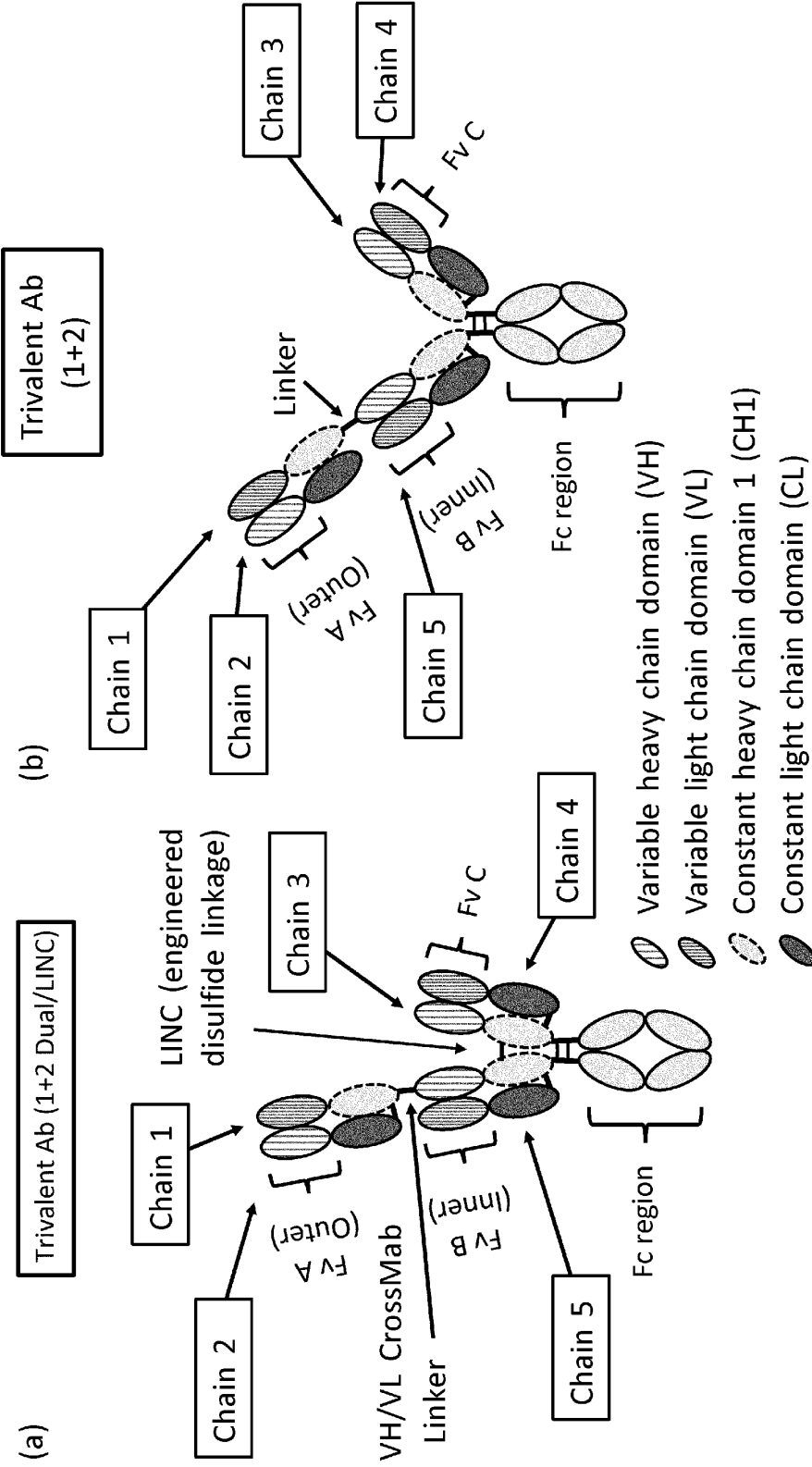
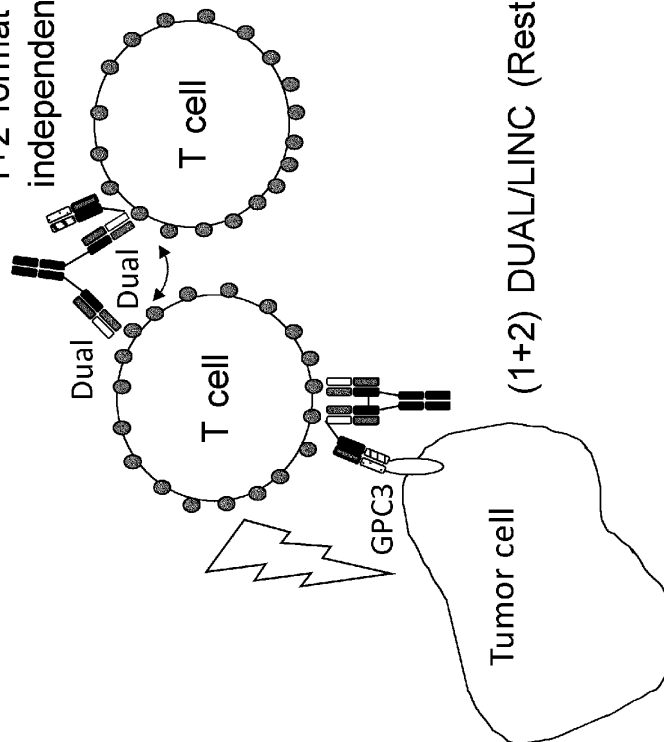


