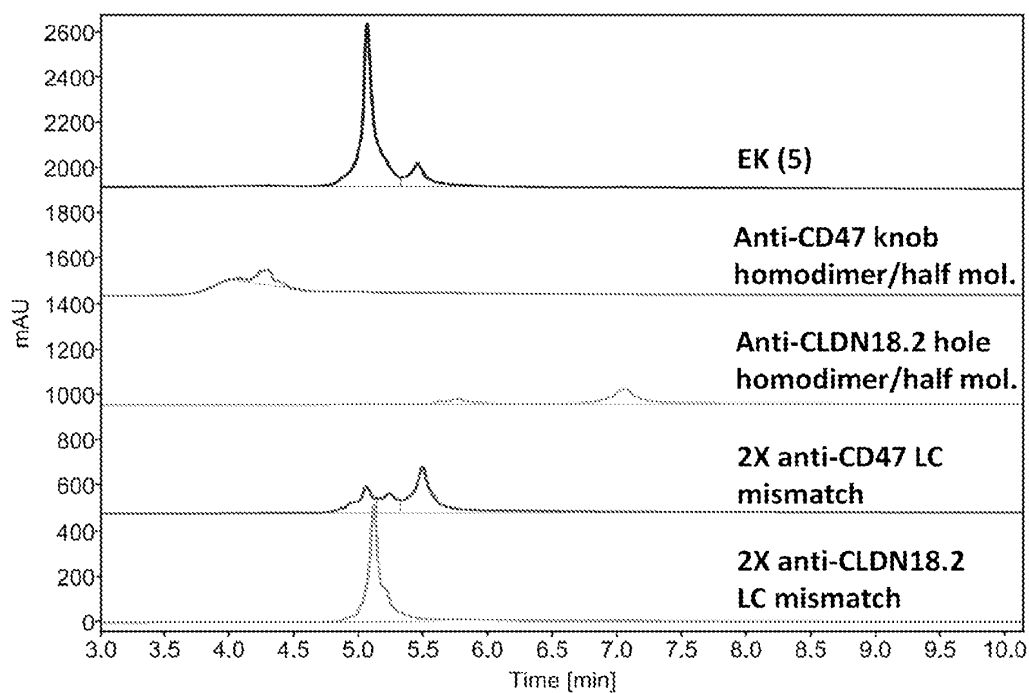
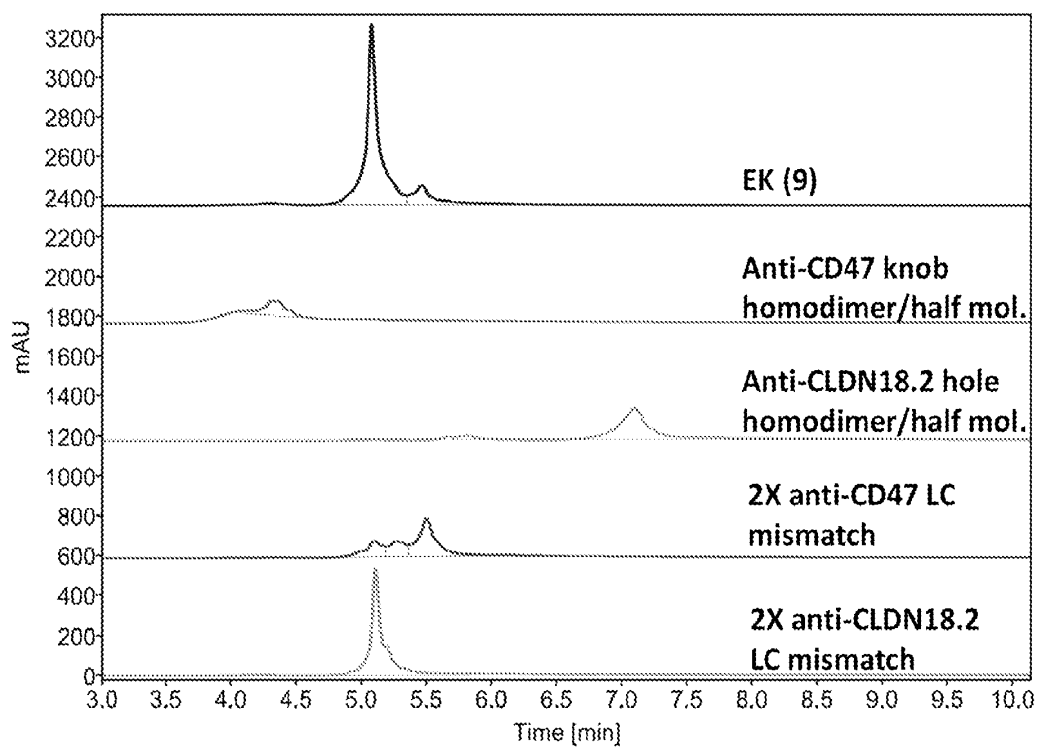


FIG. 3H

**FIG. 3I****FIG. 3J**

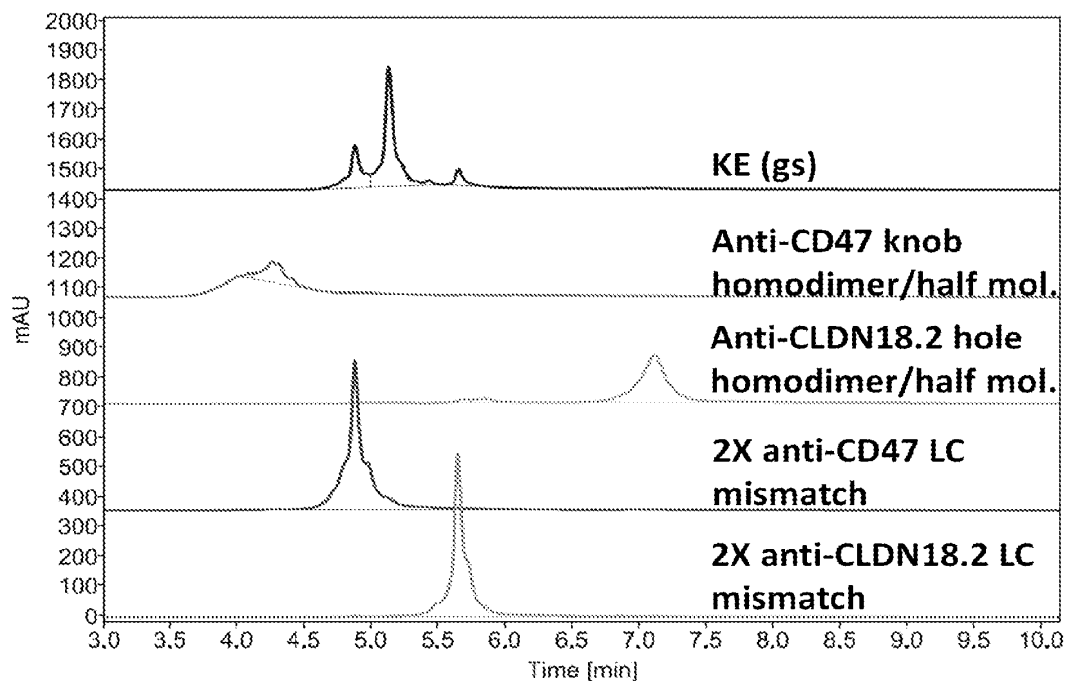


FIG. 3K

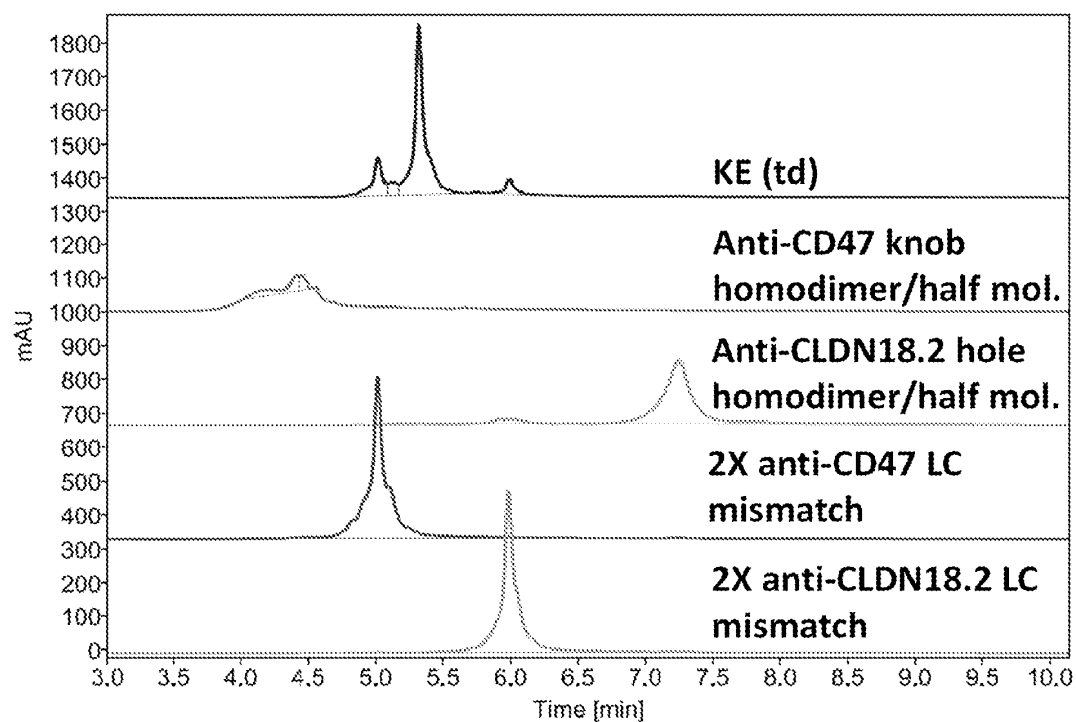


FIG. 3L

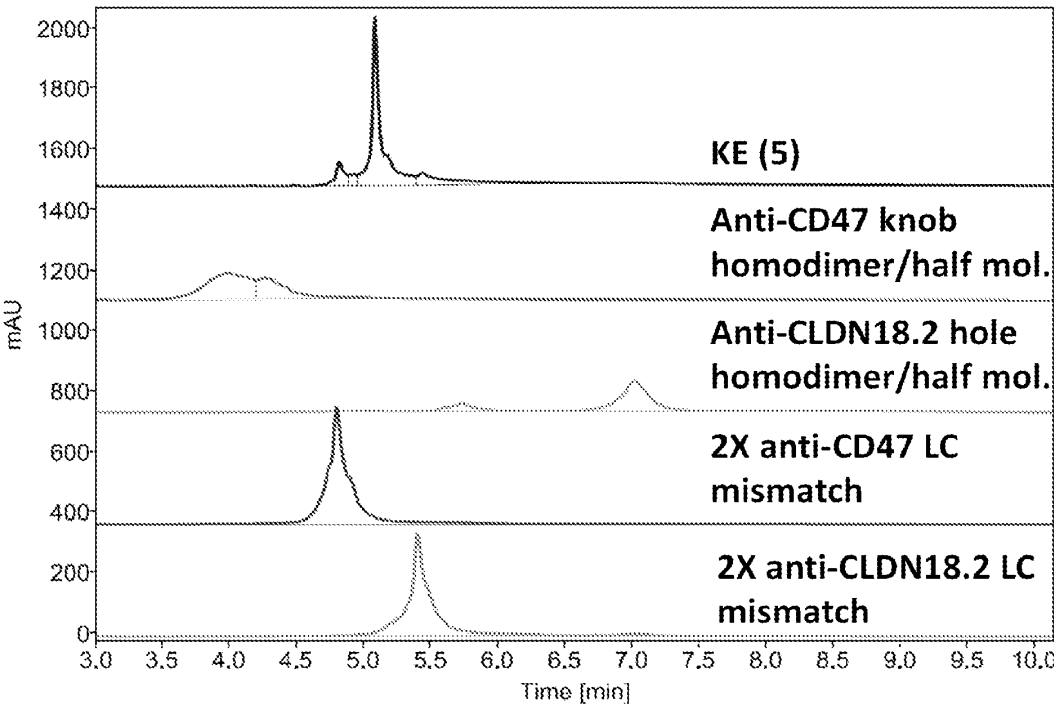


FIG. 3M

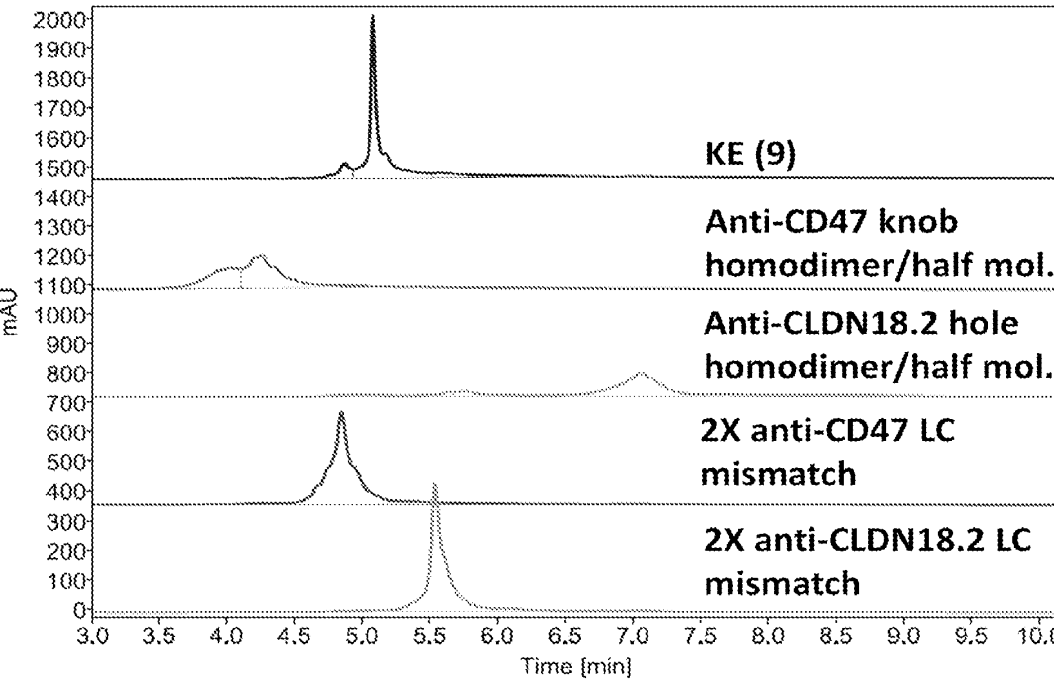
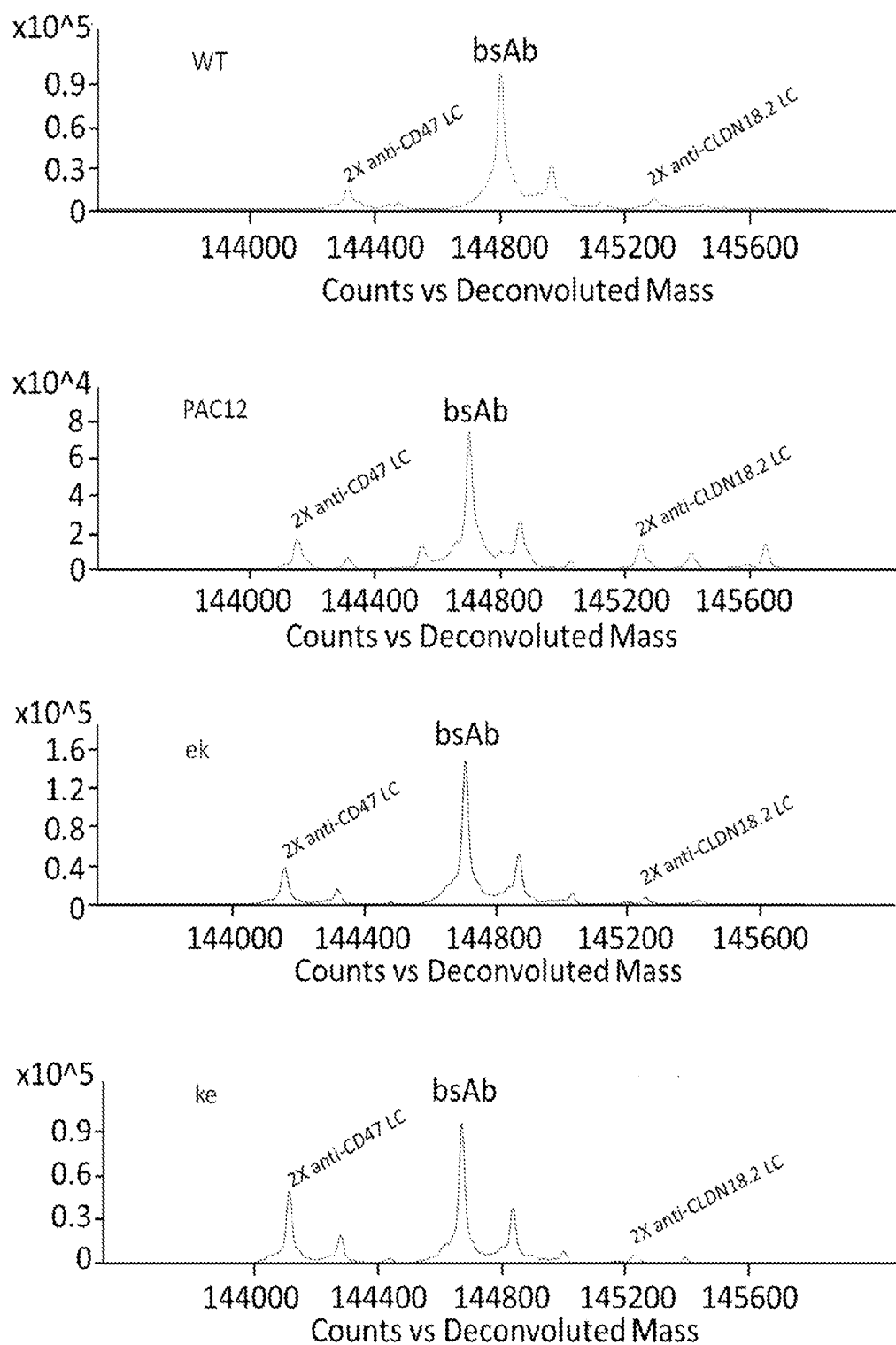
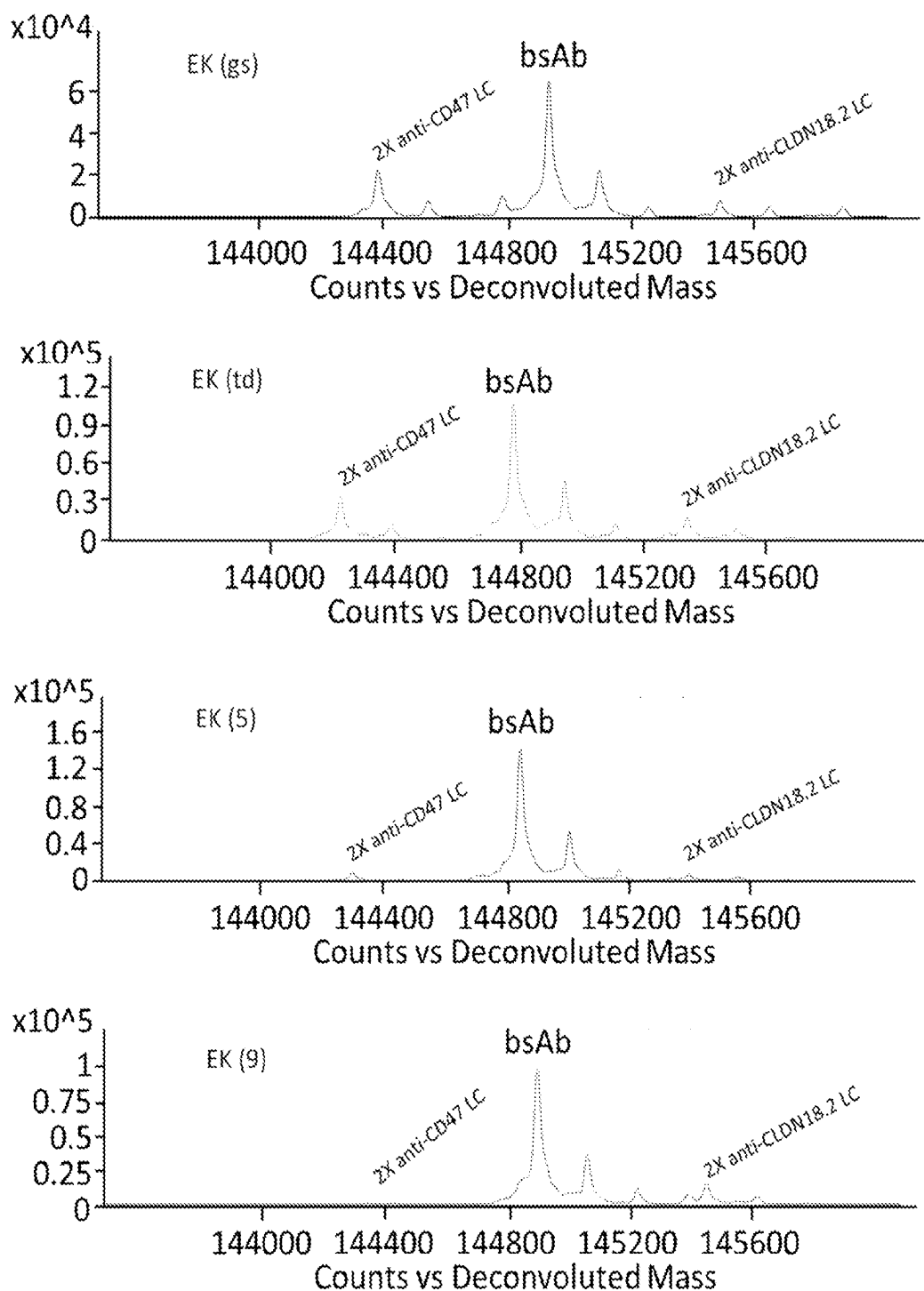


