

**Table S2. Reversion of affinity maturation patches on VRC01 and MinVRC01 reveal redundancies in affinity maturation residues that contribute to neutralization.** Variants in which affinity maturation patches on VRC01 and MinVRC01 were reverted to germline were tested on a cross-clade panel of viruses. While there was loss of neutralization breadth for L-FW3 and H-CDR2 reversions for MinVRC01, neutralization activity remained for VRC01. The data also show that the disulfide is not necessary for potency and breadth of MinVRC01. Presented values are neutralization IC<sub>50</sub> in µg/ml and colored according to the listed legend.

CLADE	VIRUS	VRC01	VRC01 LACDR1	VRC01 LΔFW3	VRC01 HΔCDR1	VRC01 HΔCDR2	VRC01 HΔFW3	VRC01 HΔdisulf	Neutralization IC <sub>50</sub> (µg/mL)
A	Q769	0.02	0.06	0.05	0.03	3.13	0.06	0.029	50
	0330.v4.c3	0.11	> 50	2.51	1.43	8.50	0.29	0.076	10
	Q461.e2	0.26	> 50	> 50	50	12.9	0.77	0.239	1
B	REJO4541.67	0.05	0.11	0.08	0.07	1.35	0.07	0.036	0.10
	JR-FL	0.03	> 50	0.24	> 50	> 50	0.10	0.033	0.01
	TRO.11	0.33	> 50	> 50	50	> 50	0.78	0.300	0.001
C	ZM197M.PB7	0.10	0.19	0.12	0.19	0.30	0.14	0.059	
	6240_08_TA5_4622	0.61	> 50	1.37	0.62	10.6	1.23	0.495	
	ZM109F.PF4	0.18	> 50	4.53	2.33	2.08	0.36	0.194	
C (T/F)	Ce2010_F5	0.28	> 50	0.83	0.33	3.47	0.53	0.166	
	Ce2060_G9	0.23	> 50	> 50	> 50	> 50	0.83	0.262	
BC	CNE52	0.34	> 50	2.00	0.83	3.17	1.30	0.206	
	CNE30	0.50	> 50	> 50	2.47	> 50	1.17	0.433	
CRF02_AG	C3347.c11	0.08	0.07	0.12	3.26	> 50	0.16	0.073	
	R2184.c04	0.09	0.37	0.10	0.35	0.28	0.20	0.061	
G	X2131	0.29	20	0.69	2.40	1.99	1.43	0.467	

  

CLADE	VIRUS	MinVRC01	MinVRC01 LACDR1	MinVRC01 LΔFW3	MinVRC01 HΔCDR1	MinVRC01 HΔCDR2	MinVRC01 HΔFW3	MinVRC01 HΔdisulf
A	Q769	0.02	11.0	> 10	0.41	> 50	0.03	0.031
	0330.v4.c3	0.17	> 50	> 50	> 50	> 50	4.76	0.510
	Q461.e2	0.28	> 50	> 50	> 50	> 50	5.89	0.522
B	REJO4541.67	0.01	> 50	> 10	> 50	> 50	0.03	0.025
	JR-FL	1.19	> 50	> 50	> 50	> 50	> 50	3.030
	TRO.11	0.44	> 50	> 50	> 50	> 50	5.07	1.551
C	ZM197M.PB7	0.03	> 50	0.17	8.90	> 50	0.06	0.066
	6240_08_TA5_4622	0.92	> 50	> 50	> 50	> 50	6.96	1.061
	ZM109F.PF4	0.52	> 50	> 50	> 50	> 50	11.4	0.418
C (T/F)	Ce2010_F5	0.15	> 50	> 50	1.81	> 50	1.39	0.153
	Ce2060_G9	0.09	> 50	> 50	> 50	> 50	0.69	0.153
BC	CNE52	0.09	> 50	> 10	0.72	> 50	1.79	0.141
	CNE30	0.40	> 50	> 50	> 50	> 50	4.92	0.936
CRF02_AG	C3347.c11	0.02	0.02	0.17	0.25	> 50	0.03	0.055
	R2184.c04	0.04	> 50	> 10	3.09	> 50	0.14	0.071
G	X2131	0.18	> 50	> 50	> 50	> 50	2.44	0.145