[Fig. 1]

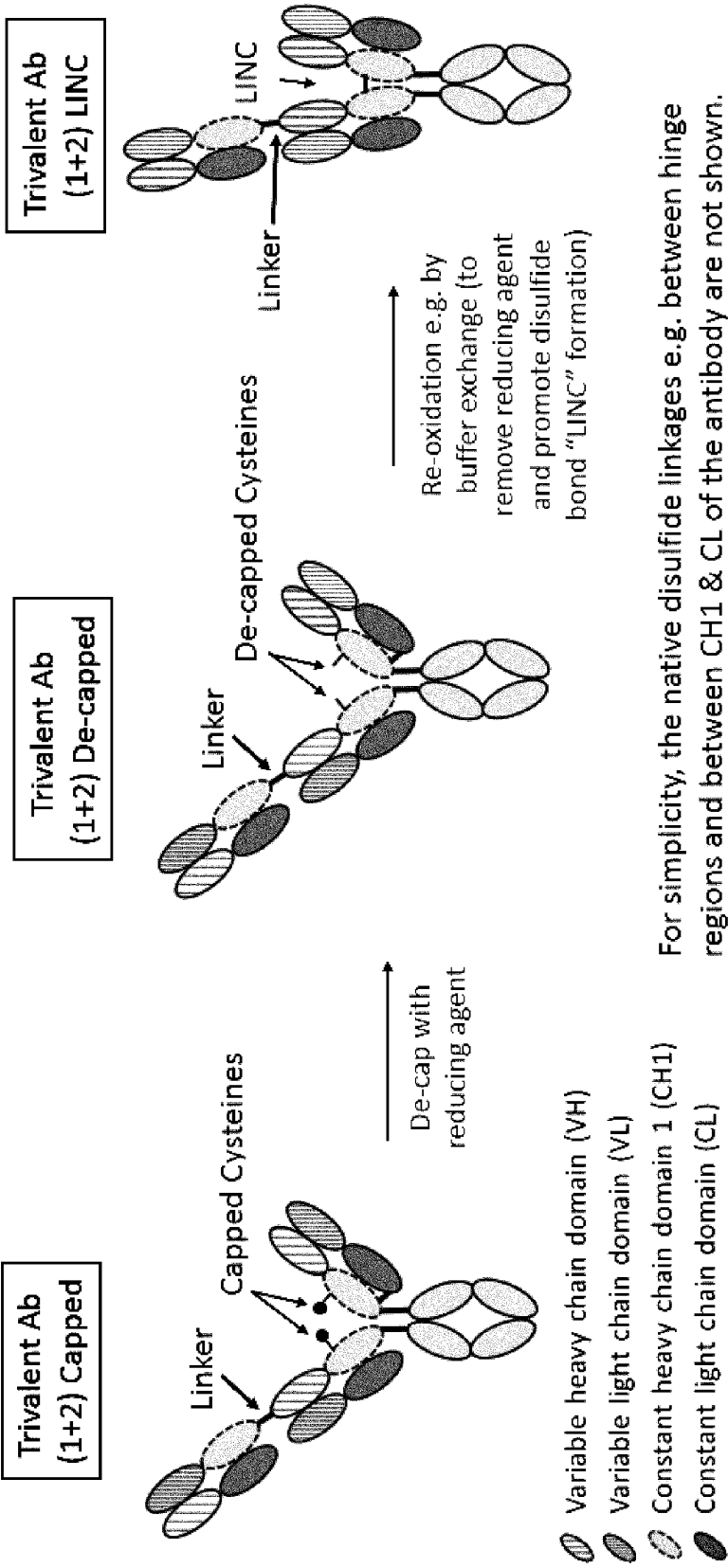


[Fig. 2a]

1+2 format (trans binding may lead to antigen independent cross-linking of immune cells)

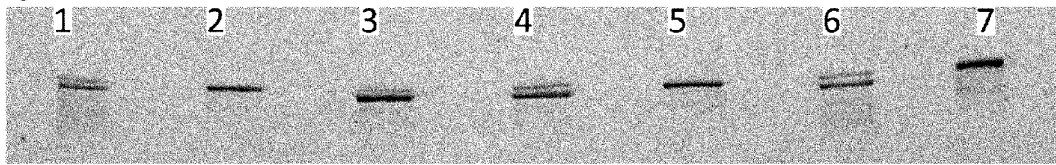


[Fig. 2b]



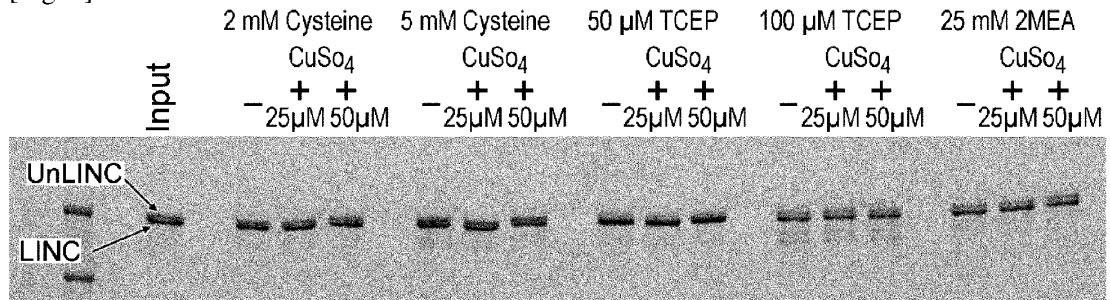
For simplicity, the native disulfide linkages e.g. between hinge regions and between CH1 & CL of the antibody are not shown.

[Fig. 3]