FIG. 3N

**FIG. 30**

**FIG. 3P**

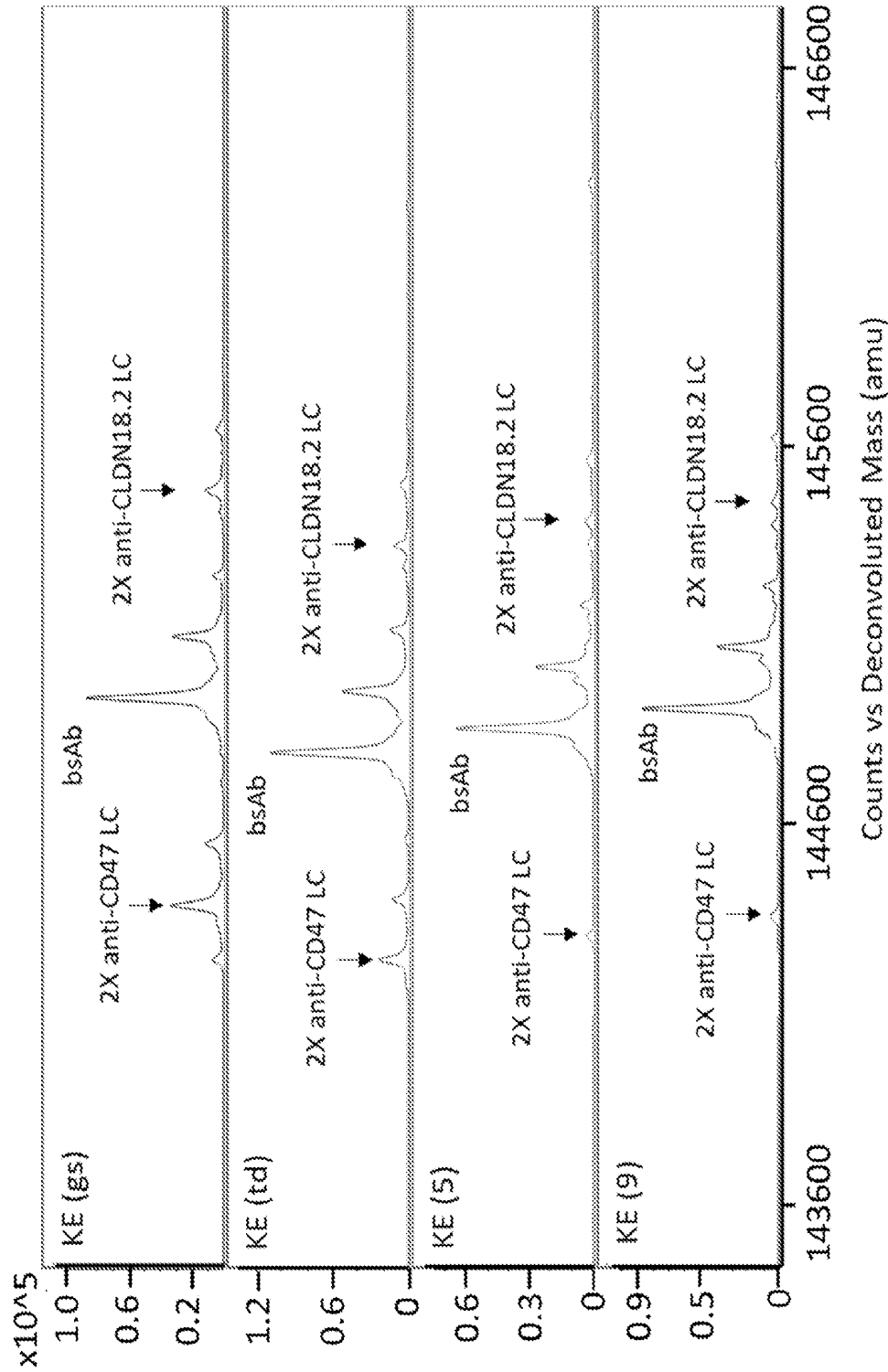


FIG. 3Q

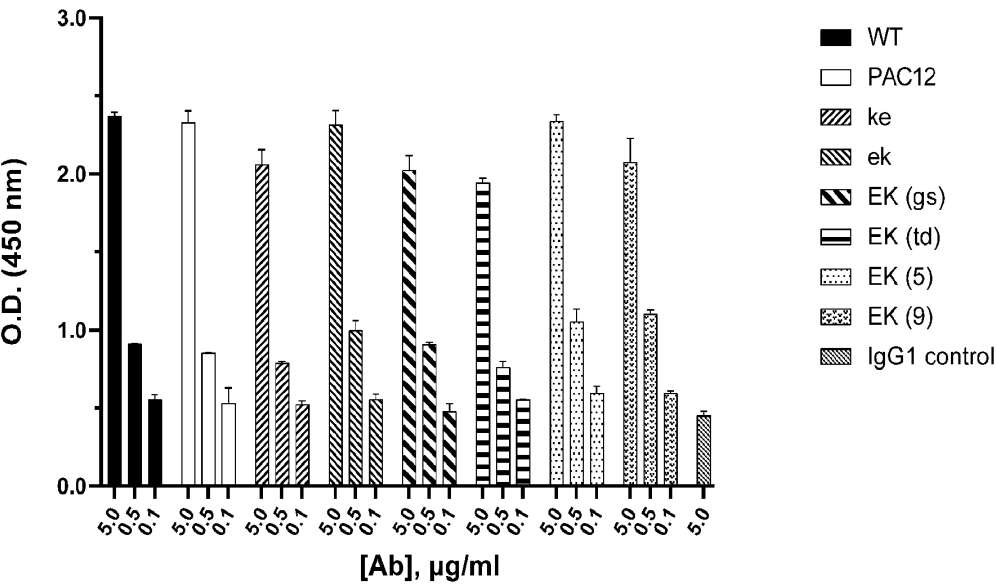


FIG. 4A

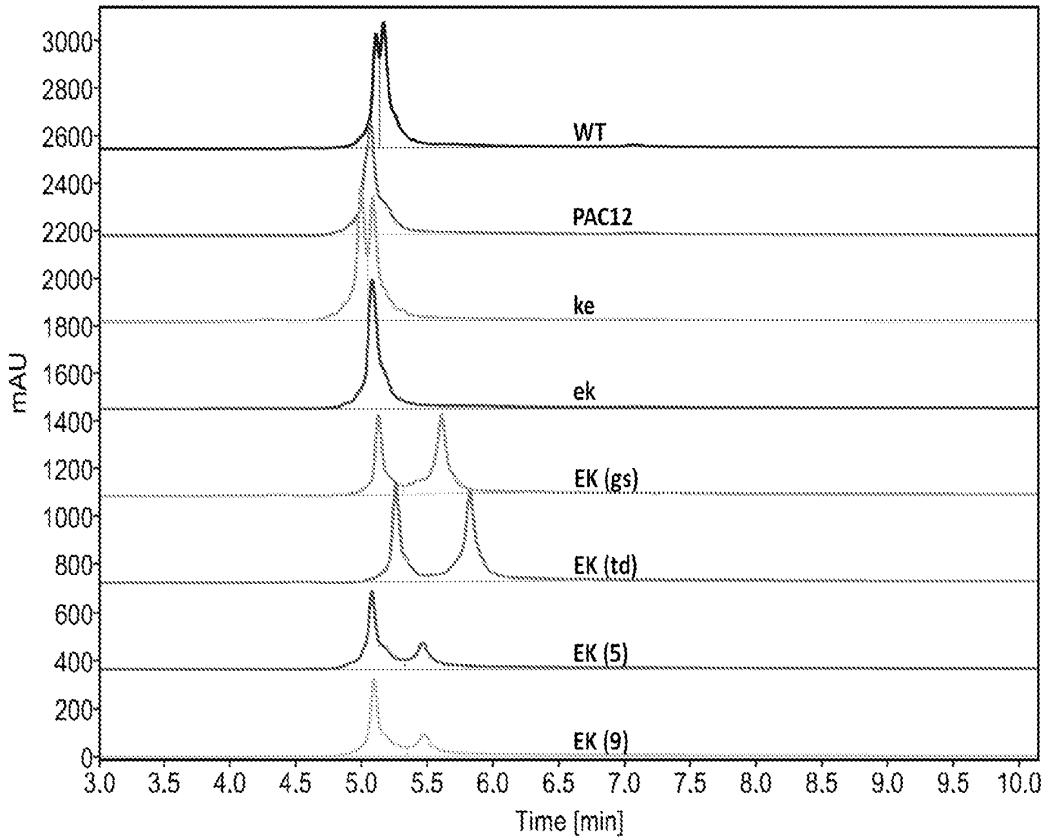
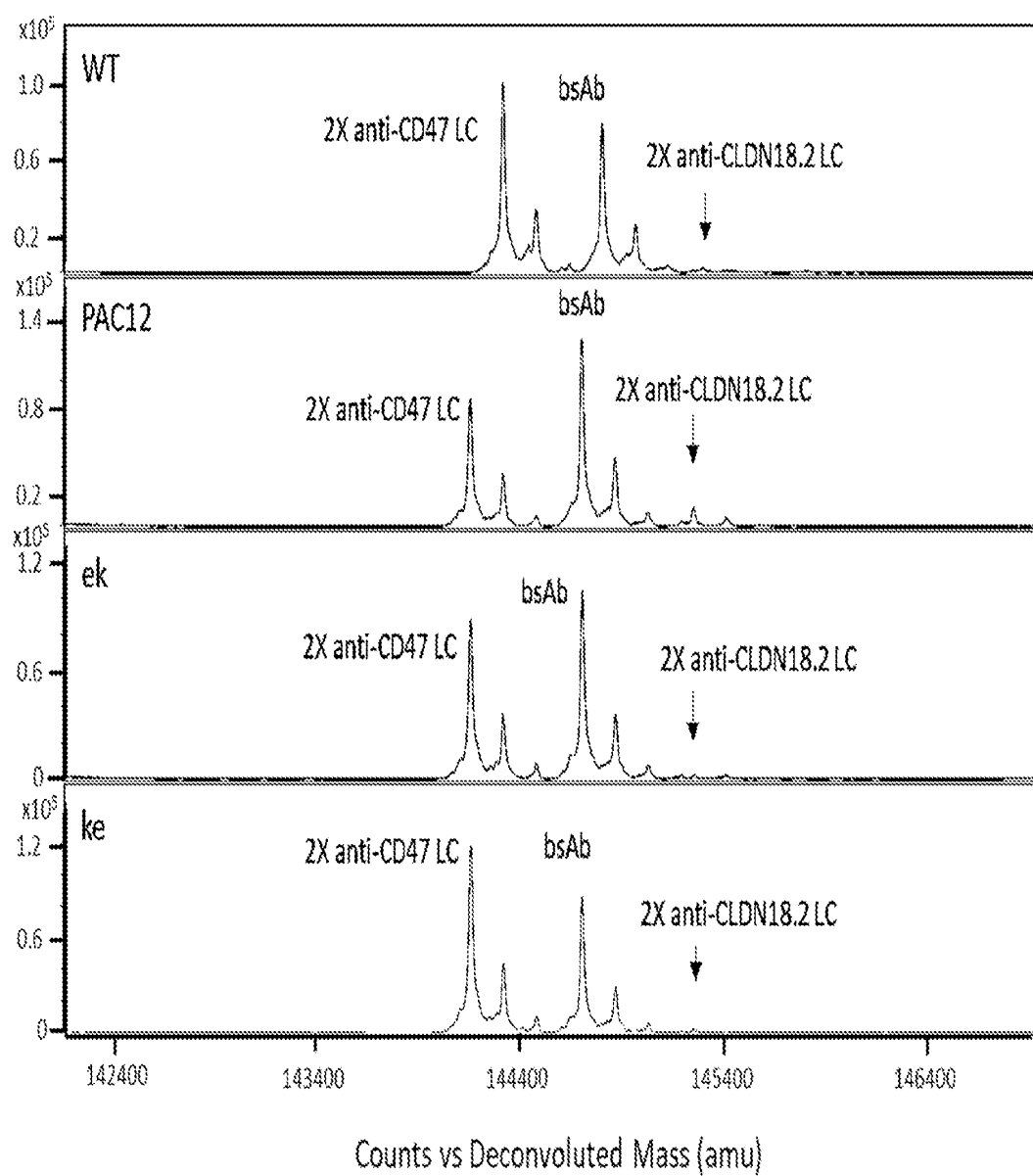
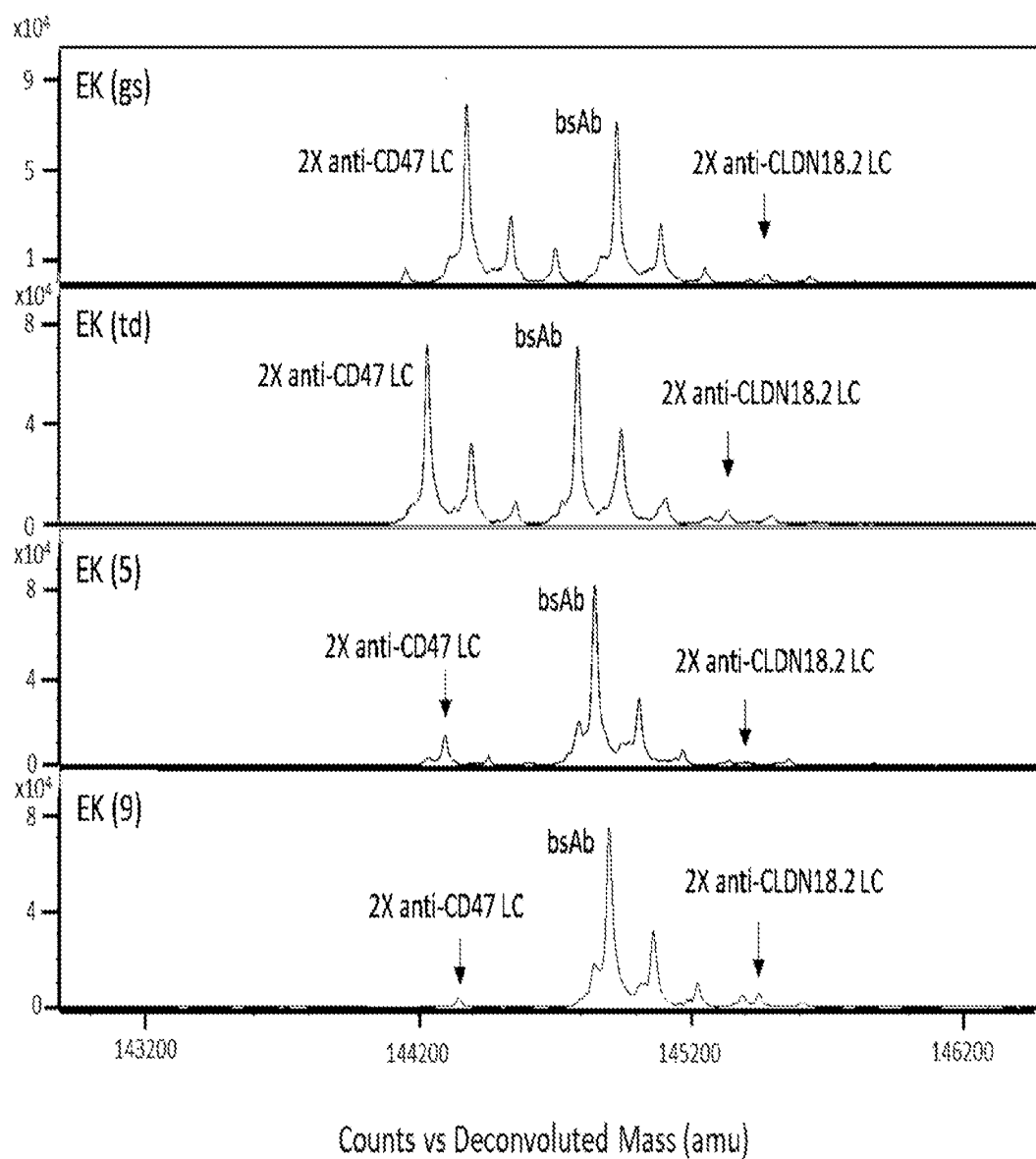


