

**Table S2.** Antibody escape mutants with highest ACE2 binding affinities.

<b>Antibody</b>	<b>% Change in Binding Affinity for Optimal Escape Mutant</b>
S309	120%
EY6A	155%
S2A4	120%
S304	155%
S2M11	117%
B38	200%
C105	178%
P2B-2F6	141%
CB6	151%
CC12.1	200%
CC12.3	200%
CV30	200%
Fab 2-4	178%
COVA2-04	200%
COVA2-39	200%
S2H13	200%
S2H14	200%
S2E12	123%
REGN10933	178%
REGN10987	200%