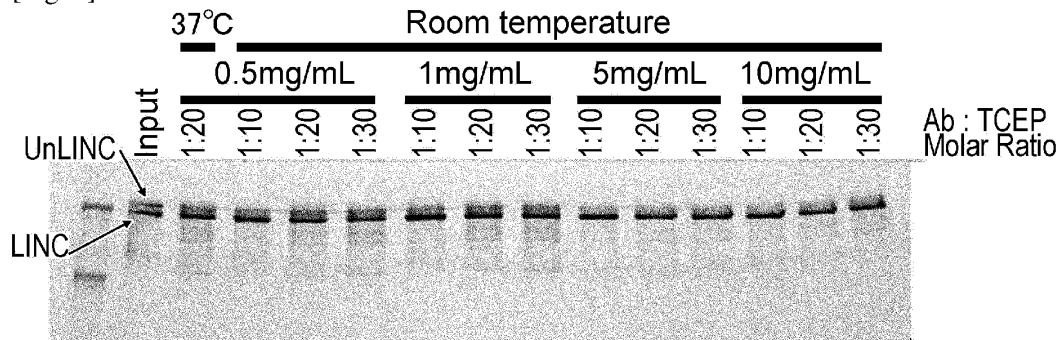


Lane	(1+2) Trivalent Ab sample name	S191C mutation	% of faster band, i.e. LINC format	% of slower band, i.e. UnLINC format
1	GPC3-DualAE05/DualAE05-FF056	Yes	63.41	36.59
2	GPC3-DualAE05/DualAE05-FF030	No	0.00	100.00
3	DLL3-DualAE05/DualAE05-FF056	Yes	78.08	21.92
4	DLL3-DualAE05/DualAE05-FF102	Yes	69.55	30.45
5	DLL3-DualAE05/DualAE05-FF117	No	0.00	100.00
6	DLL3-DualAE05/DualAE05-FF110	Yes	66.41	33.59
7	DLL3-DualAE05/DualAE05-FF115	No	0.00	100.00

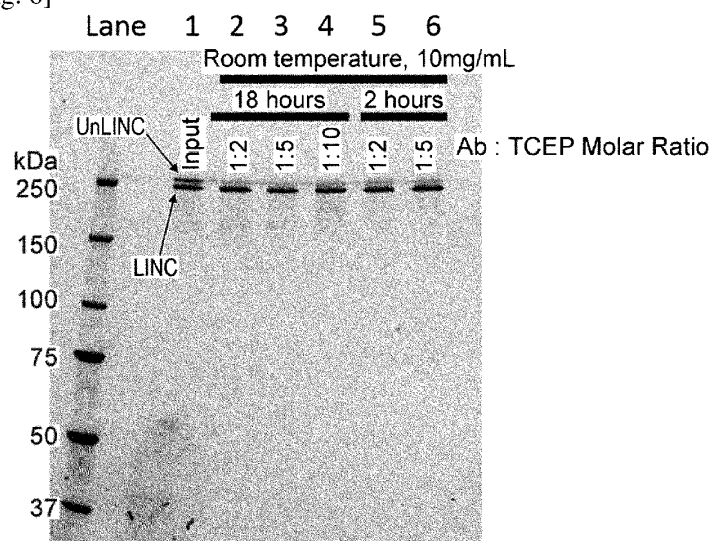
[Fig. 4]



[Fig. 5]