FIG. 4B

**FIG. 4C**

**FIG. 4D**

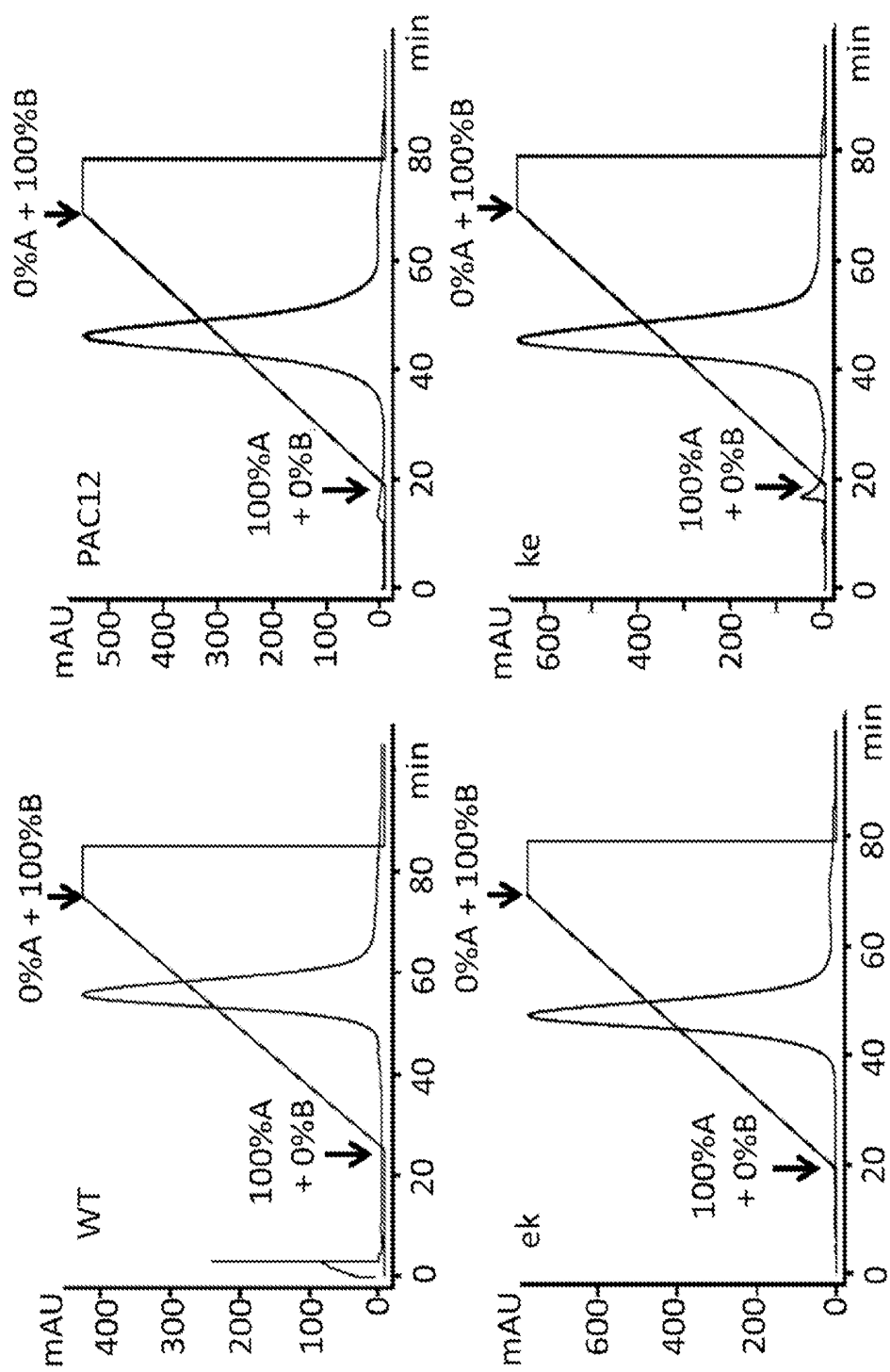


FIG. 4E

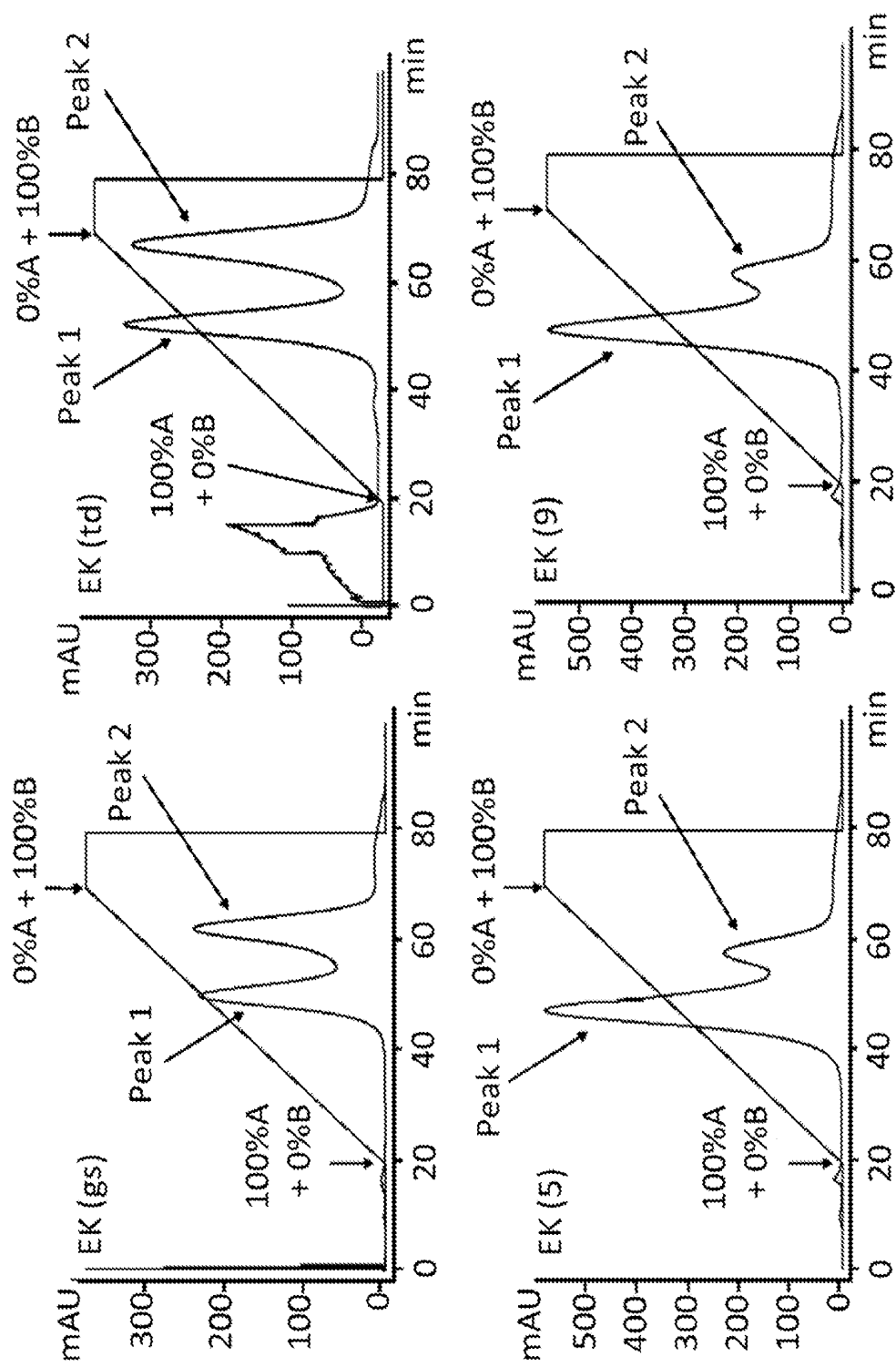


FIG. 4F

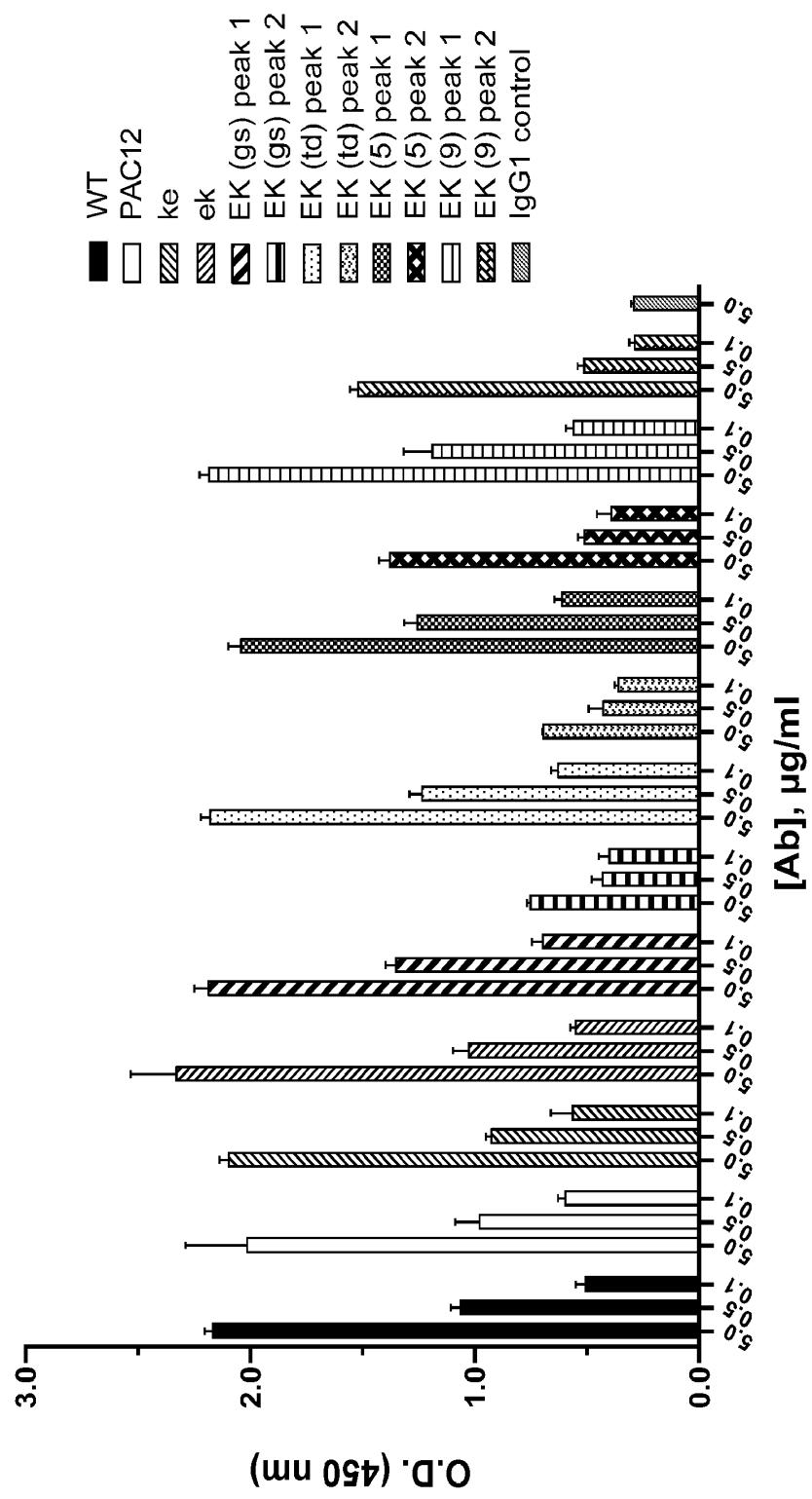
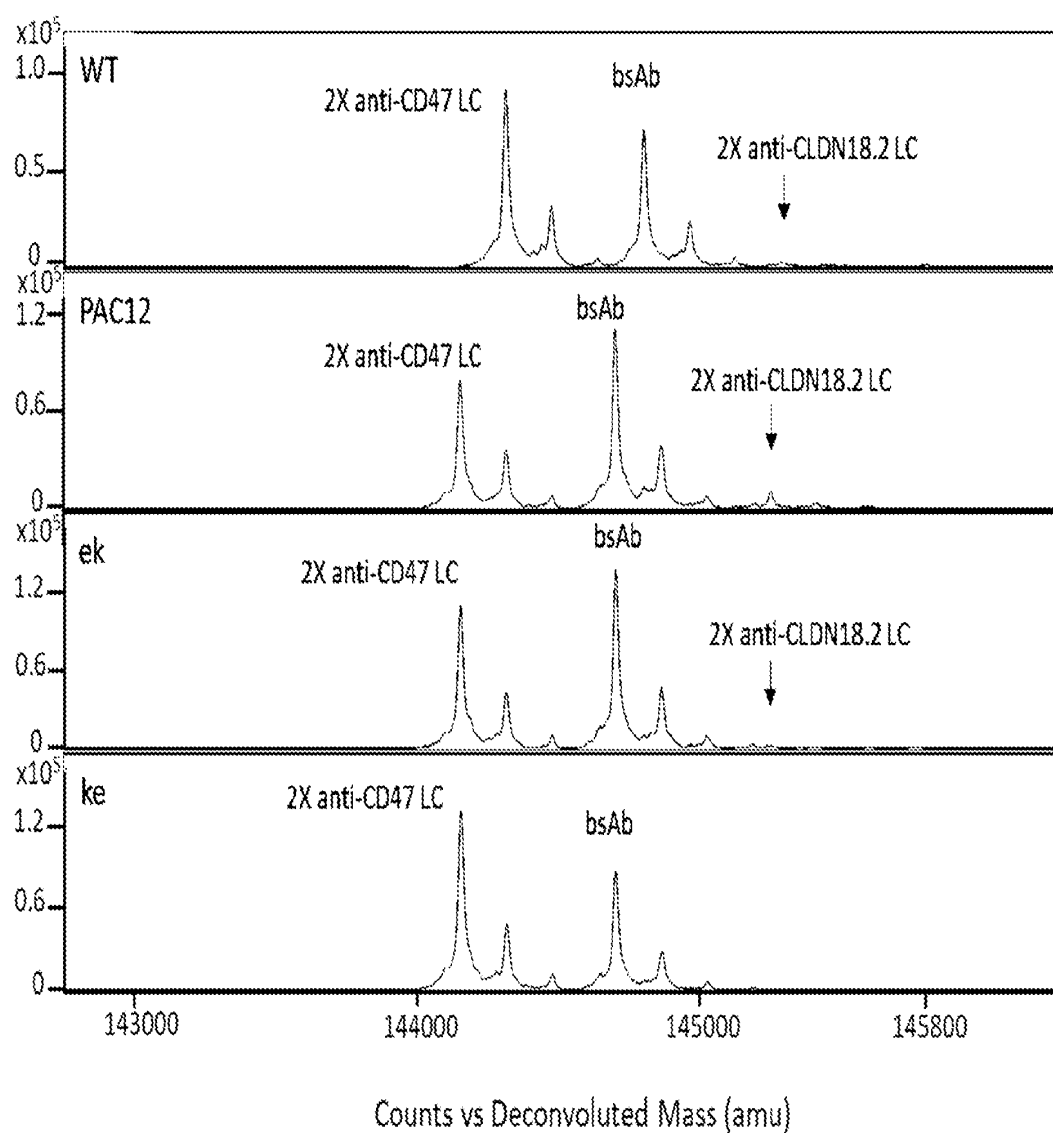
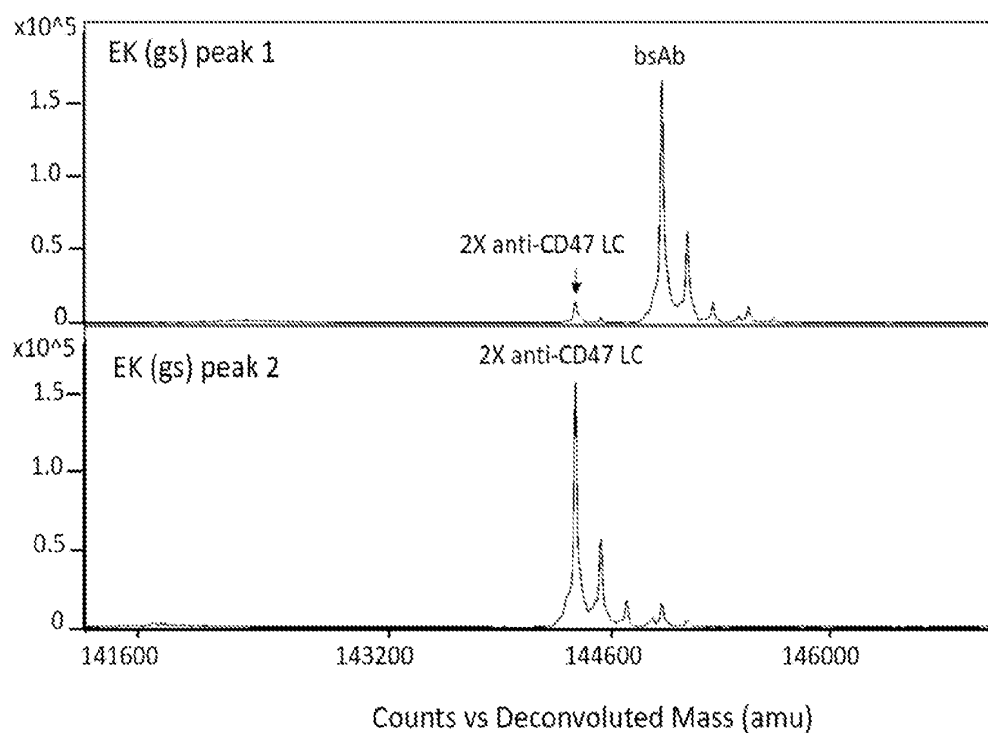
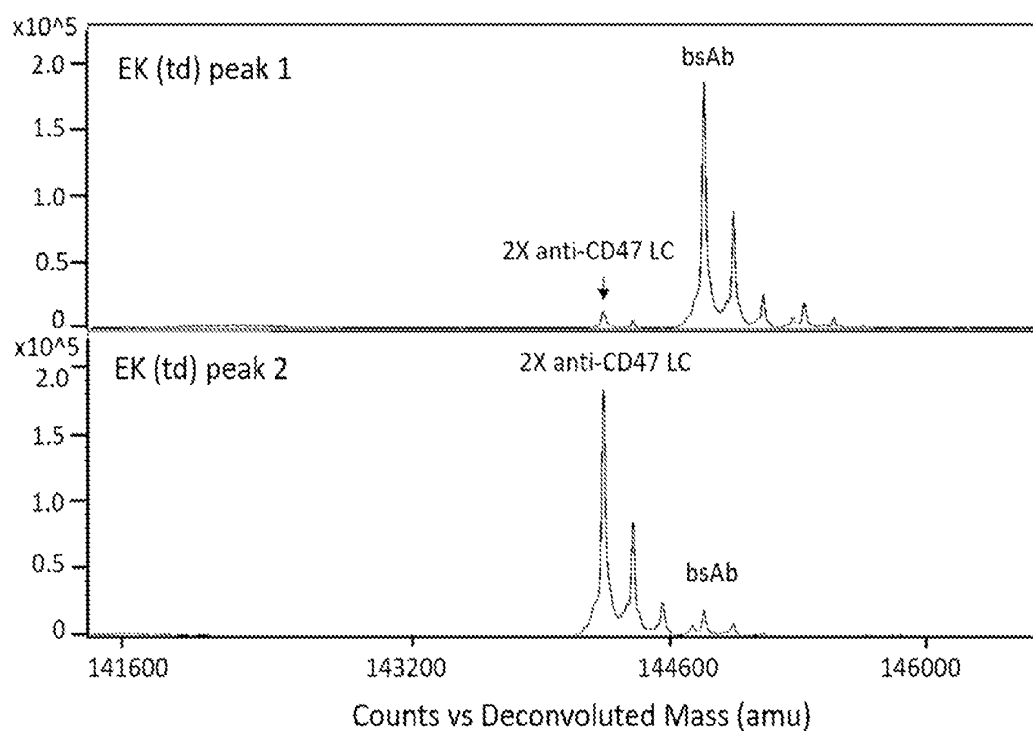
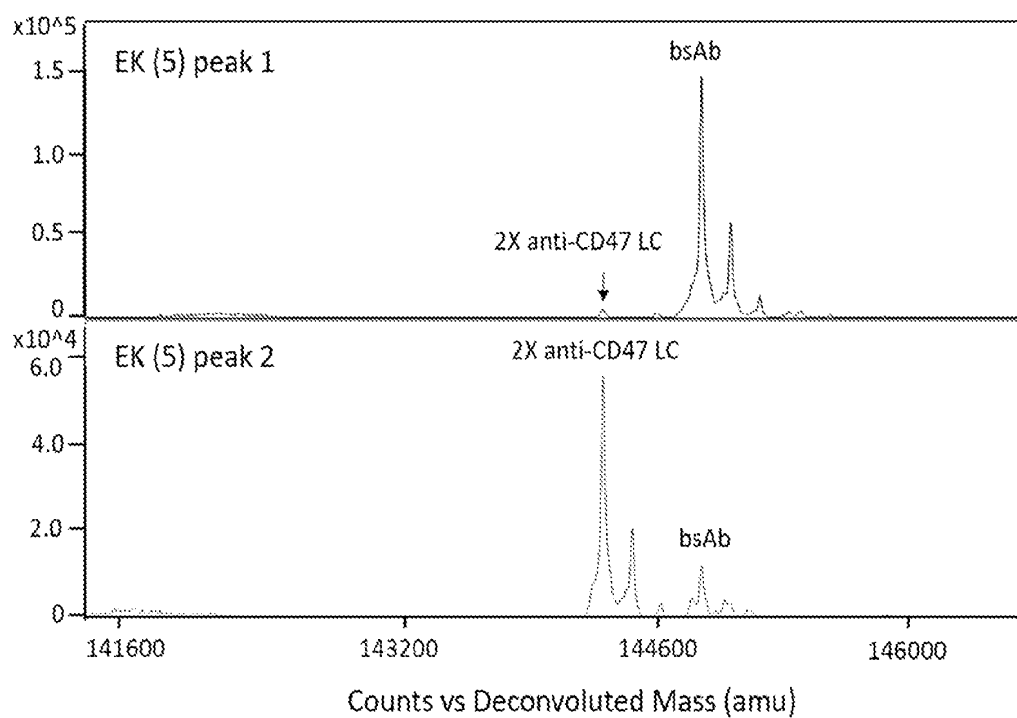
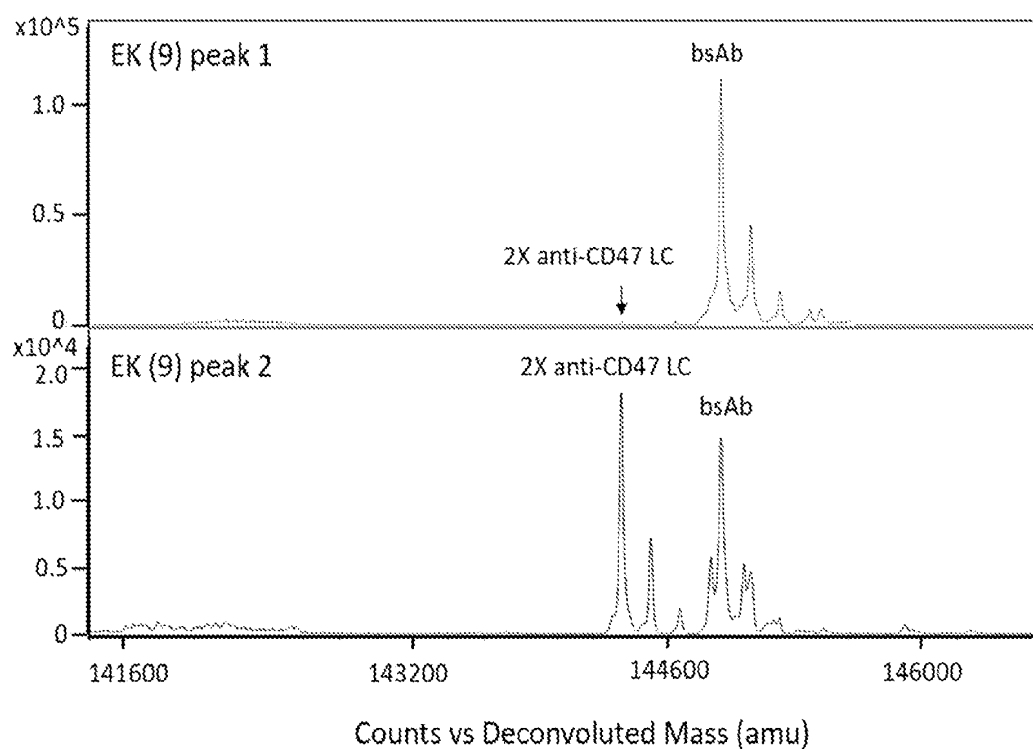


