

Supplemental Table 3. Genetic features of potent cross-neutralizing human mAbs

mAb	Light chain gene segment		V _L mutation #		Heavy chain gene segment			AA Junction	HCDR3 Length	V _H mutation #	
	V _L	J	NT	AA	V _H	D	J _H			NT	AA
1C19	1-5	1	9	3	3-48	5-12	4	CAAGRRLTFAYW	10	22	11
1M7	3-20	1	6	3	3-66	3-10	7	CARFGPPYSQEEGVFHGAFD VW	20	15	6
1N5	2-24	2	9	5	1-69	5-24	1	CAGEMATVFEYFQYW	13	24	19