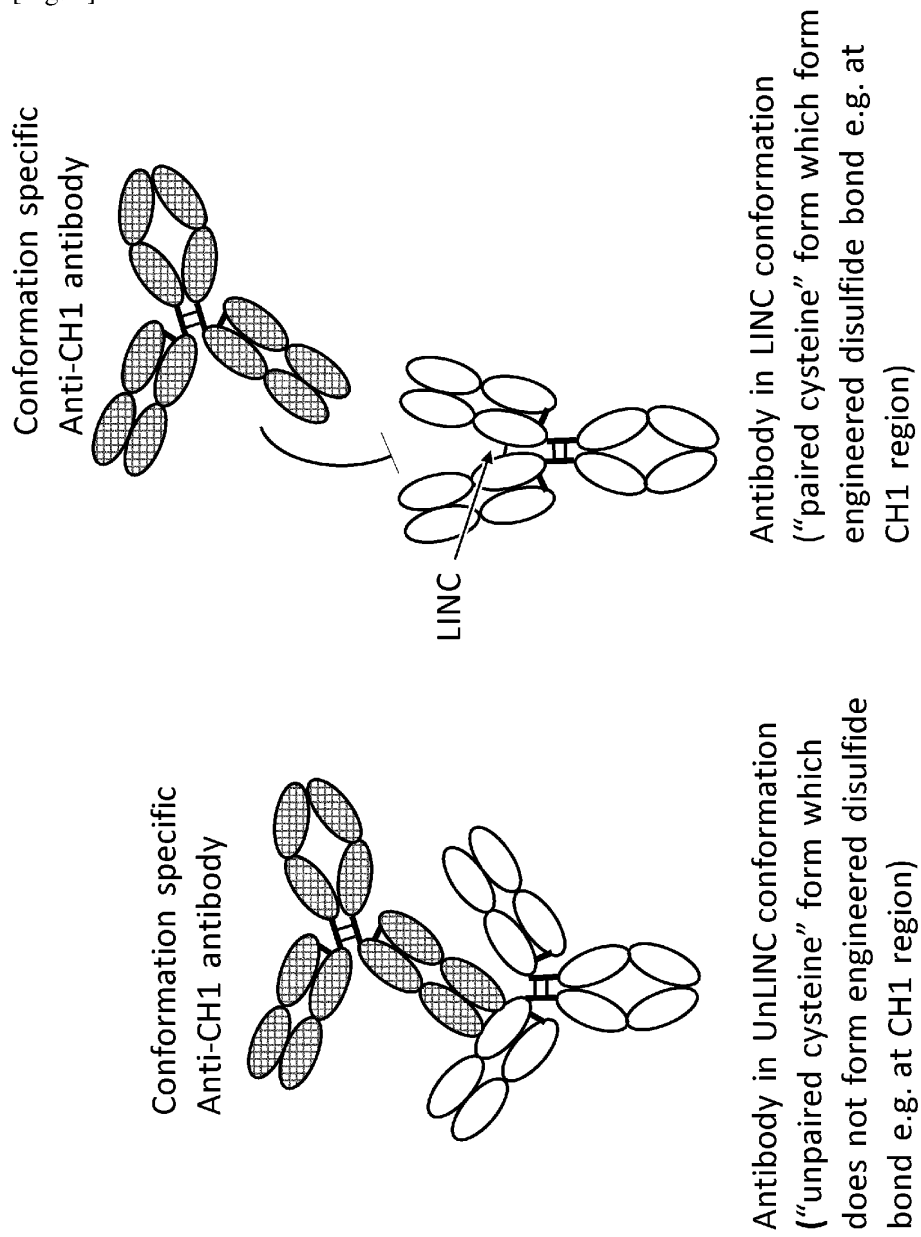


[Fig. 6]