FIG. 4G

**FIG. 4H**

**FIG. 4I****FIG. 4J**

**FIG. 4K****FIG. 4L**

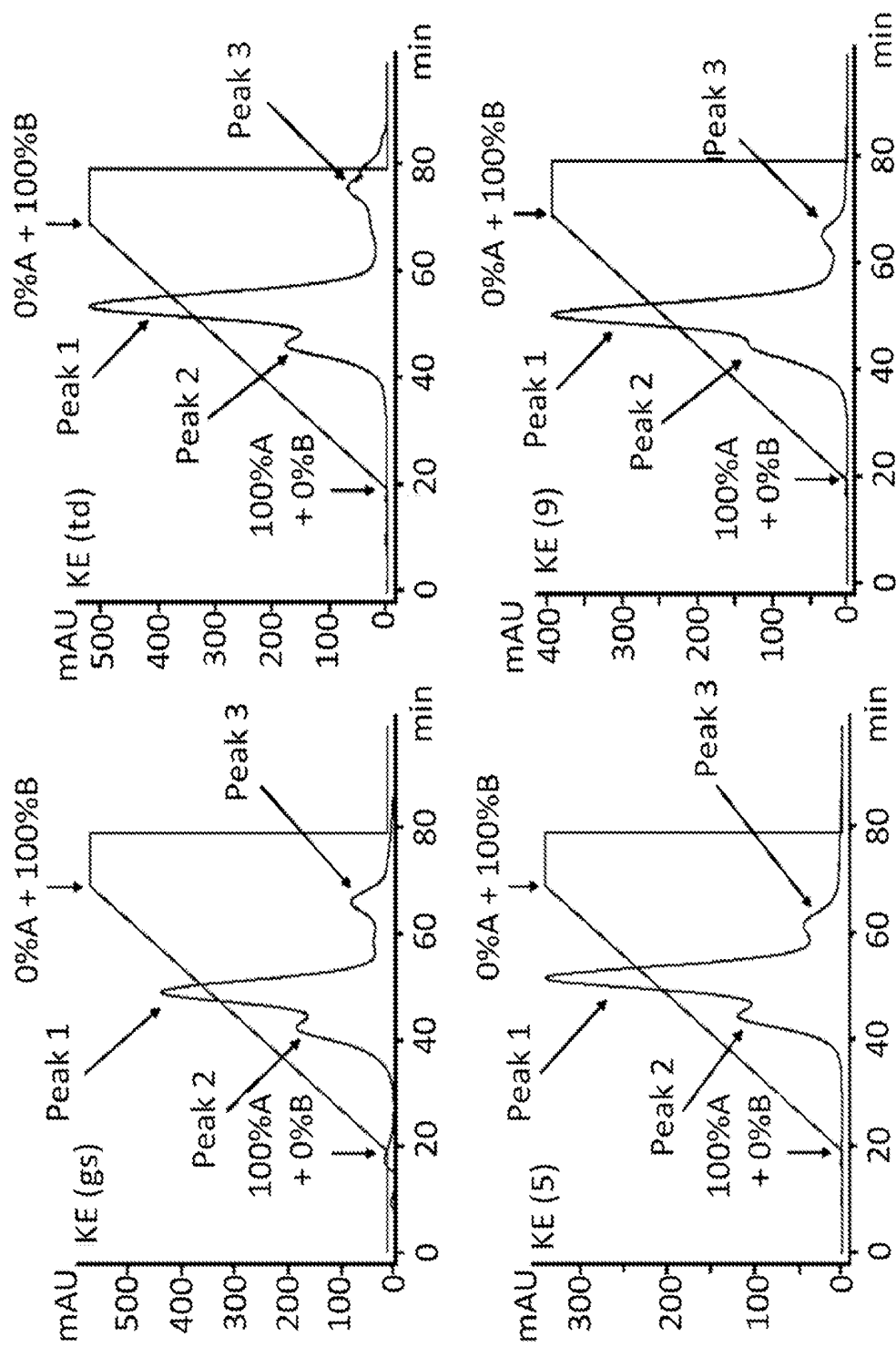


FIG. 5A

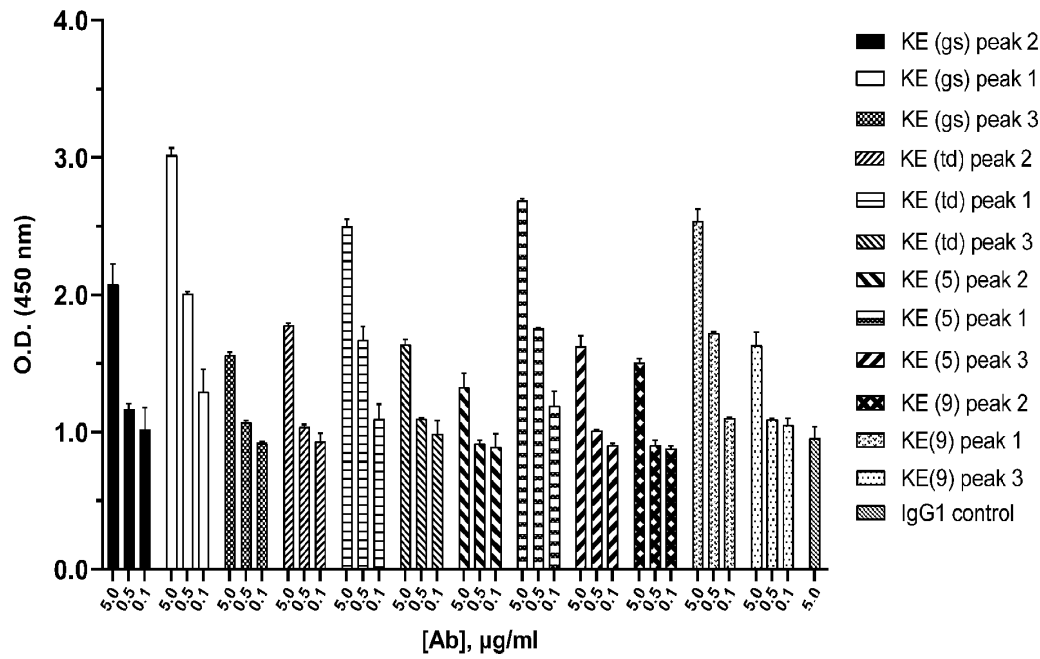


FIG. 5B

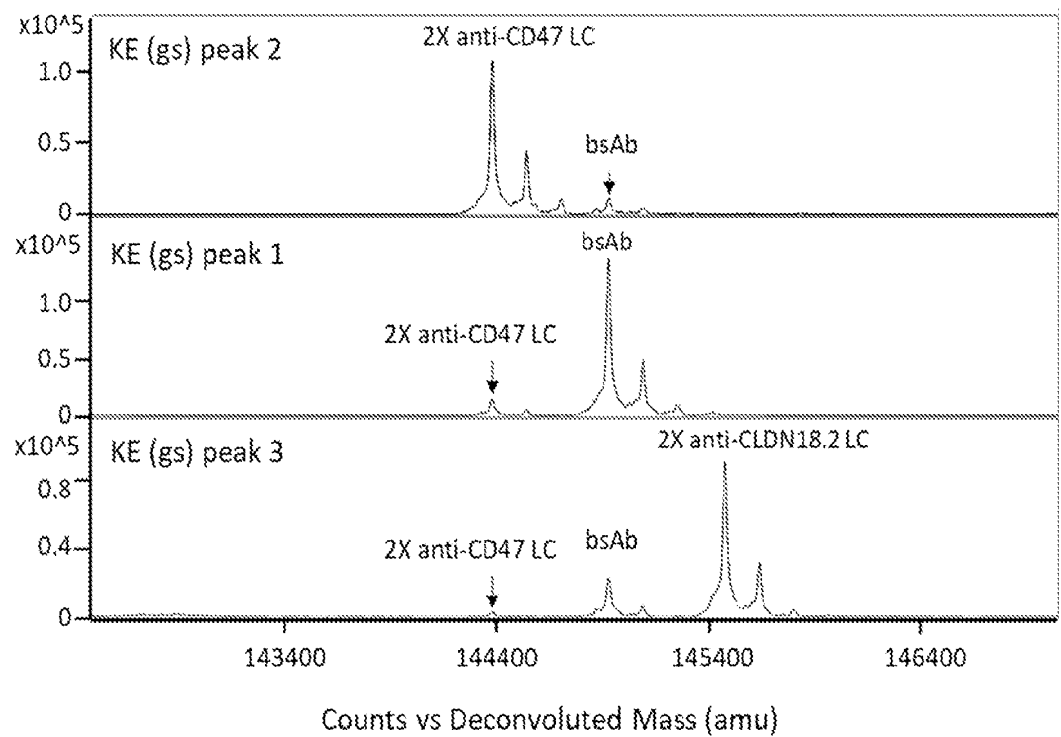
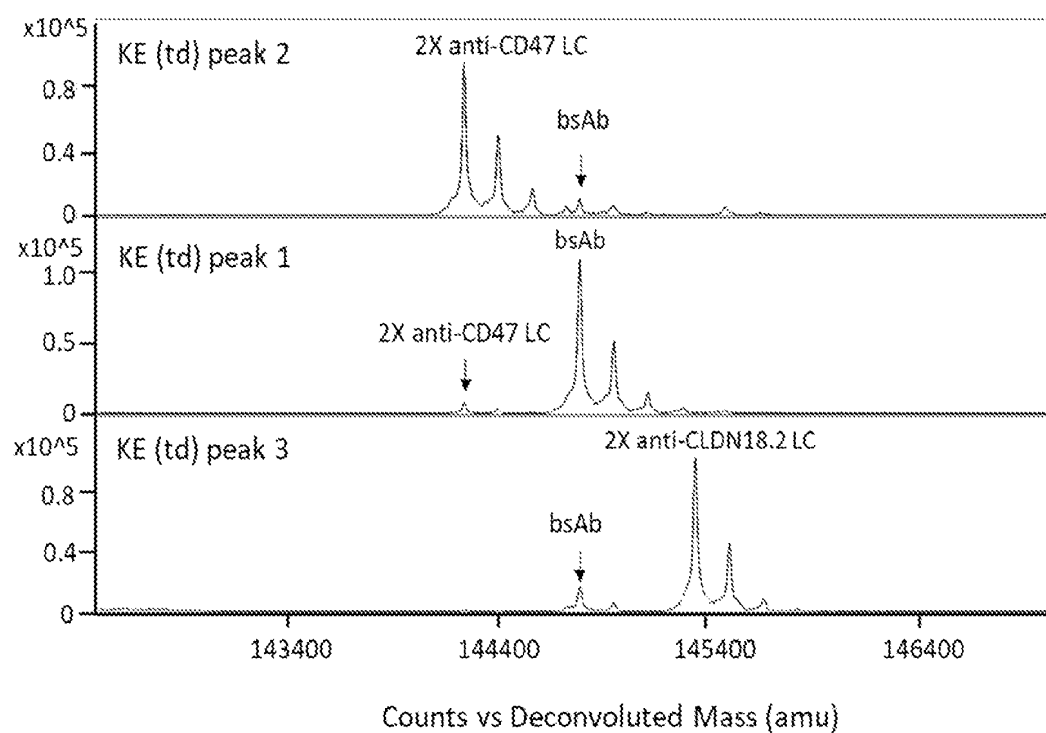
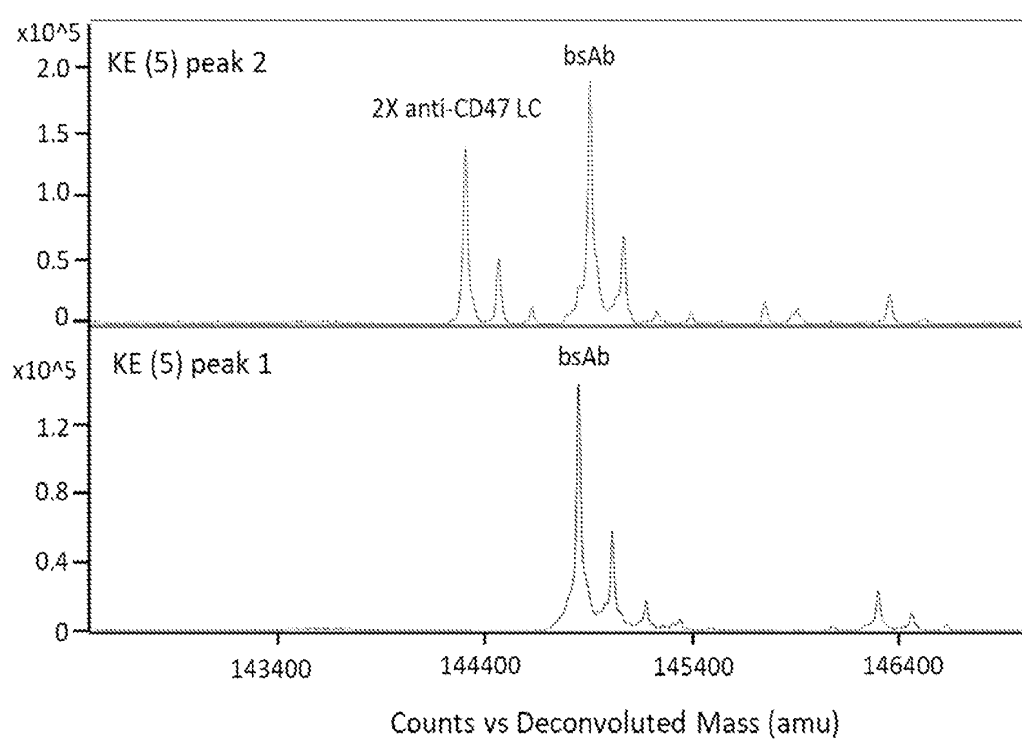
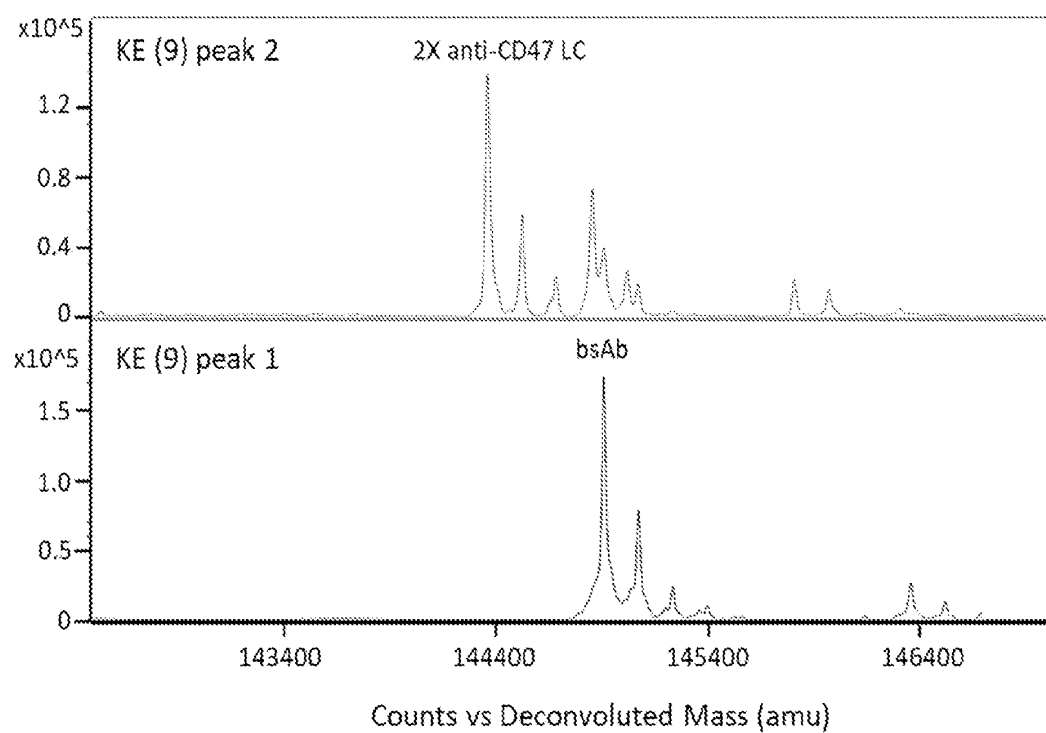


FIG. 5C

**FIG. 5D****FIG. 5E**

**FIG. 5F**

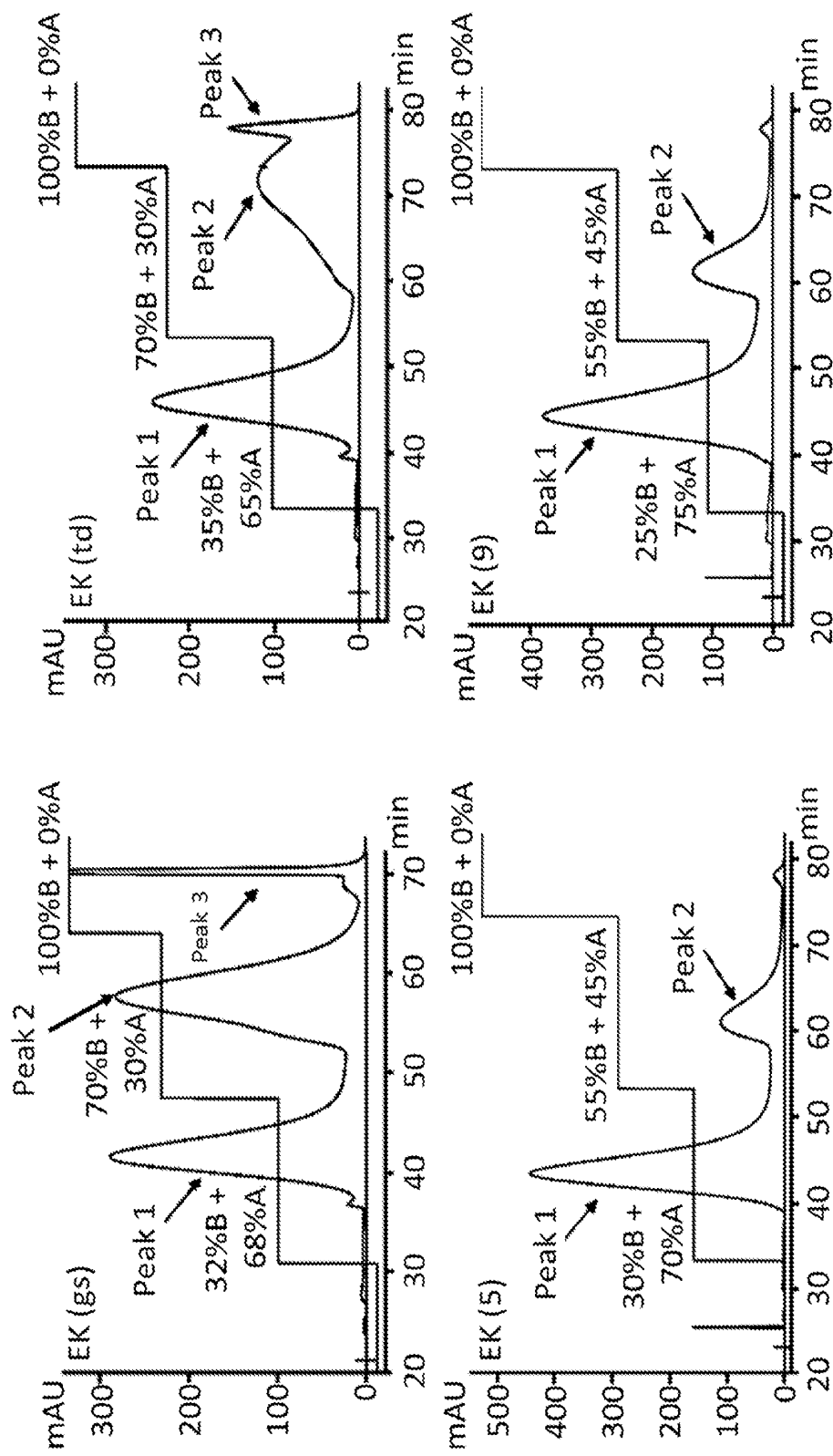


FIG. 6A

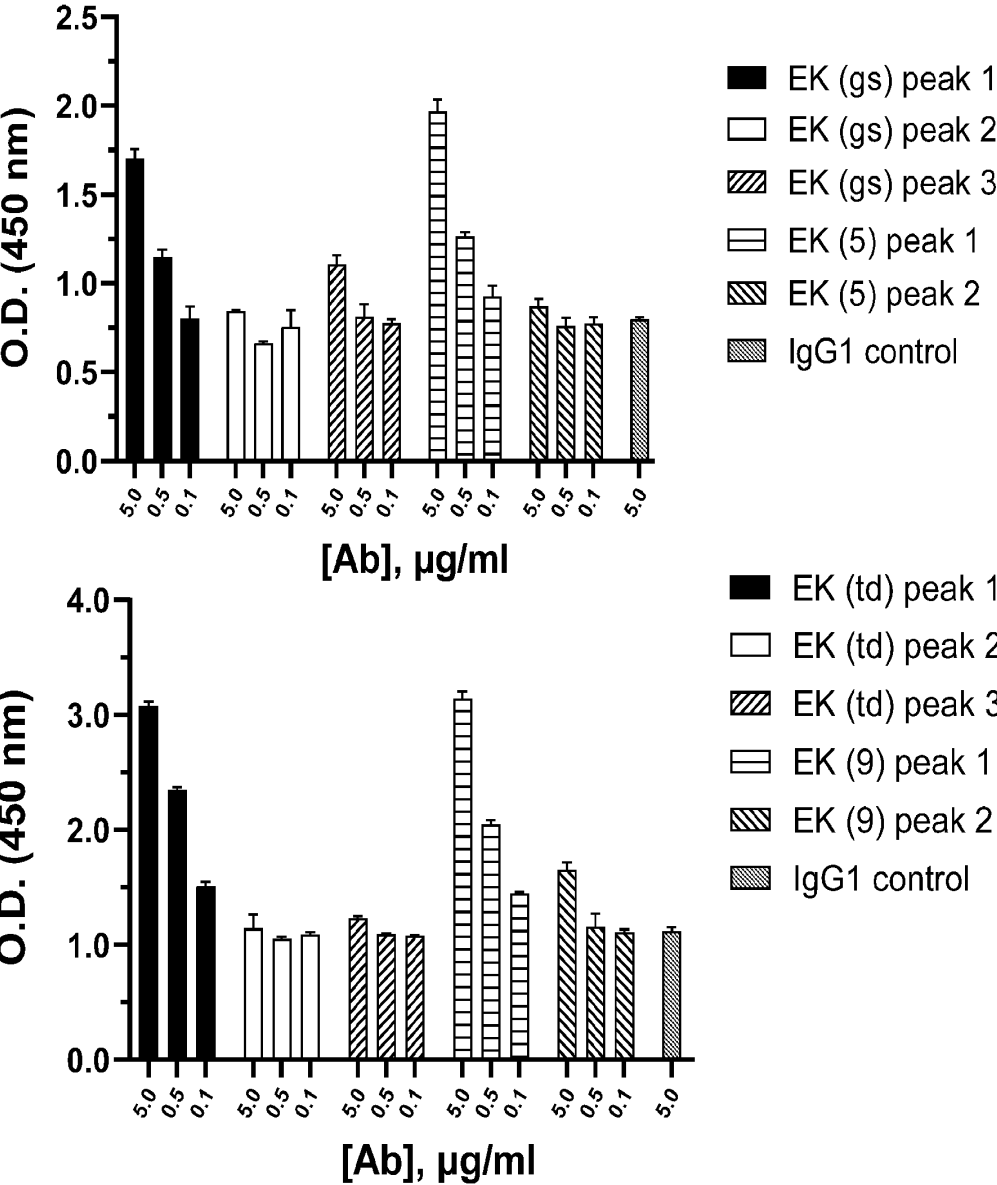
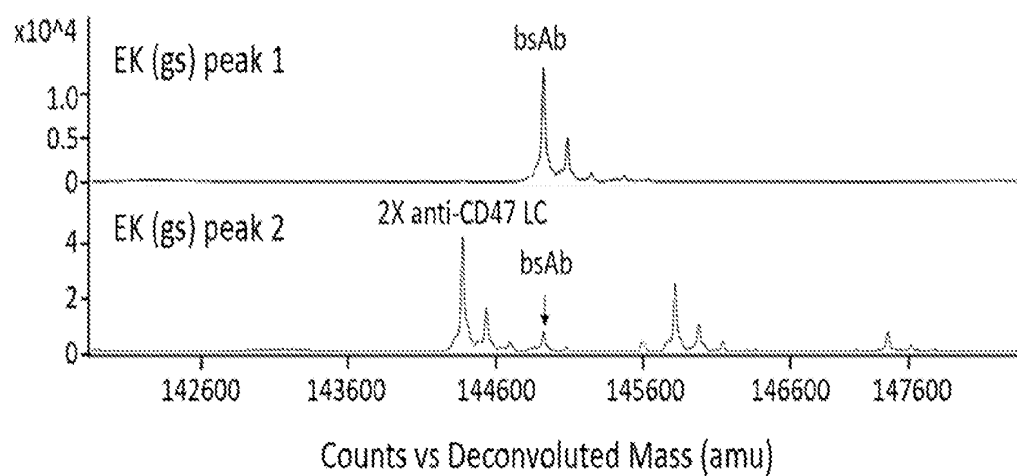
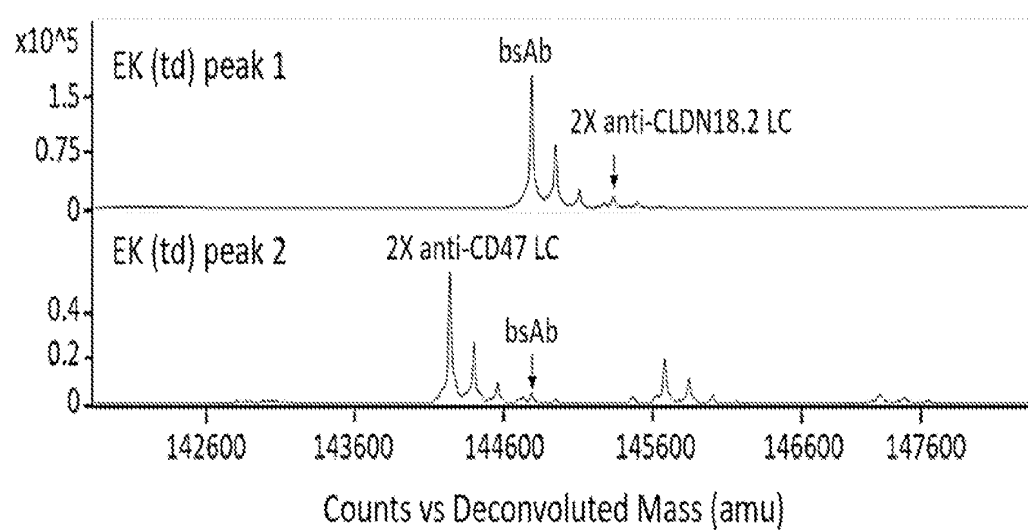
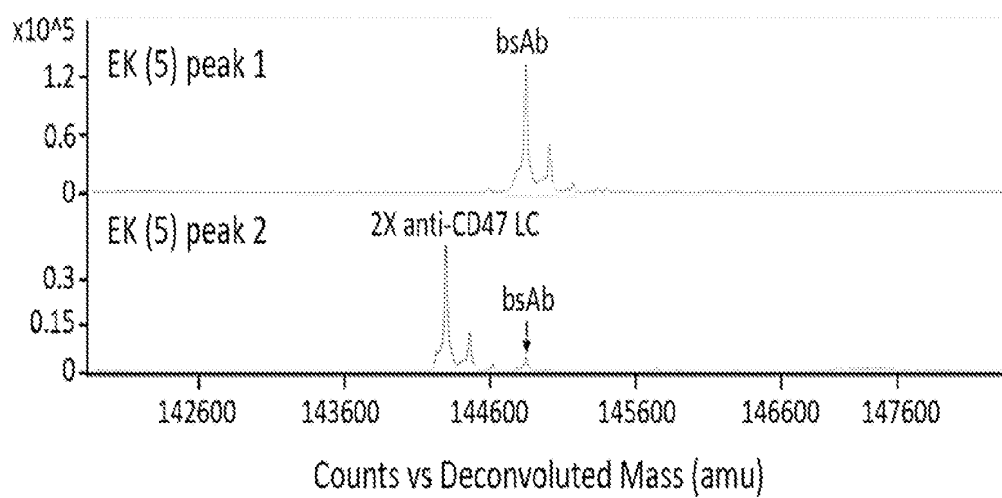
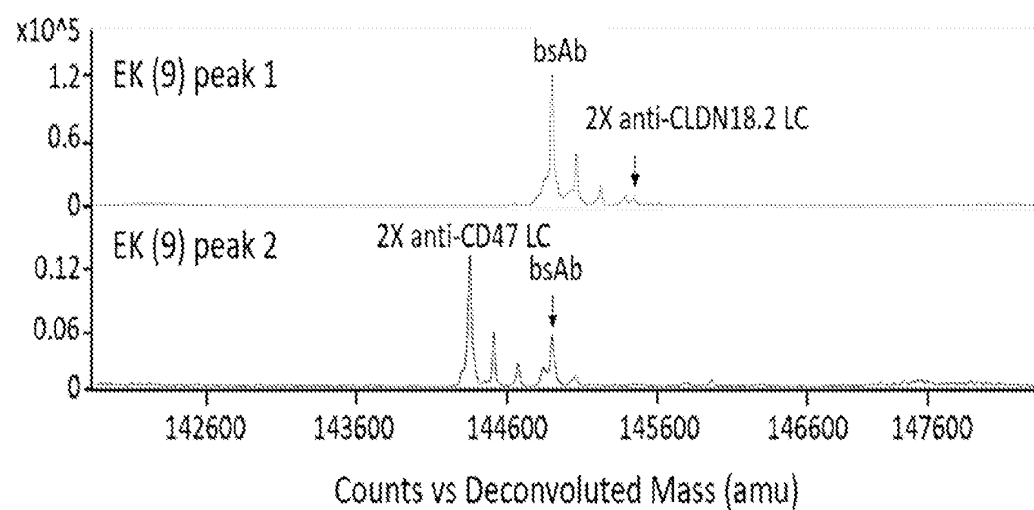


FIG. 6B

**FIG. 6C****FIG. 6D**

**FIG. 6E****FIG. 6F**

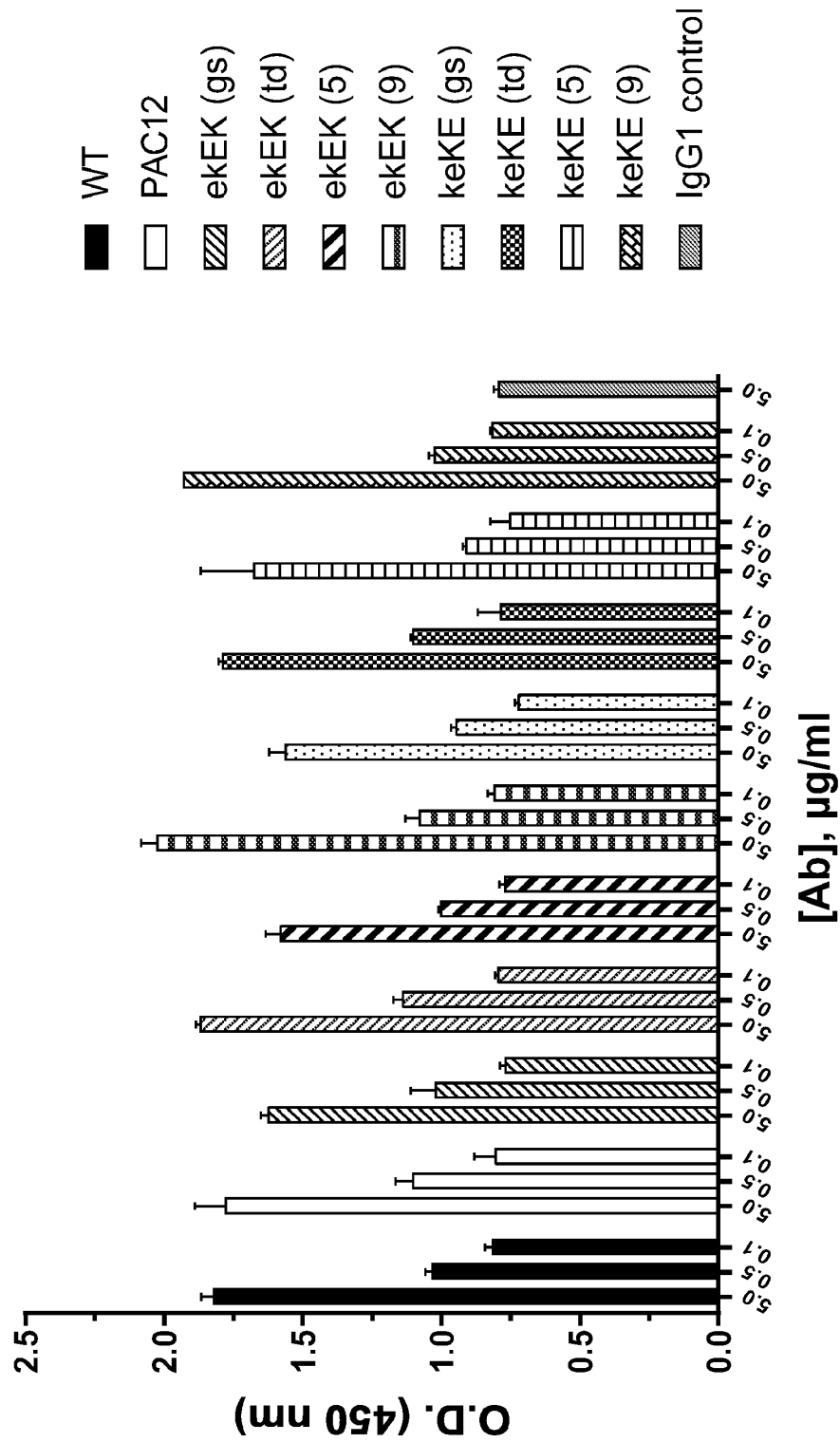


FIG. 7A

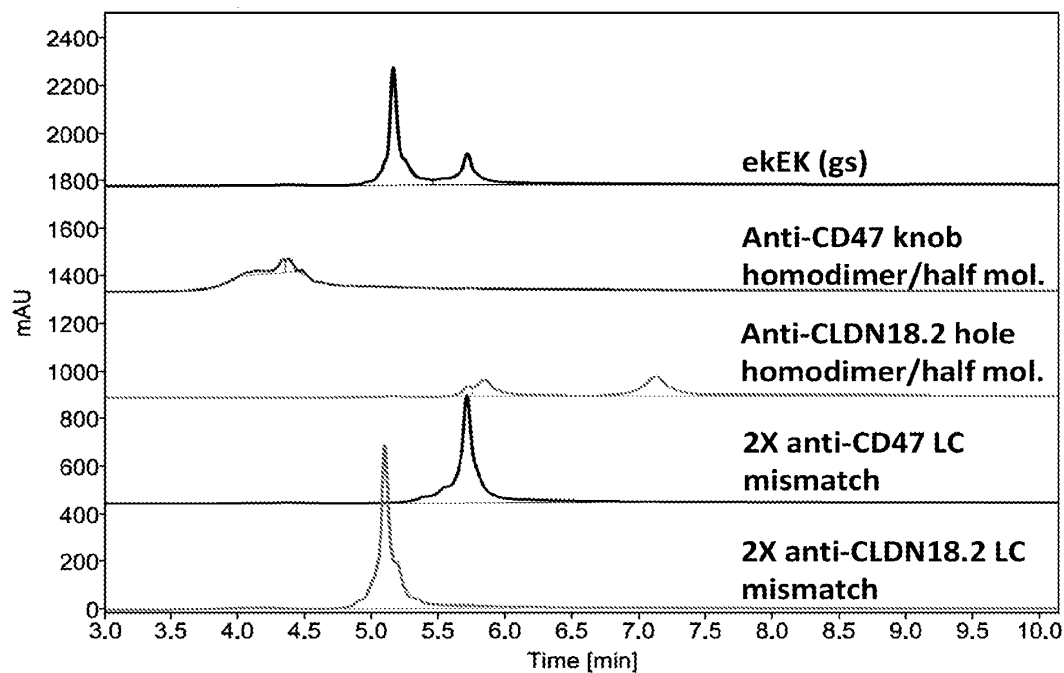


FIG. 7B

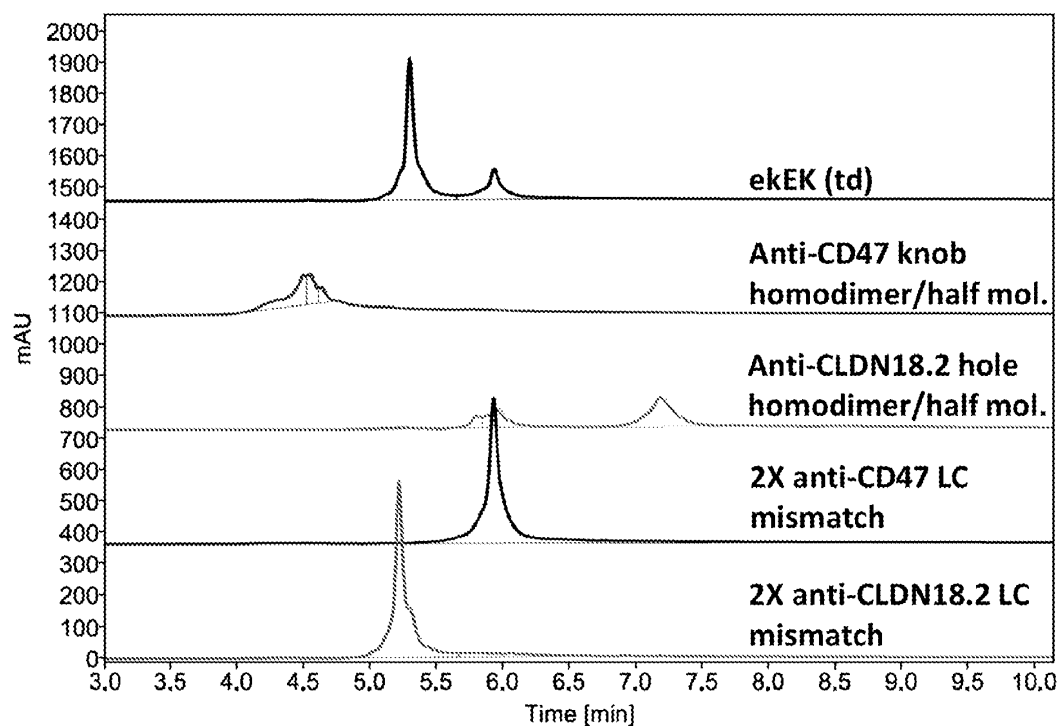
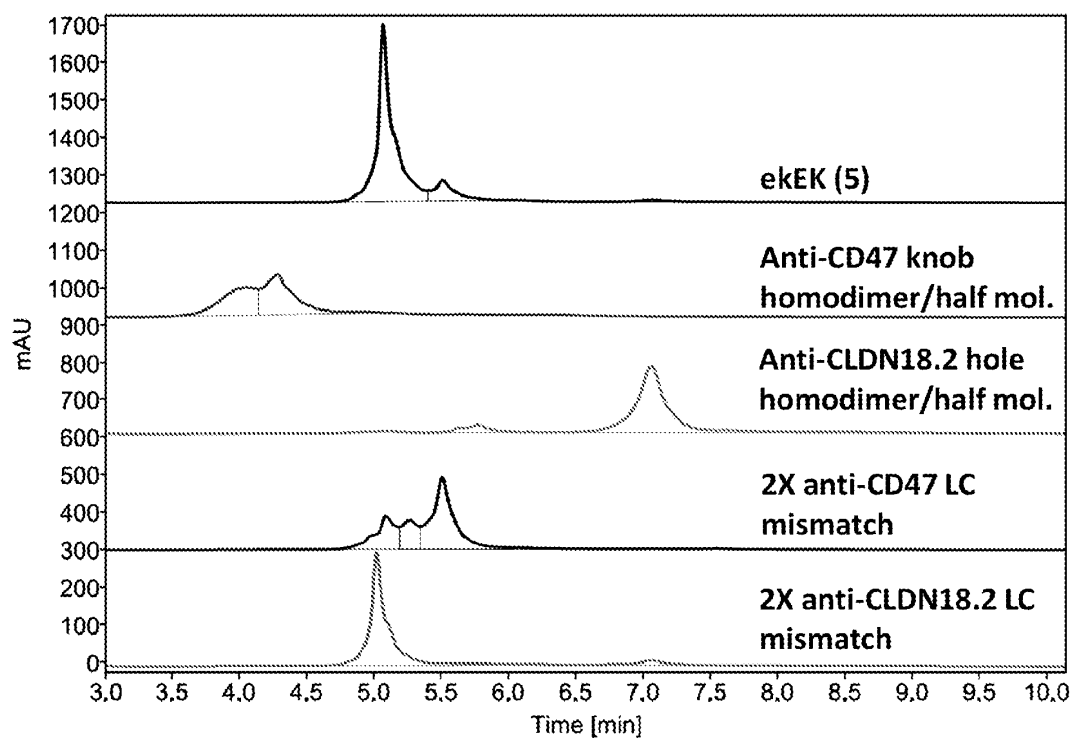
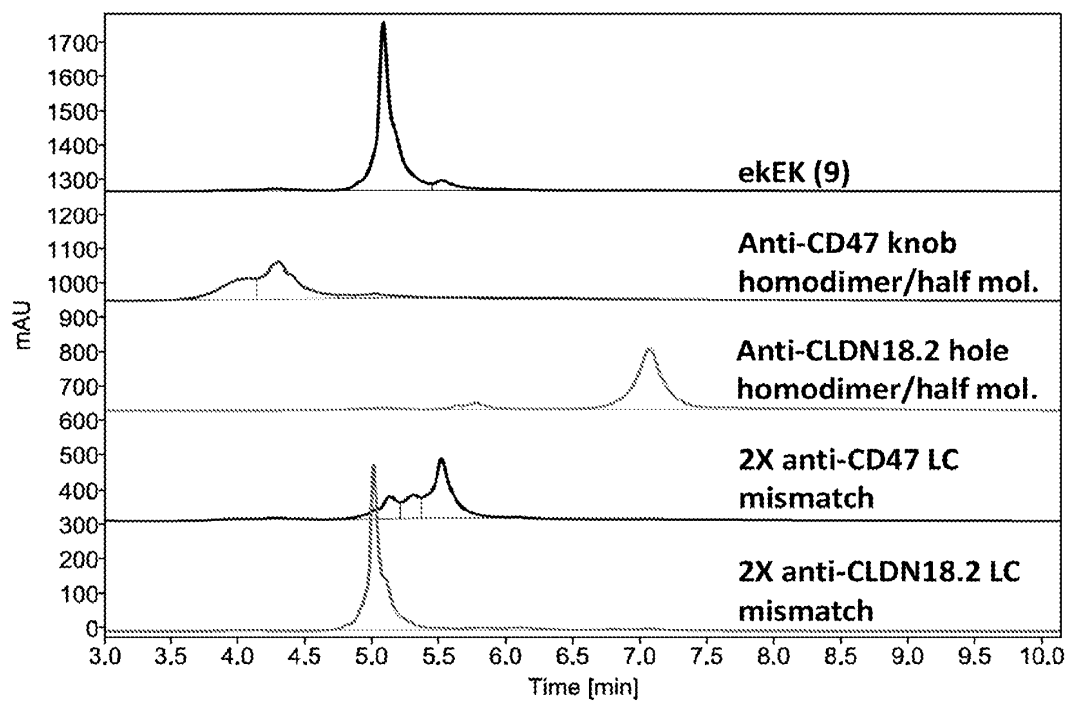


FIG. 7C

**FIG. 7D****FIG. 7E**

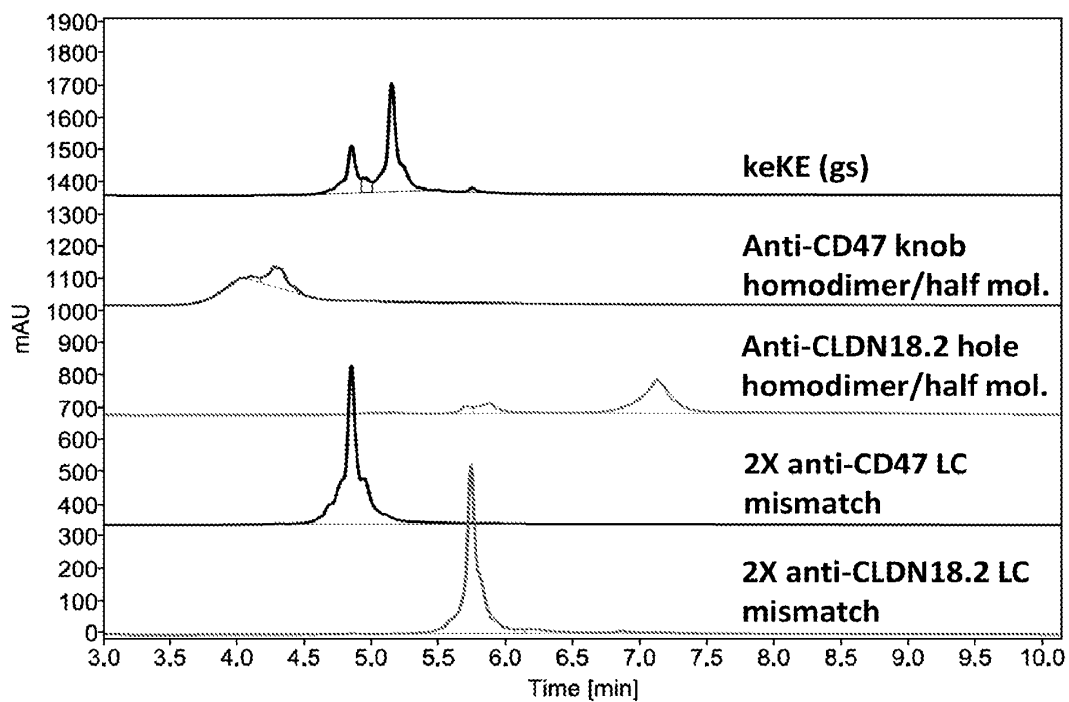


FIG. 7F

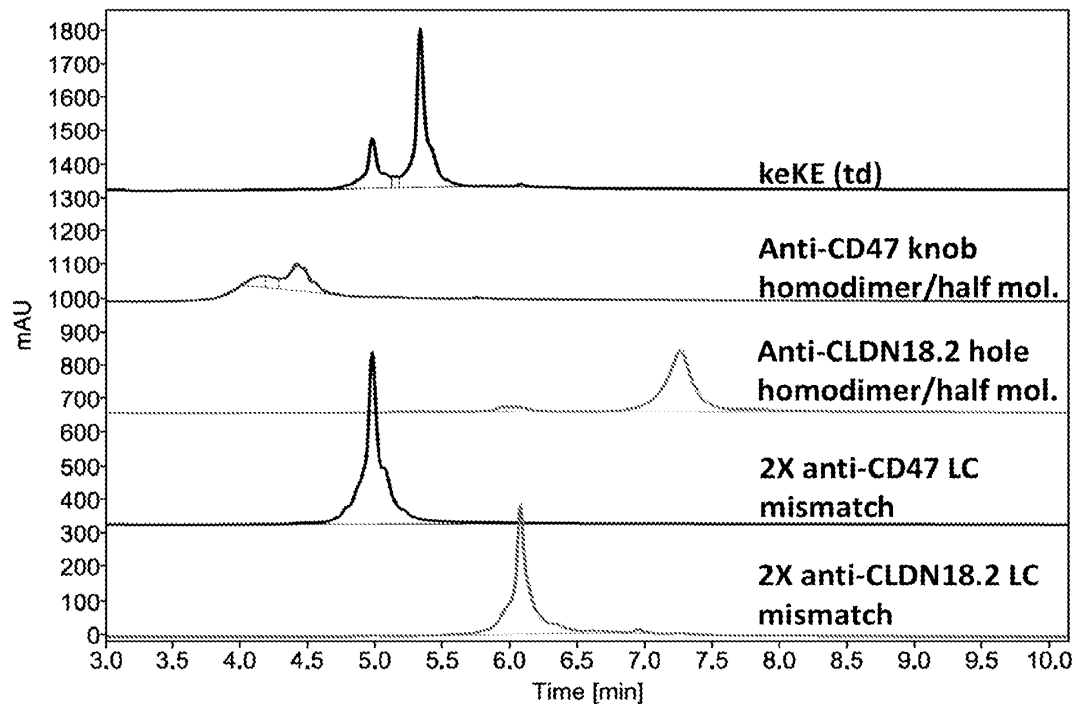


FIG. 7G

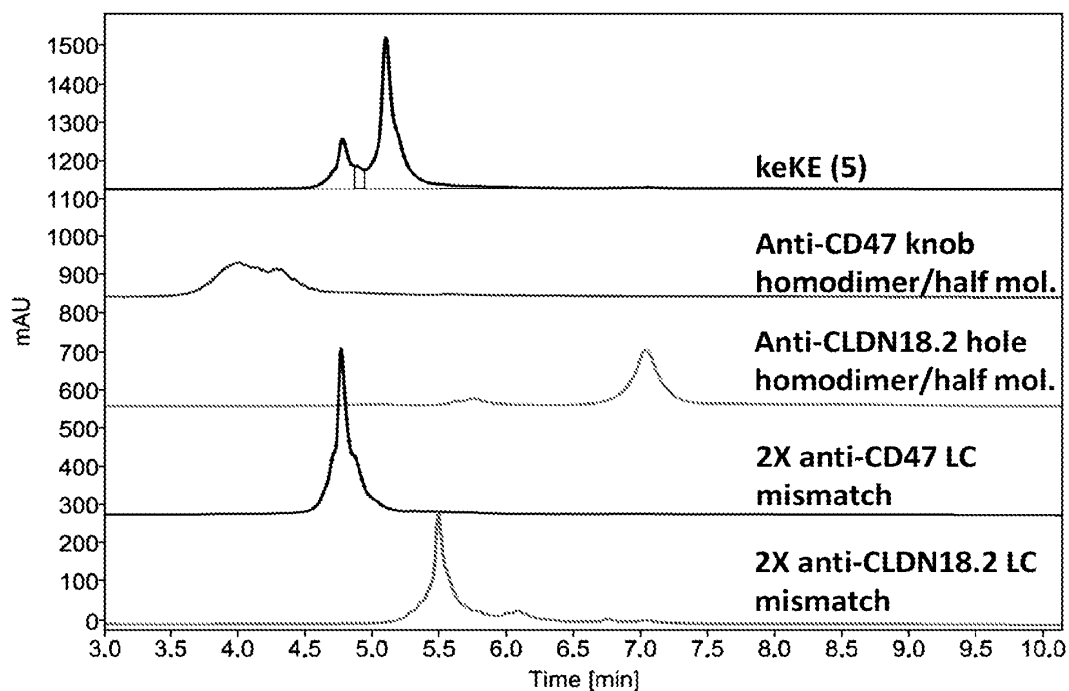


FIG. 7H

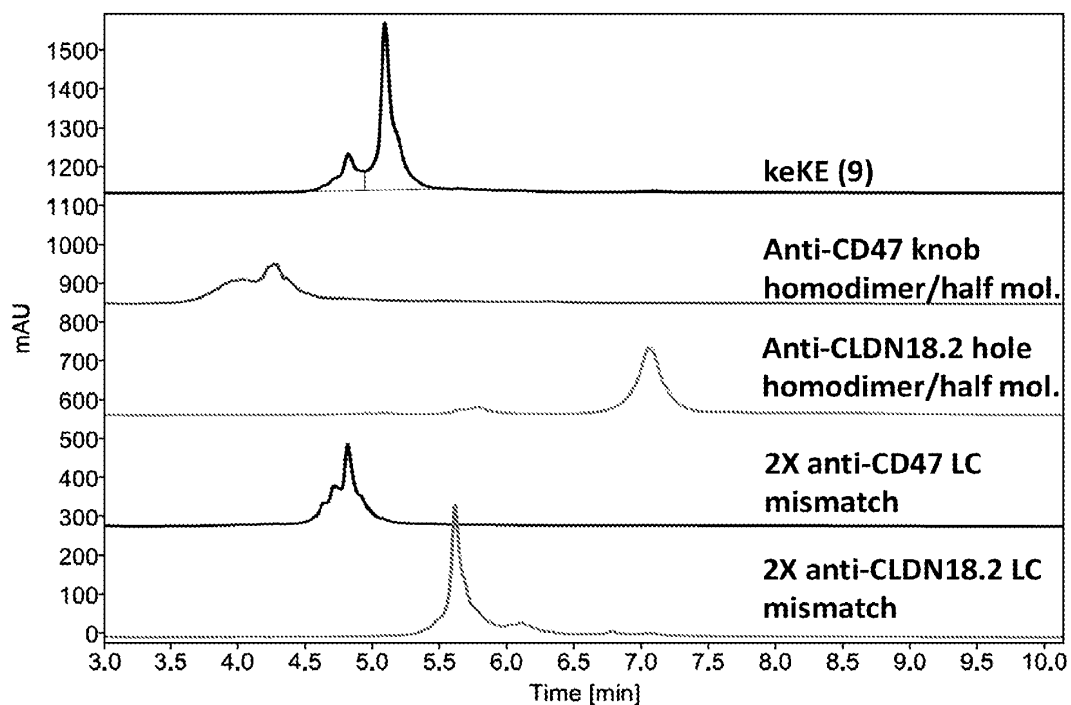
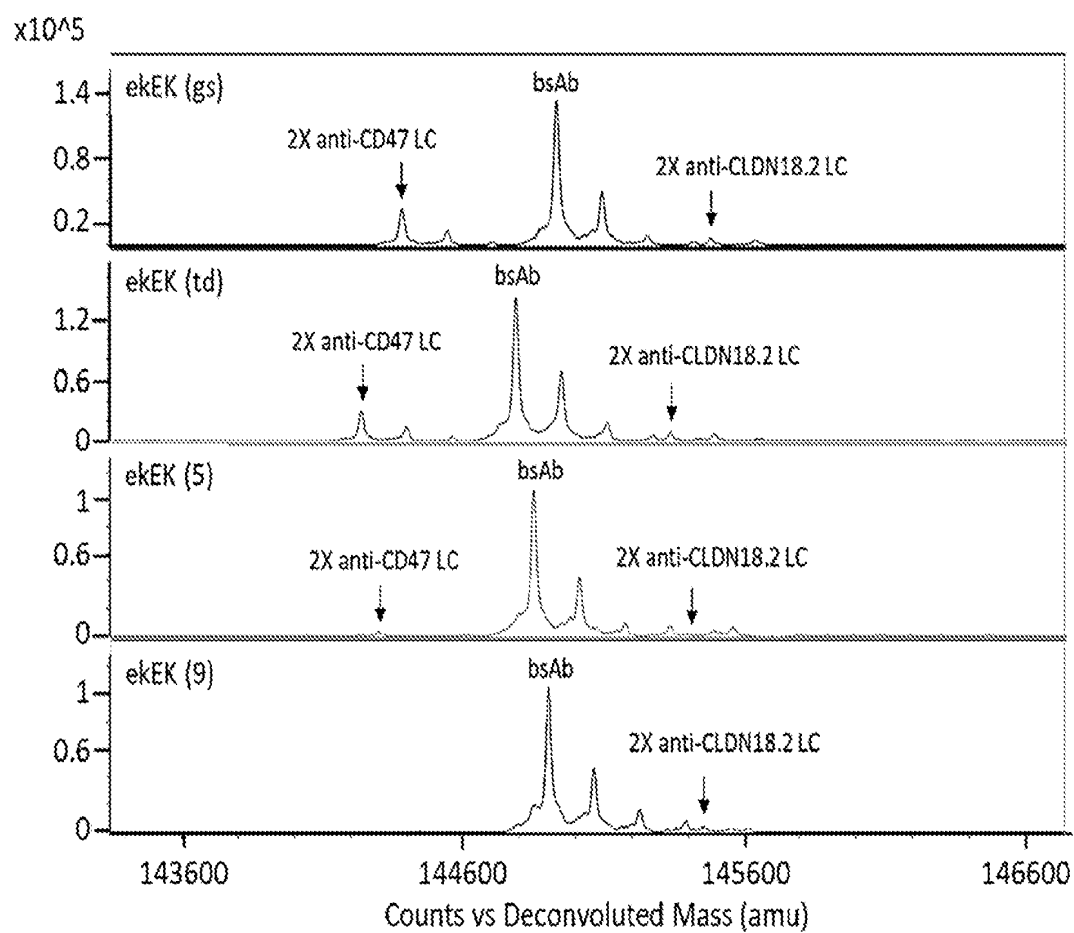
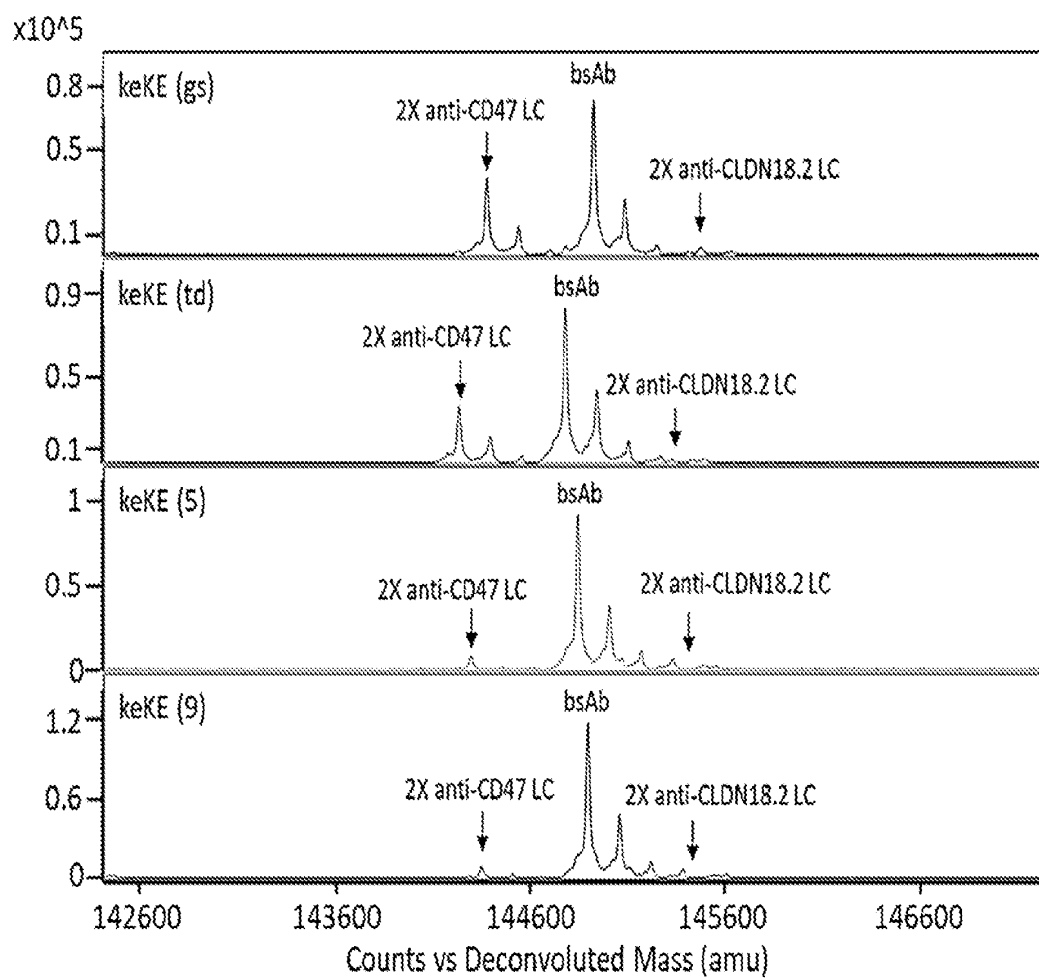