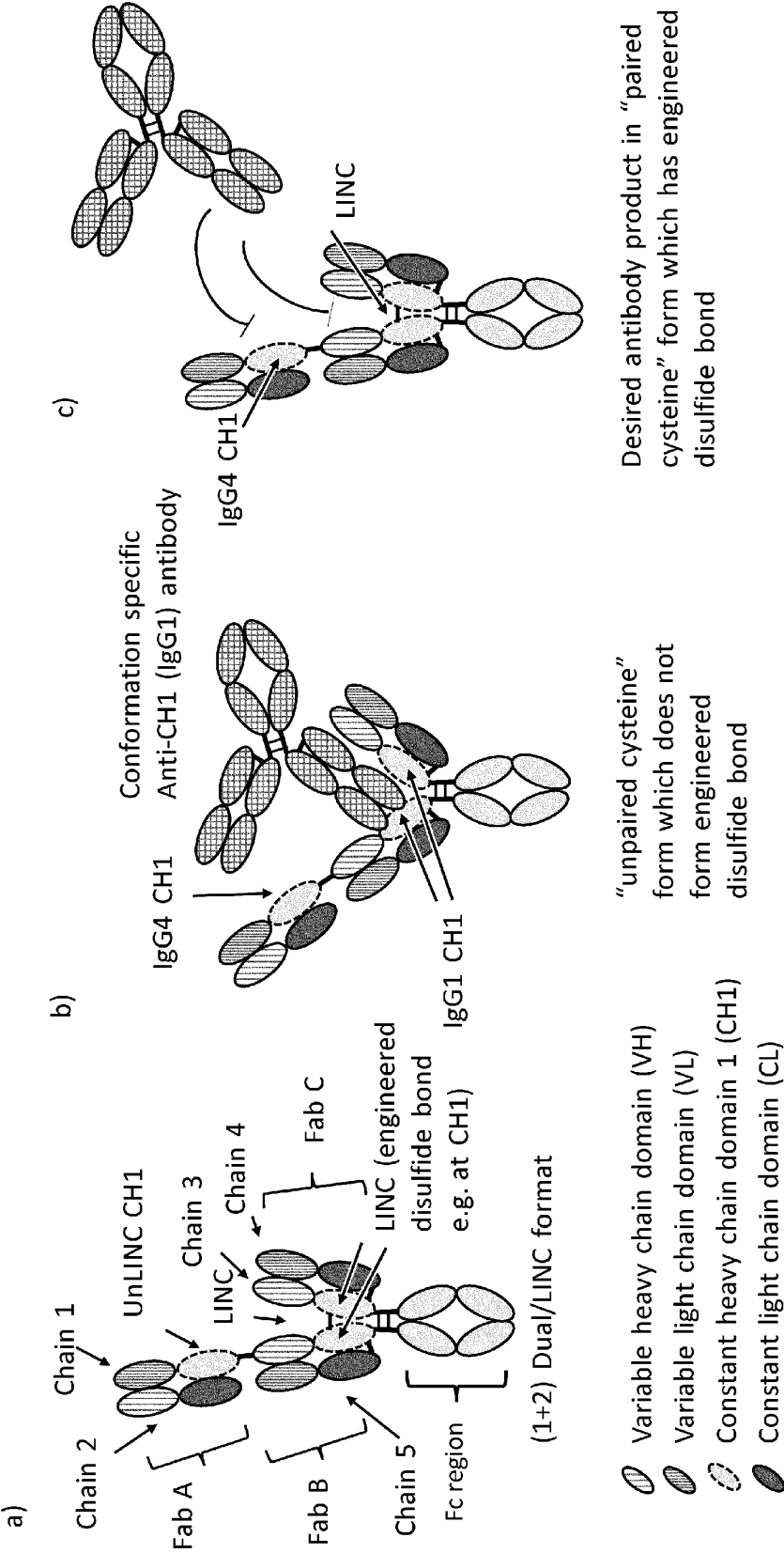


Lane	Sample	Ab: TCEP molar ratio	TCEP incubation time	% of faster band, i.e. LINC	% of slower band, i.e. UnLINC
1	DLL3-DualAE05/DualAE05-FF056	-	-	63	37
2		1:2	18 hours	90	10
3		1:5	18 hours	93	7
4		1:10	18 hours	91	9
5		1:2	2 hours	87	13
6		1:5	2 hours	89	11

[Fig. 7]



[Fig. 8]