FIG. 7I

**FIG. 7J**

**FIG. 7K**

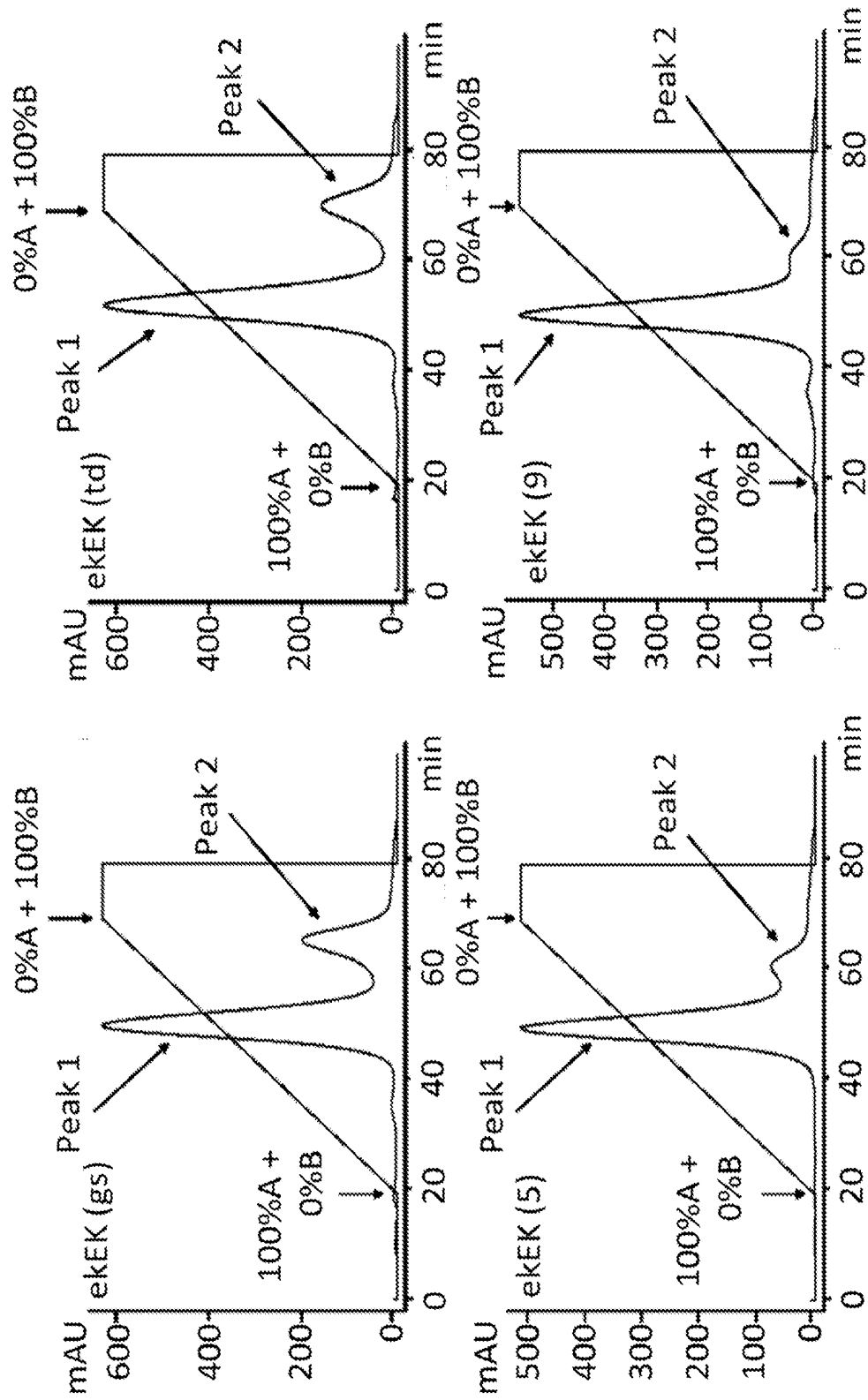


FIG. 8A

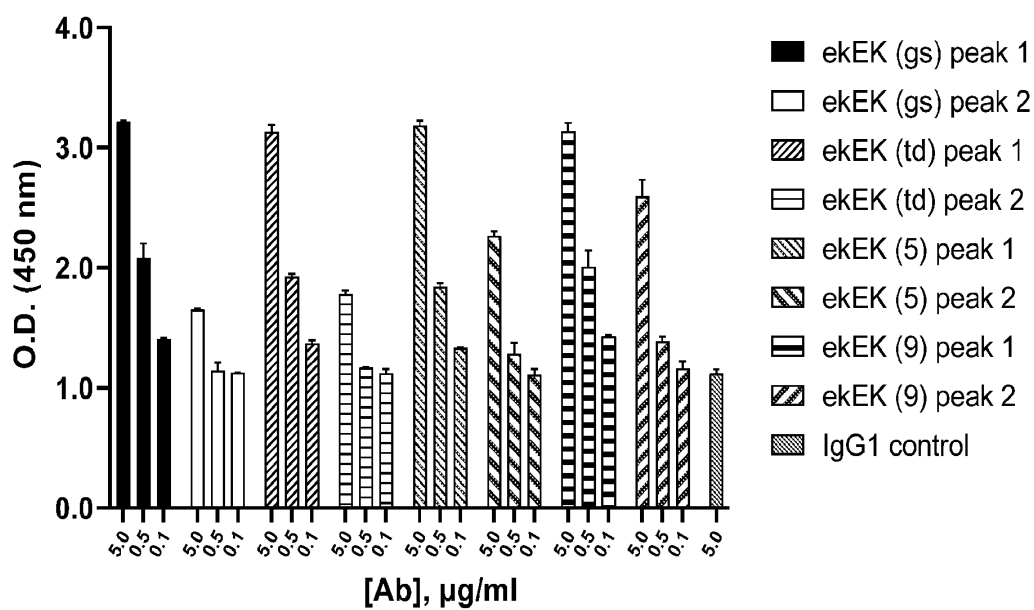


FIG. 8B

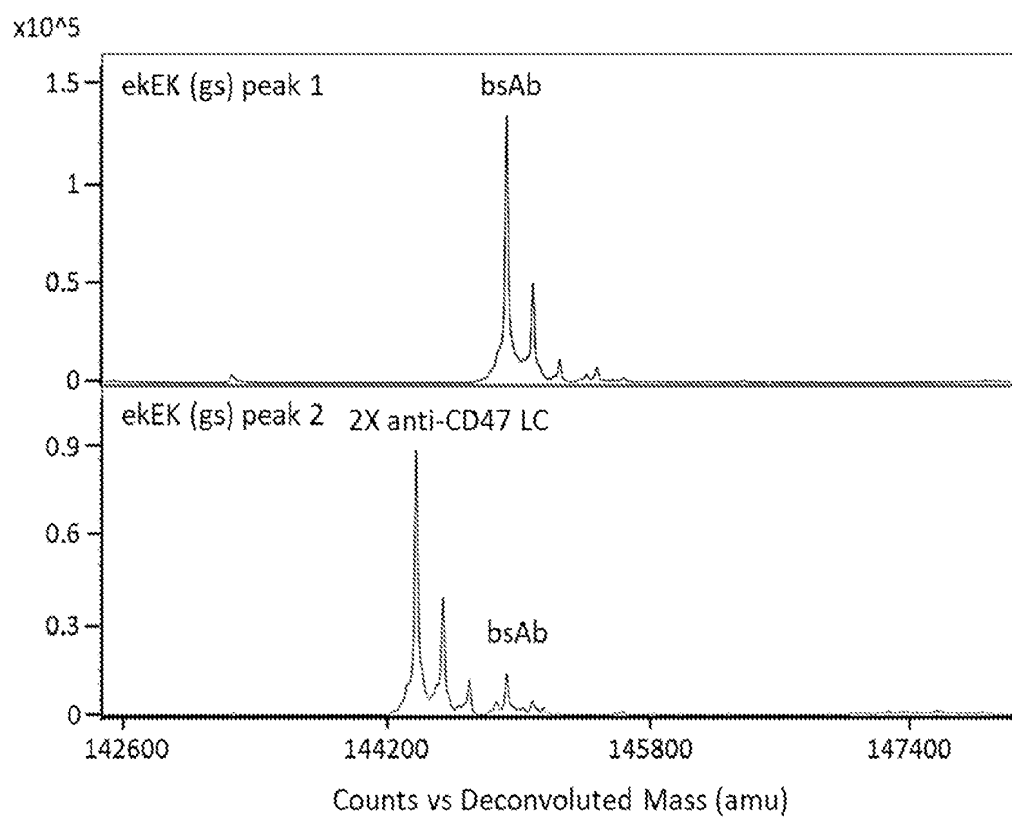
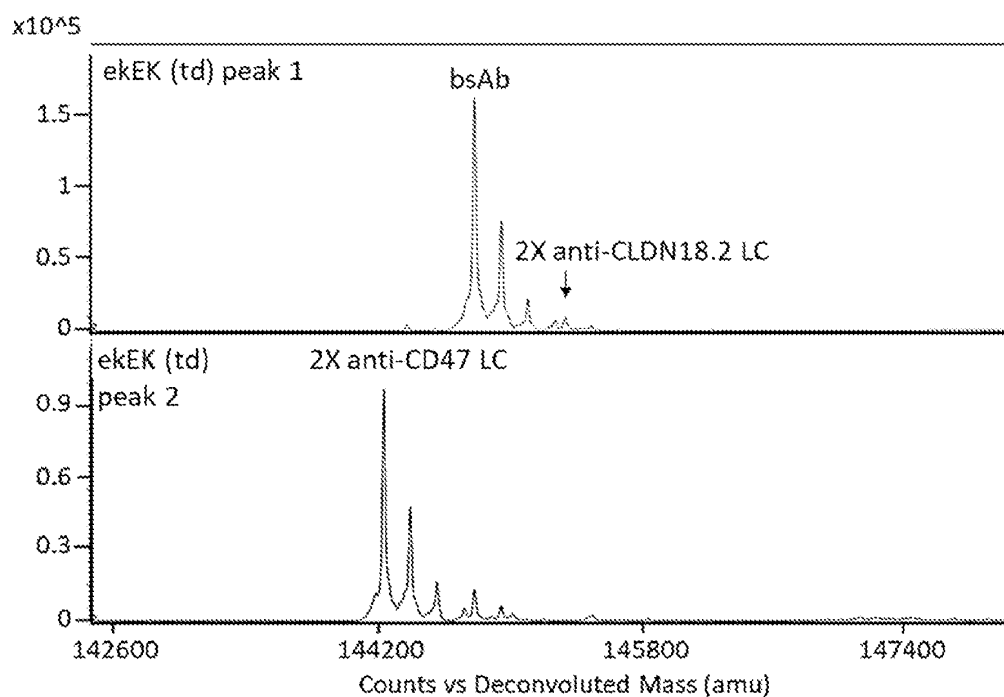
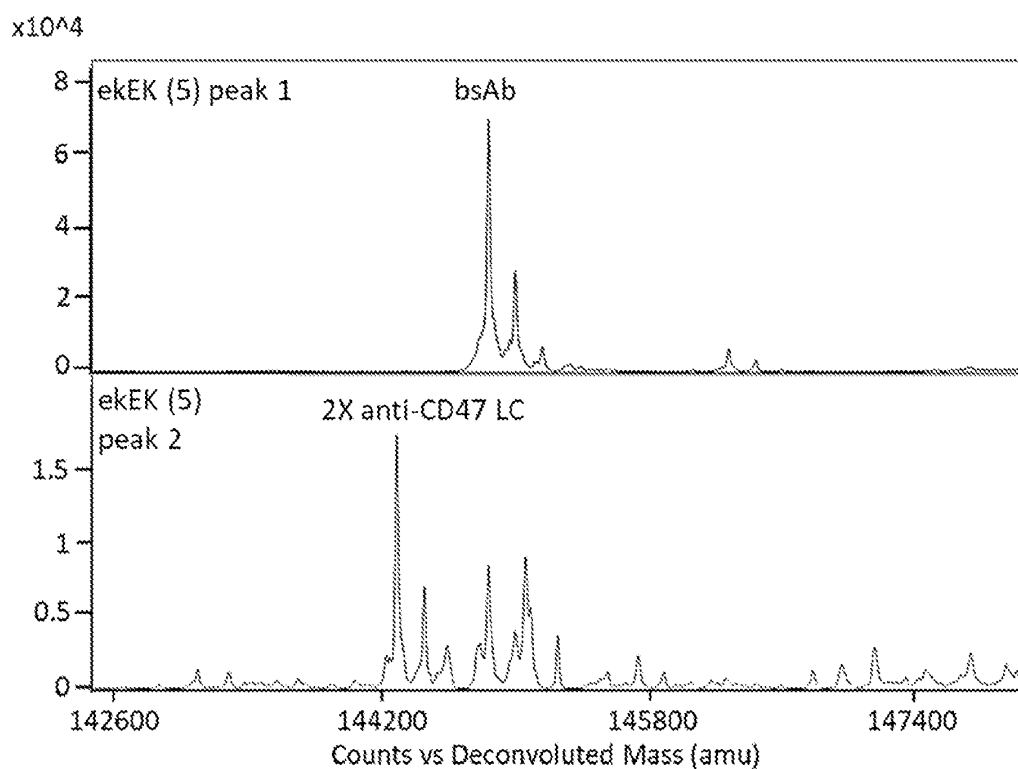
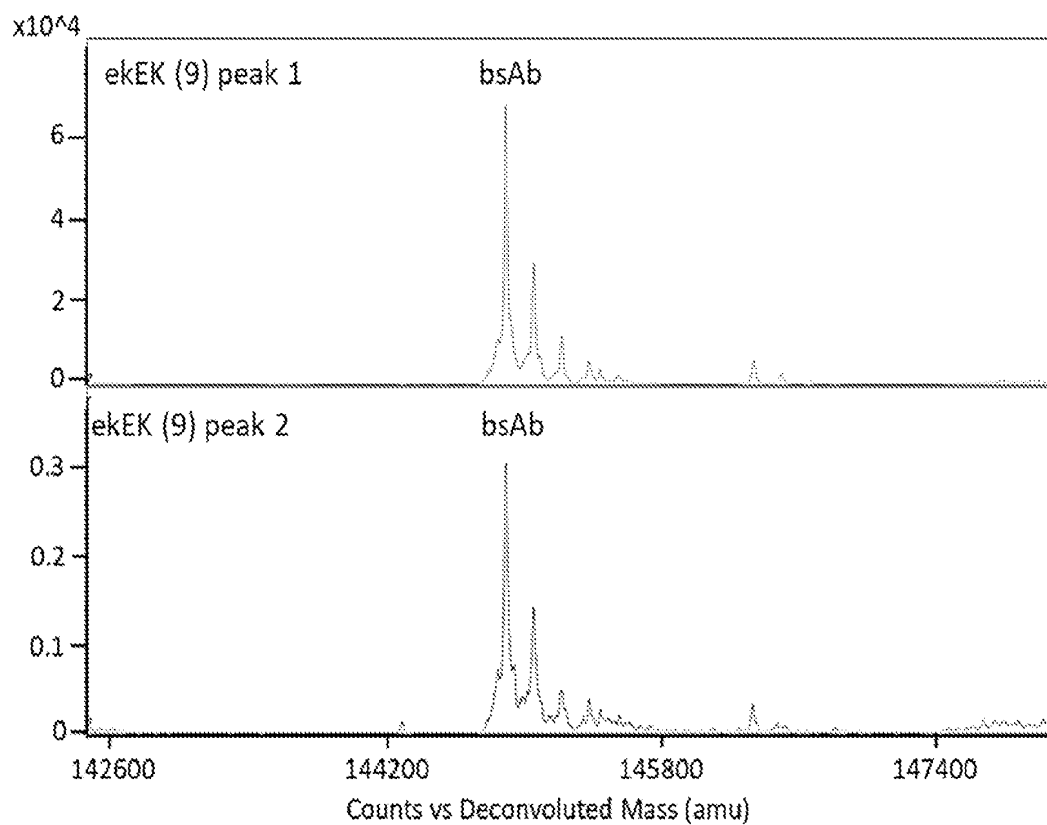


FIG. 8C

**FIG. 8D****FIG. 8E**

**FIG. 8F**

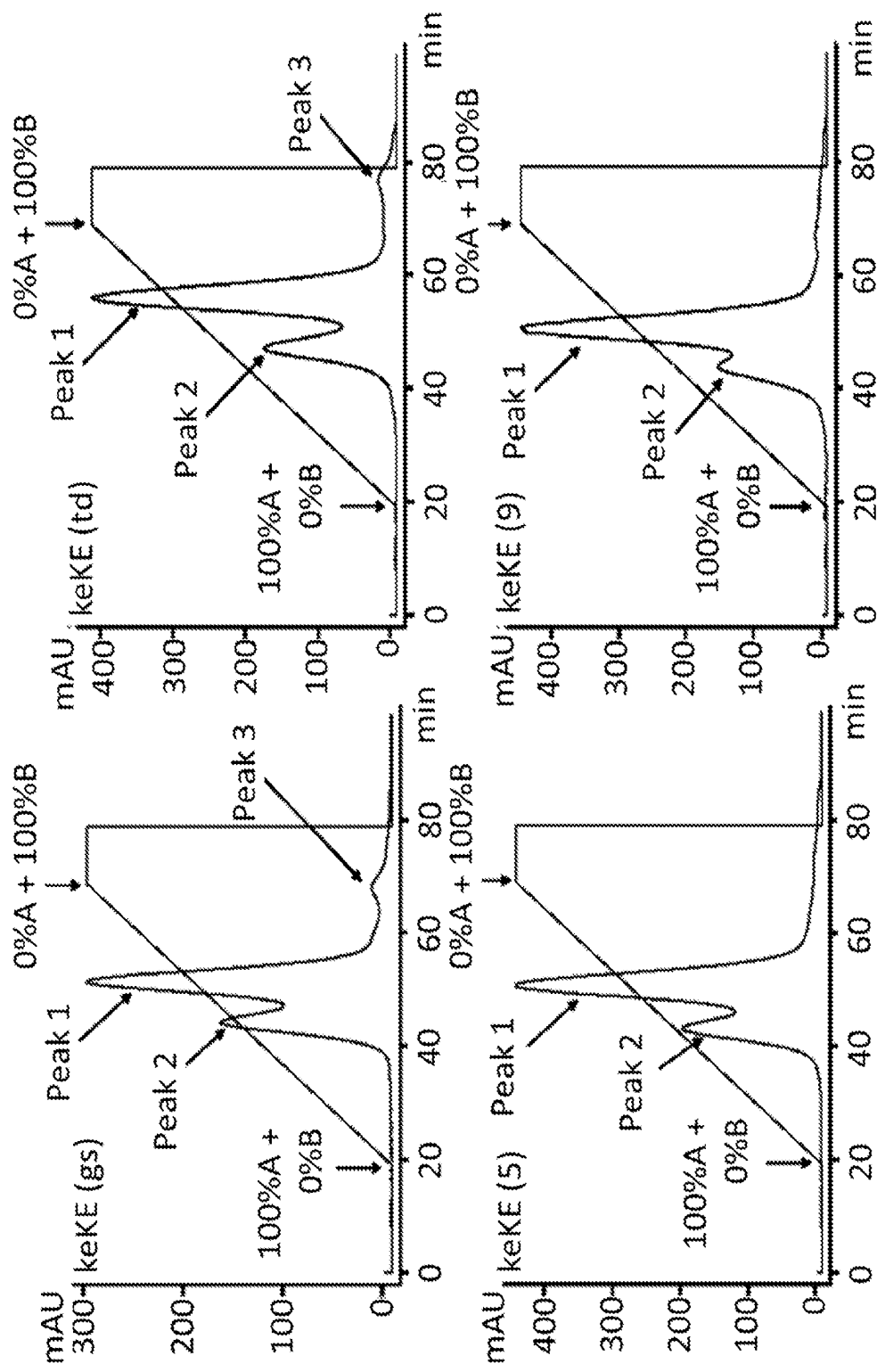


FIG. 8G

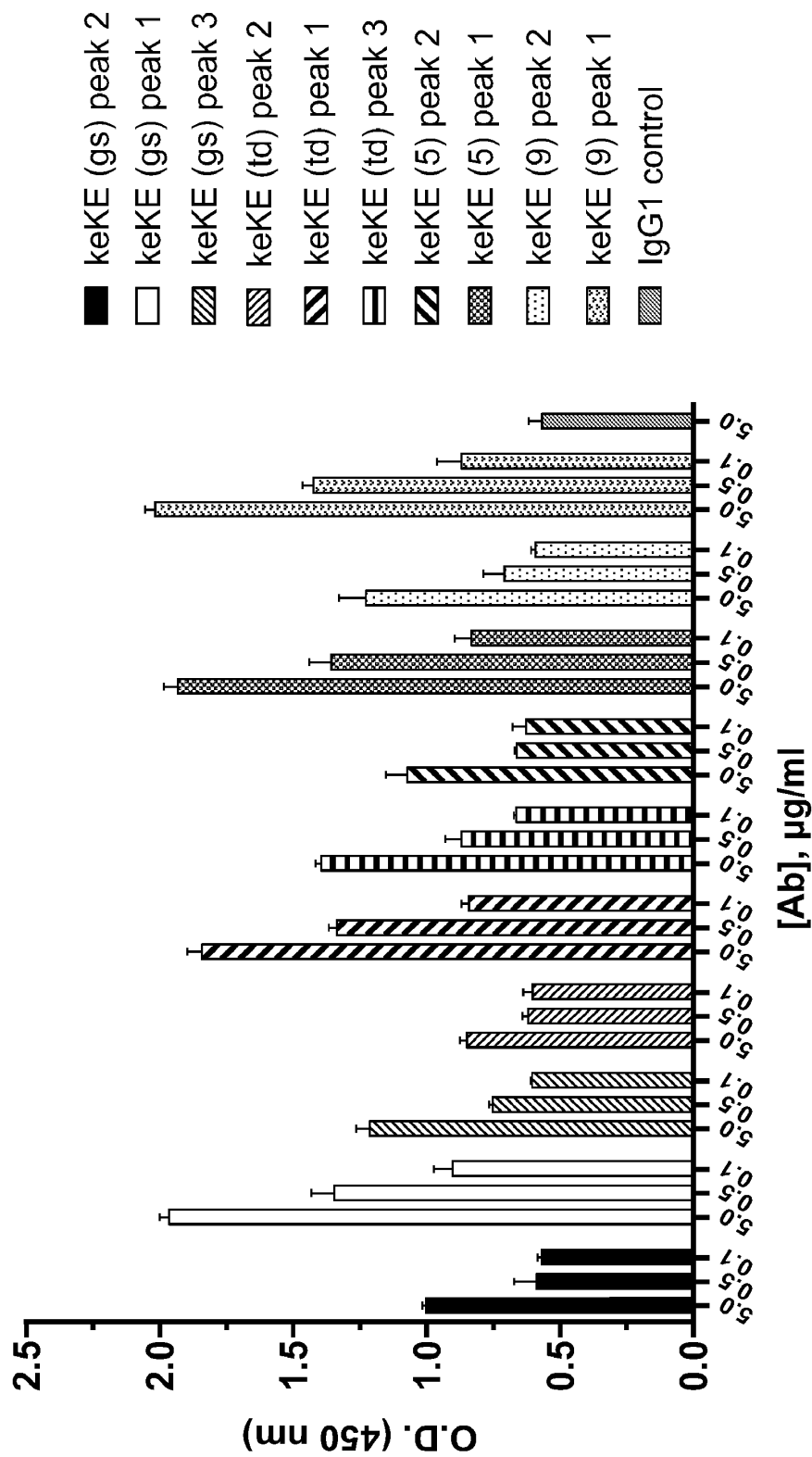
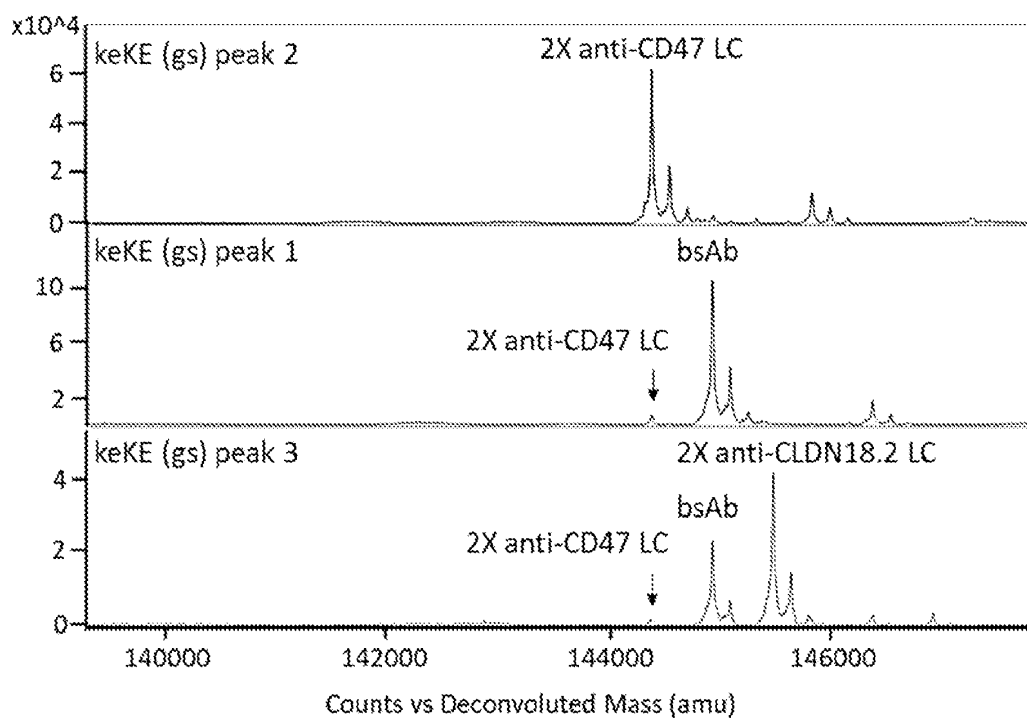
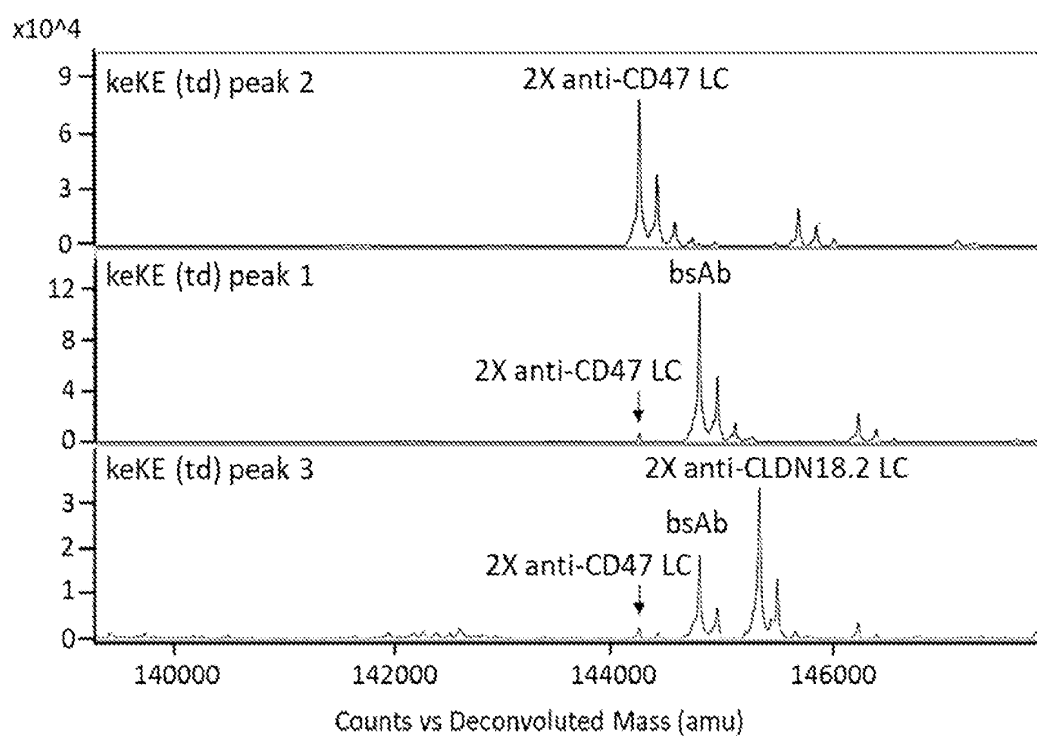
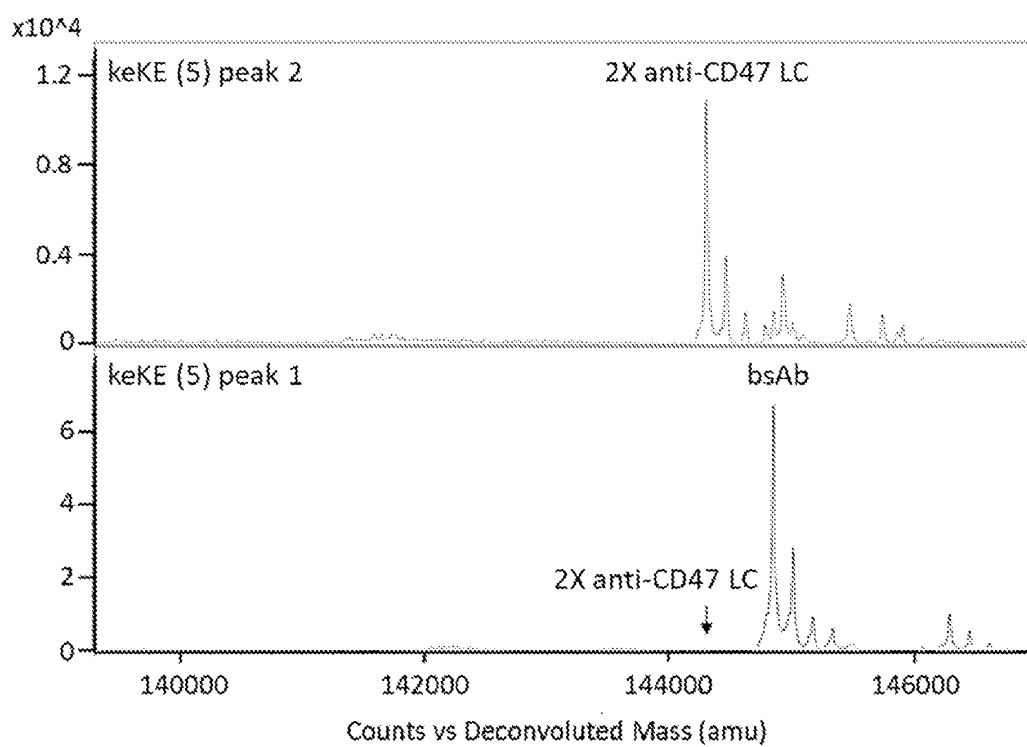
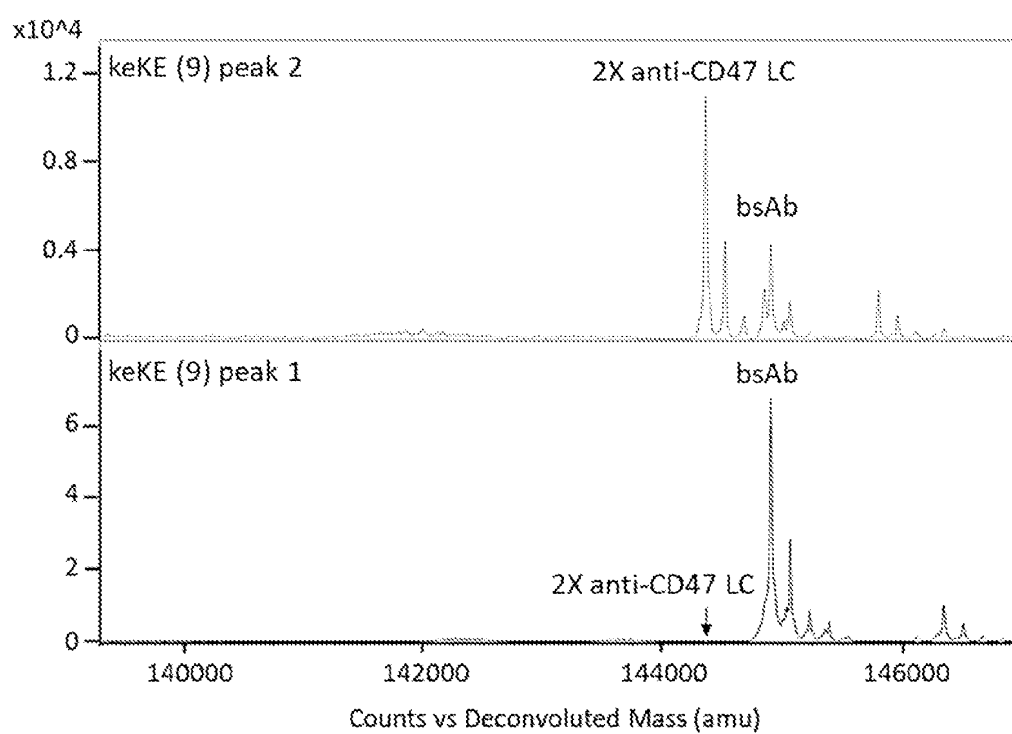


FIG. 8H

**FIG. 8I****FIG. 8J**

**FIG. 8K****FIG. 8L**

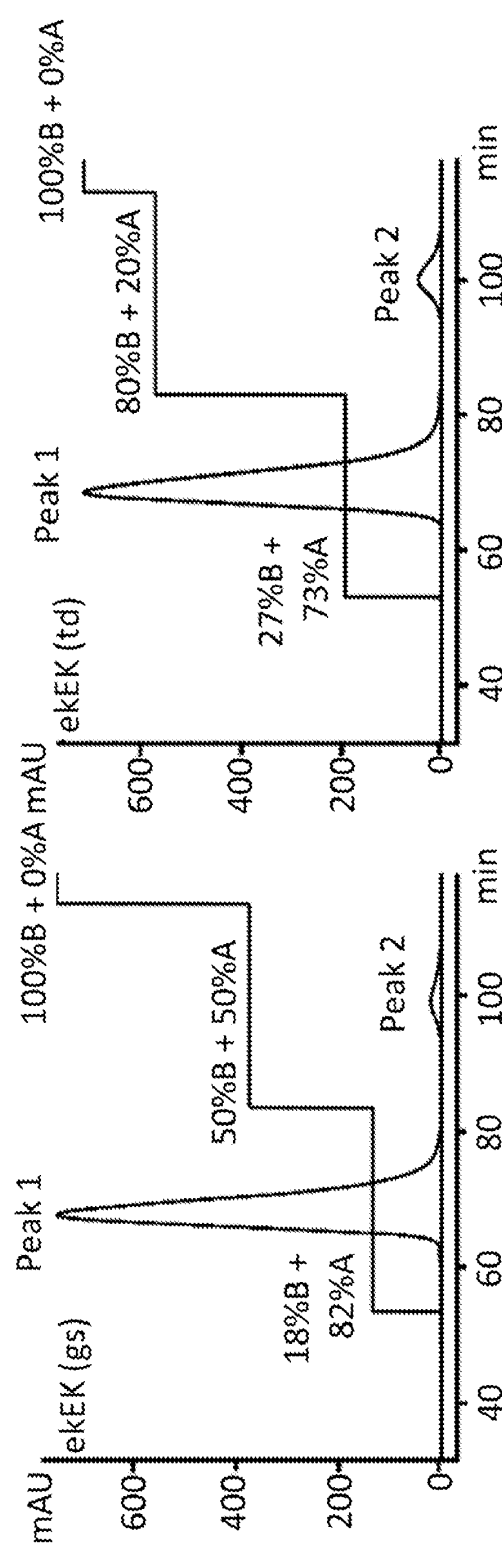


FIG. 9A

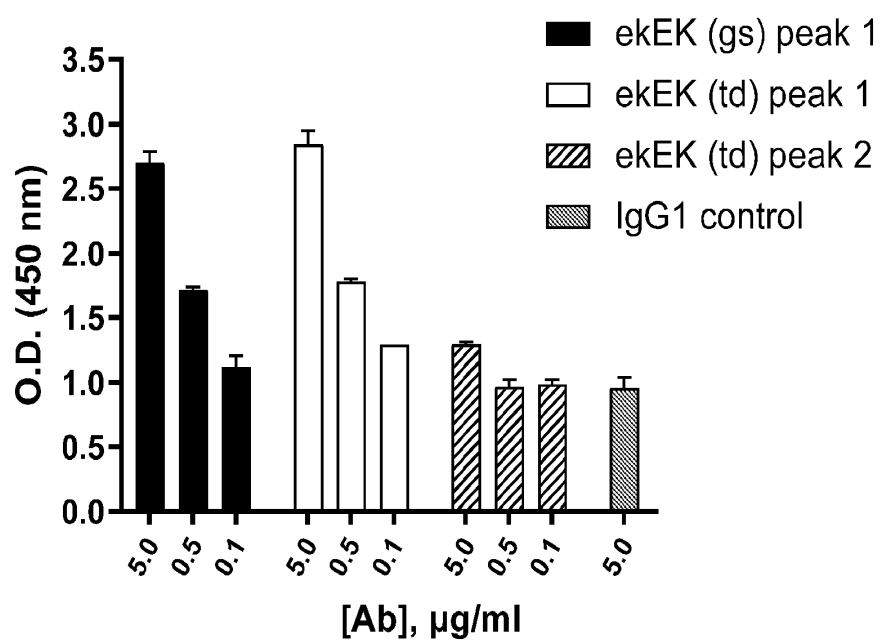


FIG. 9B

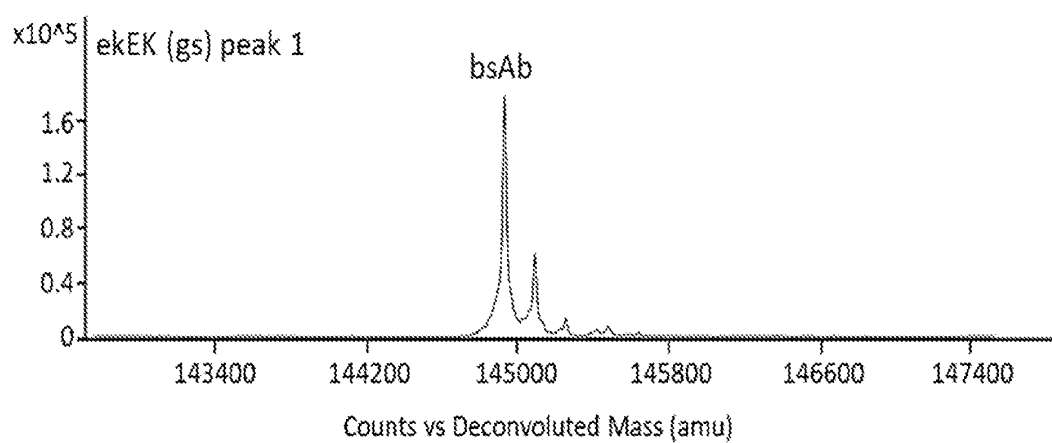
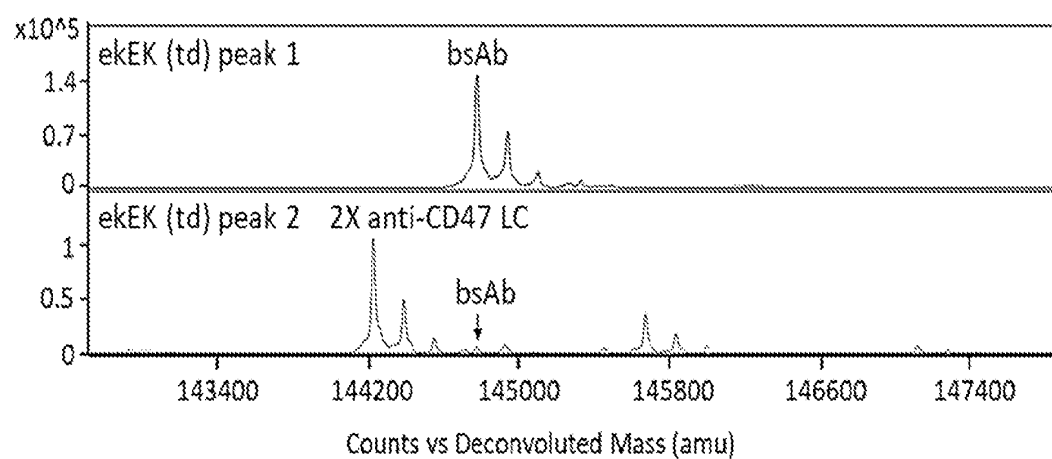