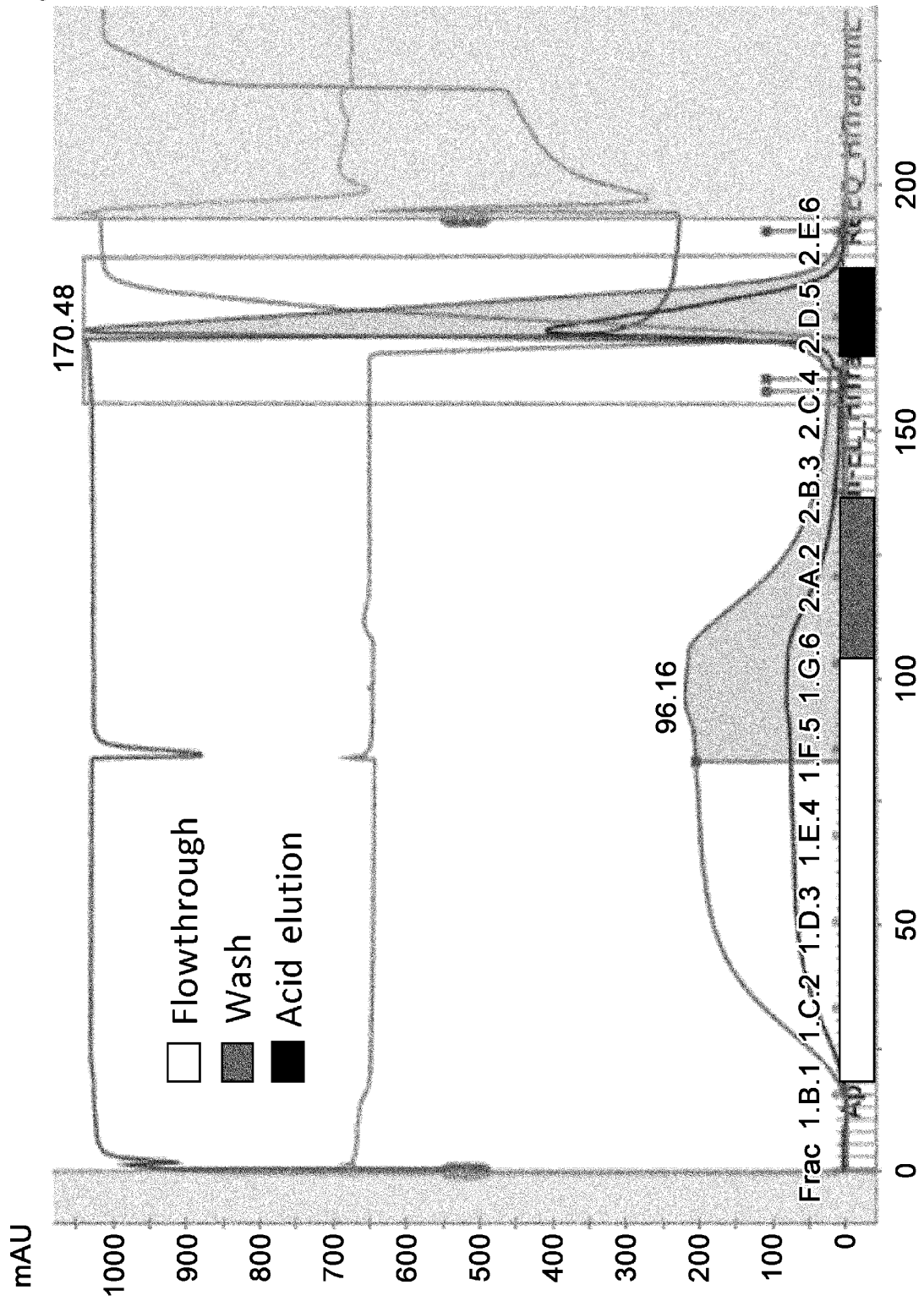
Antibody	Diagram	Legend
DualAE05-SG1201, IgG1_001, or IgG4_001		Variable heavy chain domain (VH) Variable light chain domain (VL) Constant heavy chain domain 1 (CH1) Constant light chain domain (CL)
DualAE05-SG1202		Variable heavy chain domain (VH) Variable light chain domain (VL) Constant heavy chain domain 1 (CH1) Constant light chain domain (CL)
DualAE05-SG1202k/SG1201hV11		Variable heavy chain domain (VH) Variable light chain domain (VL) Constant heavy chain domain 1 (CH1) Constant light chain domain (CL)

DualAE05-SG1201,
IgG1_001, or
IgG4_001

[Fig. 9b]

Antibody Name	SEQ ID NO.		
	Chain 1	Chain 2	Chain 3
DuaIAE05-SG1201	153	157	
DuaIAE05-SG1202	154	157	
DuaIAE05-SG1202k/SG1201hV11	155	157	156
IgG4_001	158	159	
IgG1_001	160	161	

[Fig. 10a]



[Fig. 10b]