FIG. 9C

**FIG. 9D**

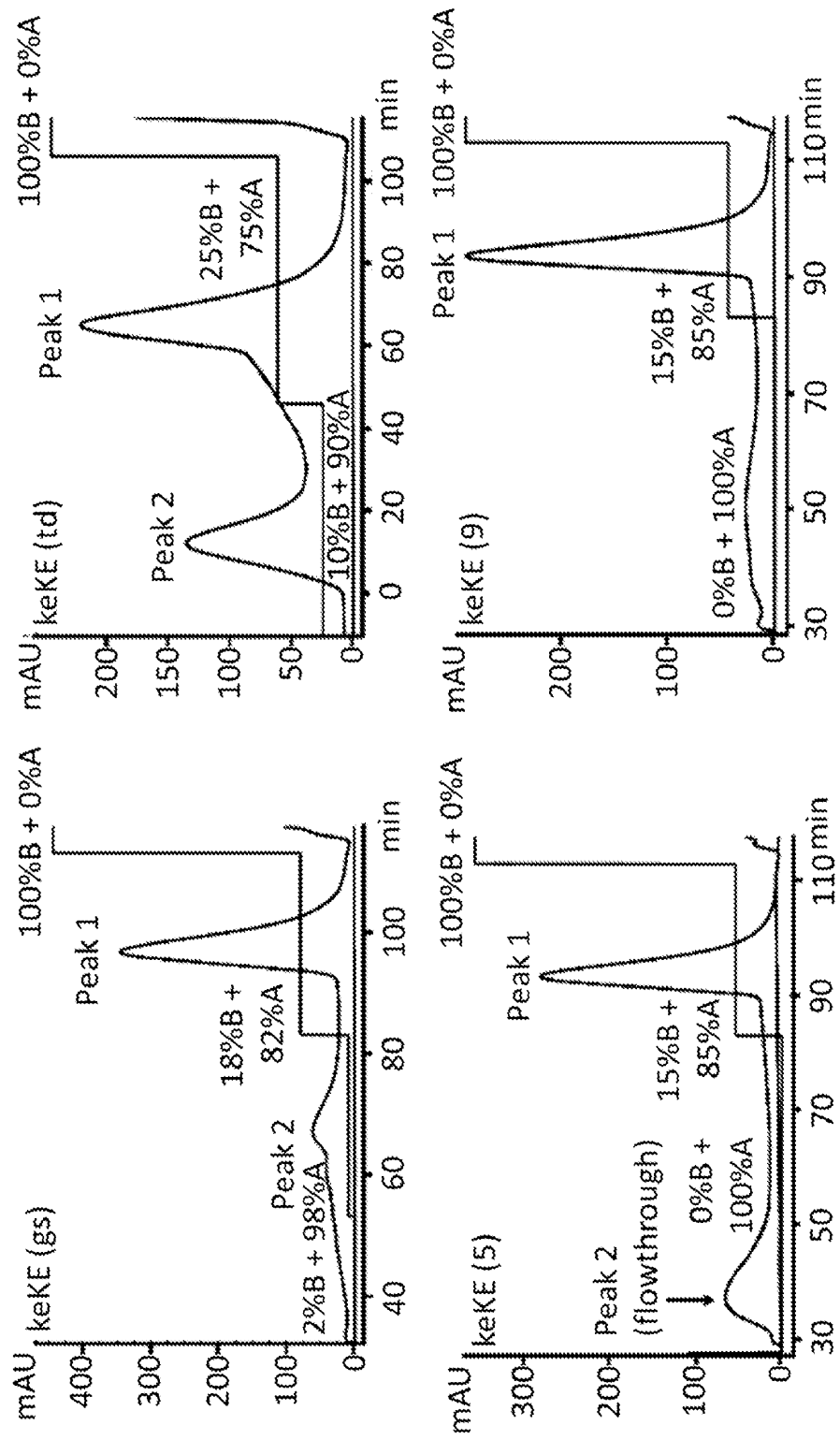


FIG. 9E

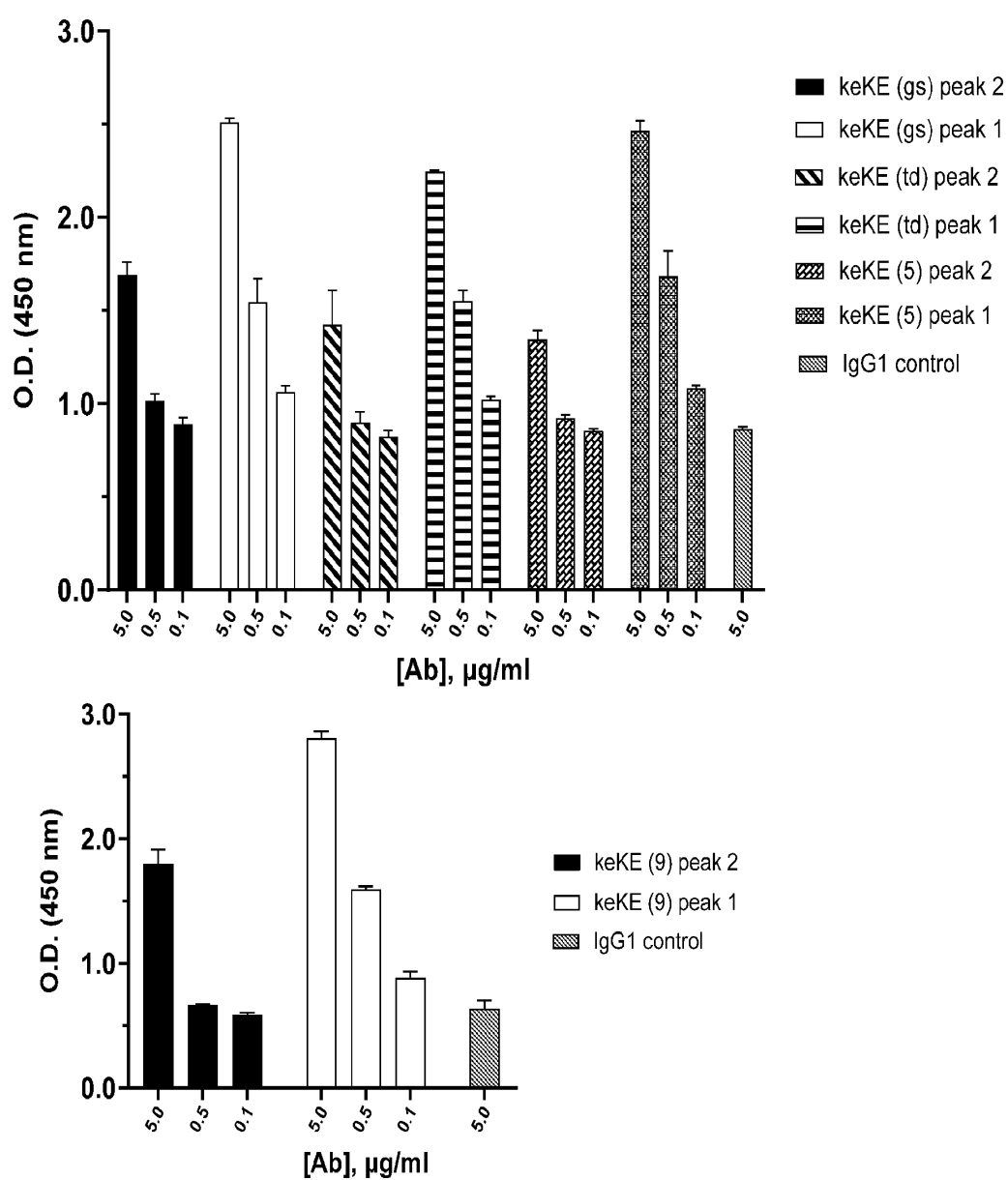


FIG. 9F

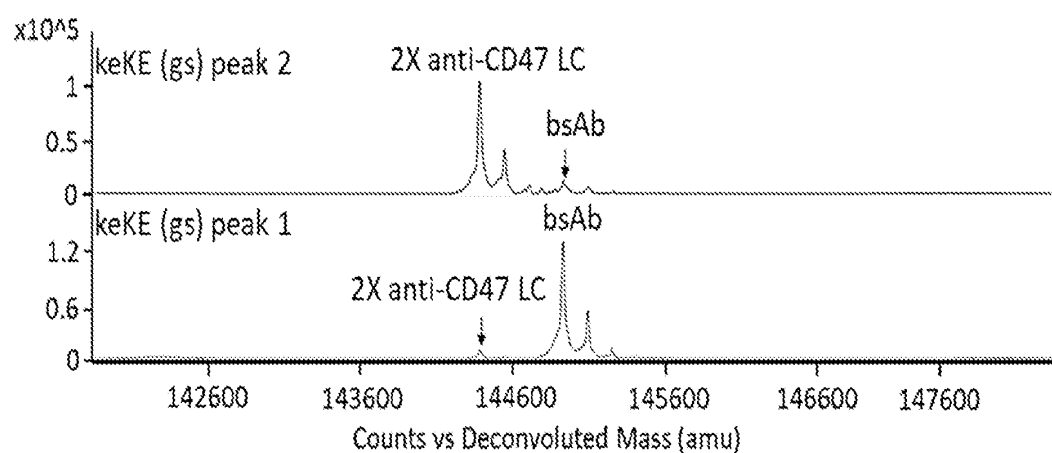


FIG. 9G

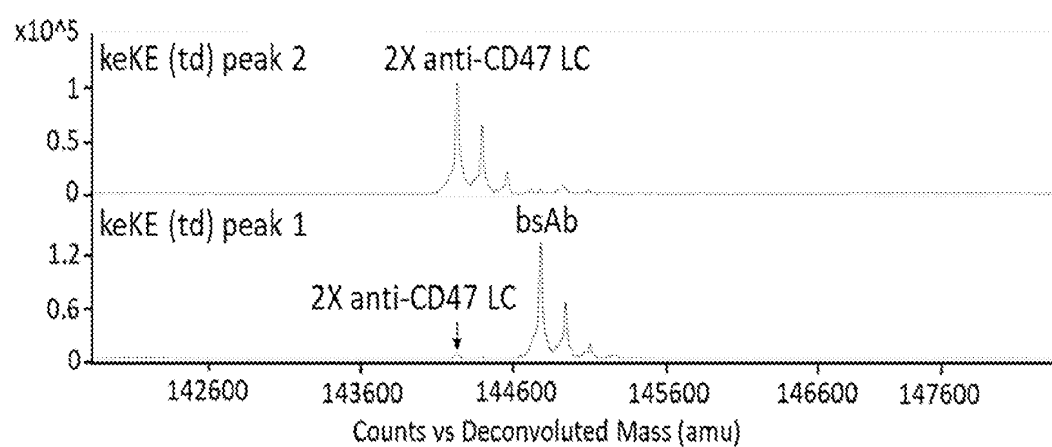
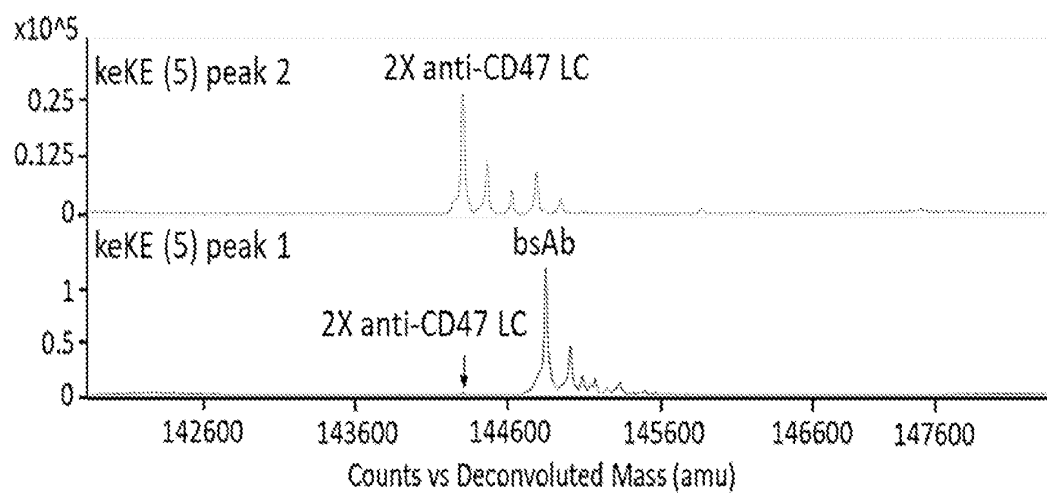
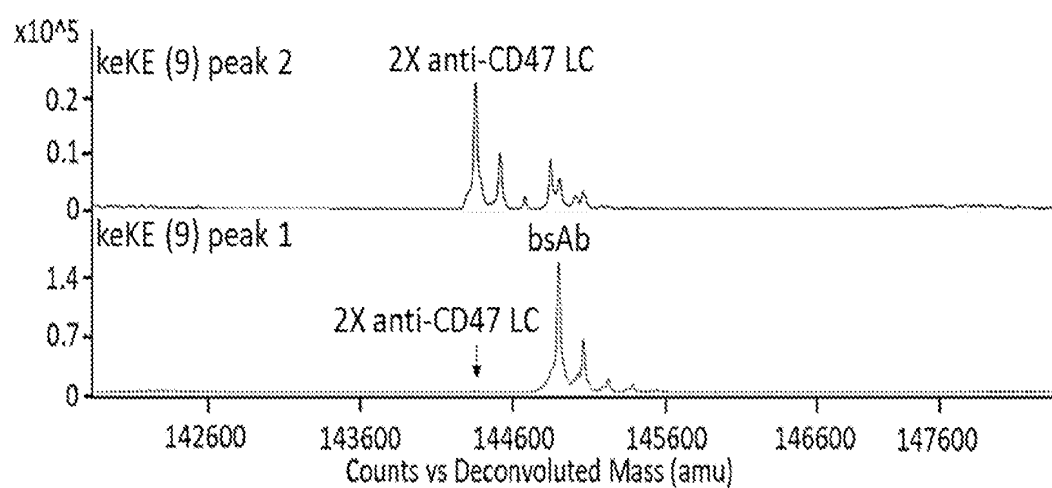


FIG. 9H

**FIG. 9I****FIG. 9J**

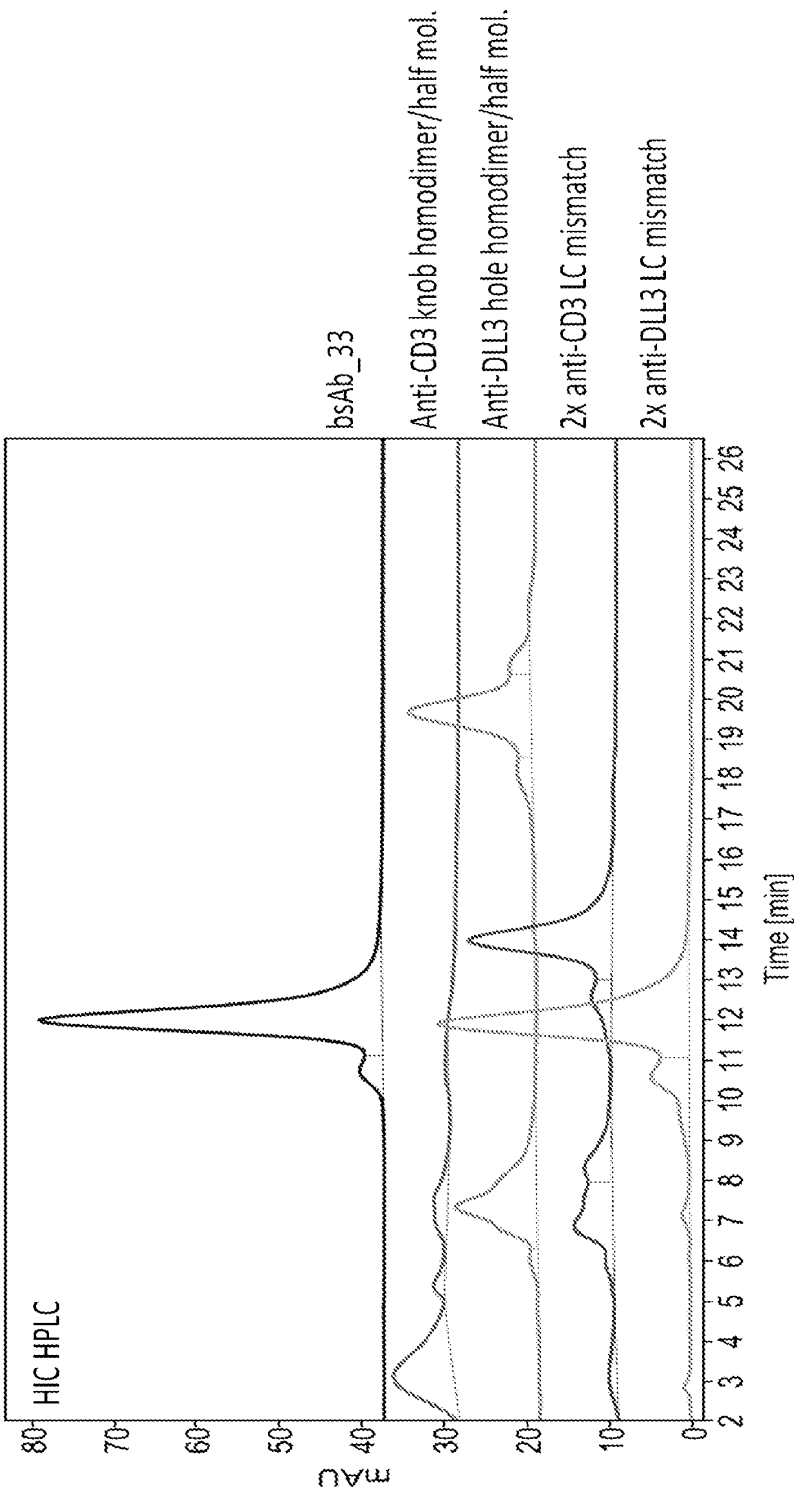


FIG. 10A

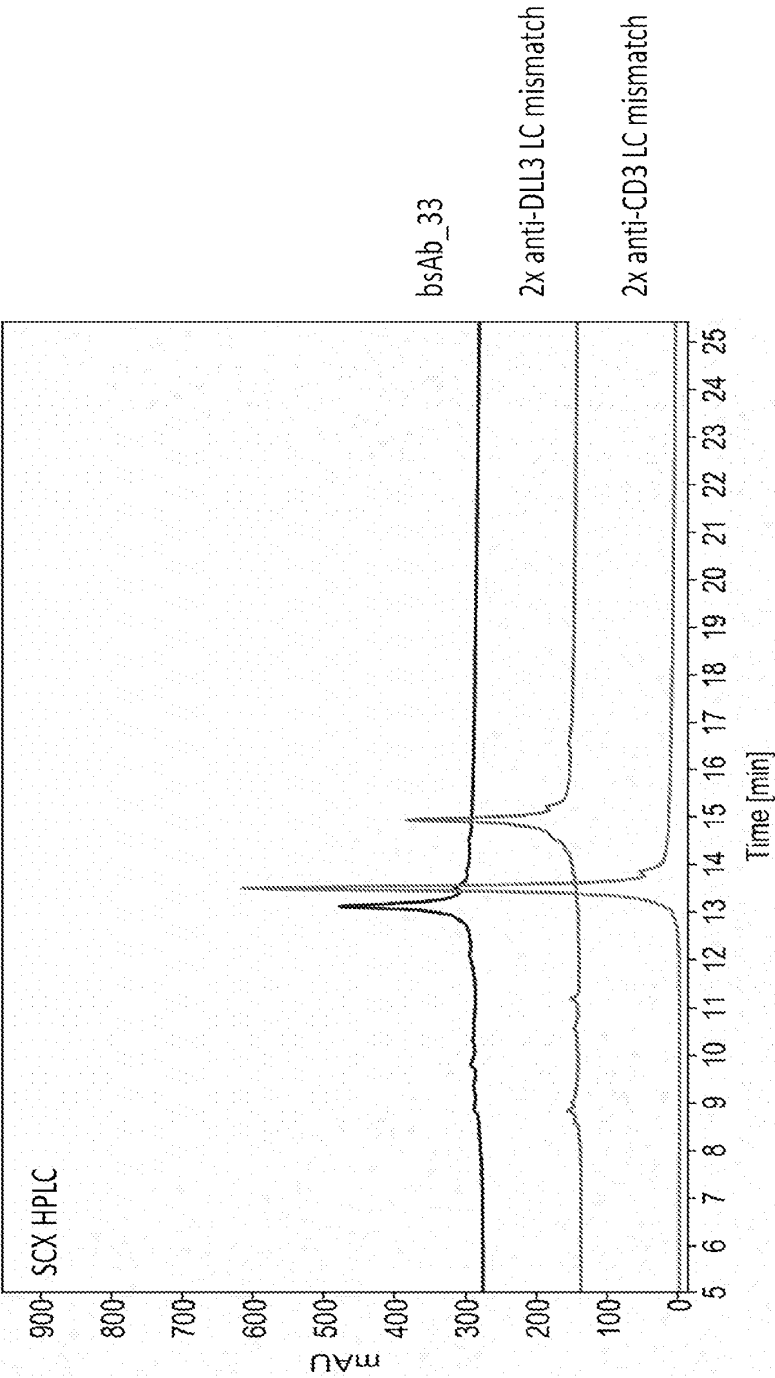


FIG. 10B

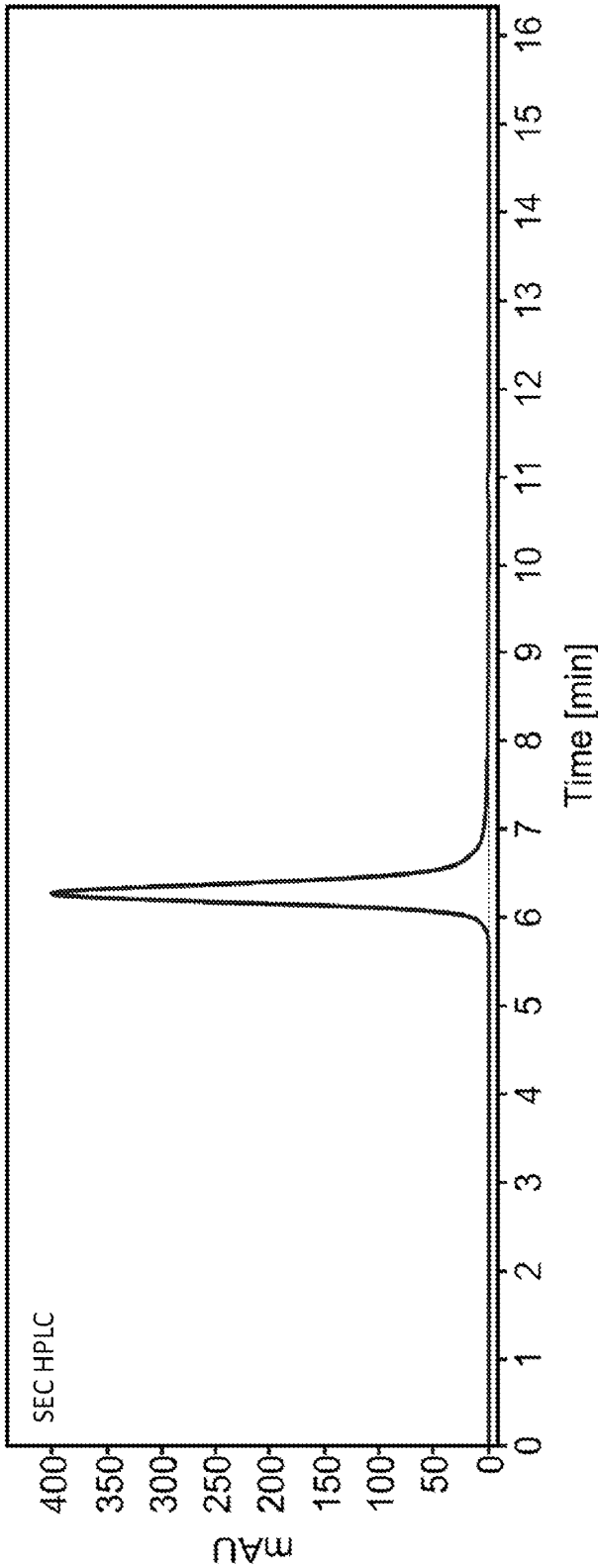


FIG. 10C

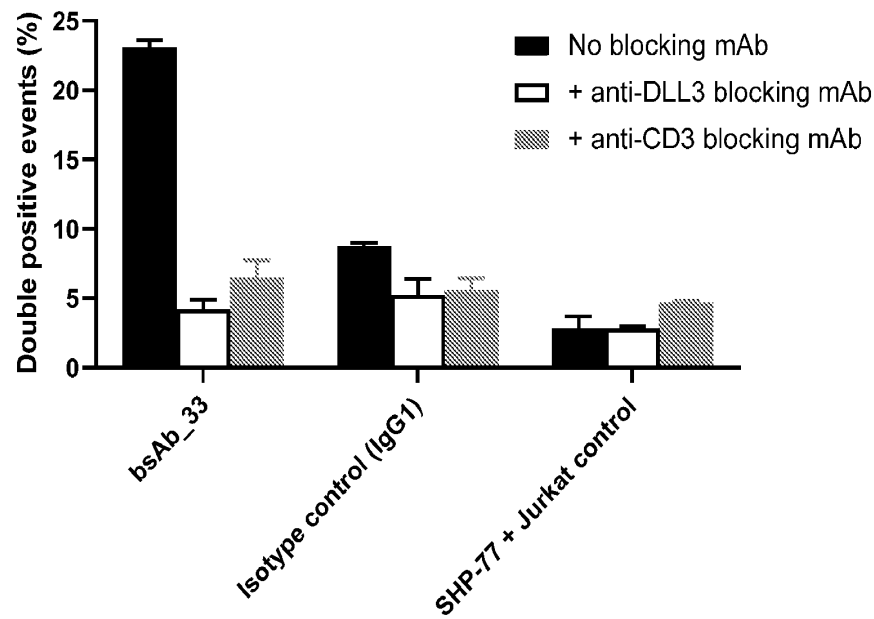


FIG. 11A

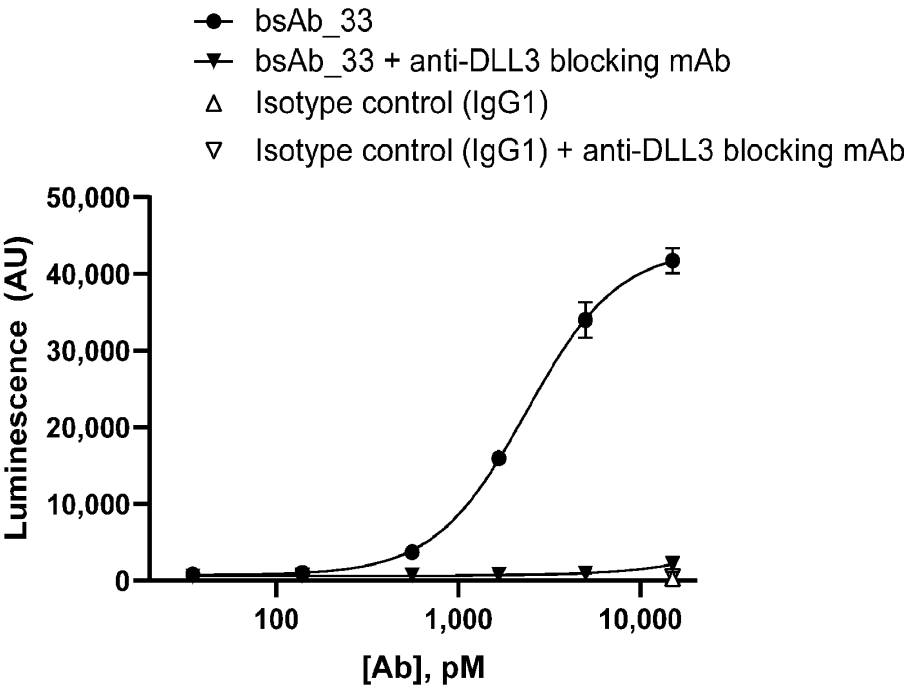


FIG. 11B