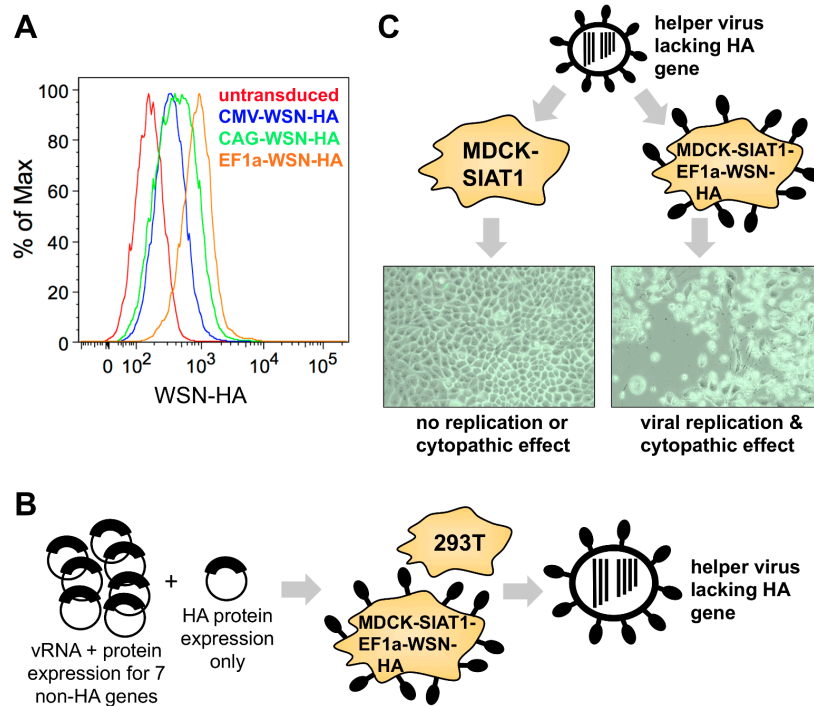
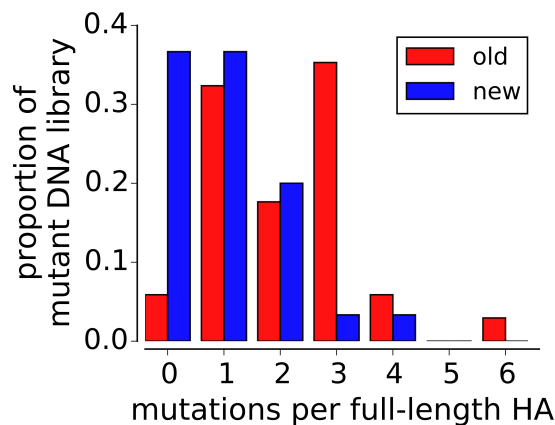


# Supplementary Materials: Accurate Measurement of the Effects of All Amino-Acid Mutations on Influenza Hemagglutinin

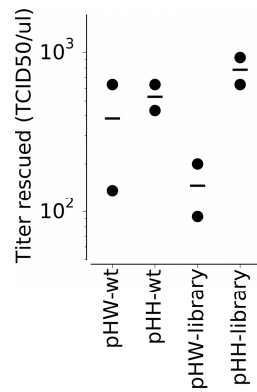
Michael B. Doud and Jesse D. Bloom



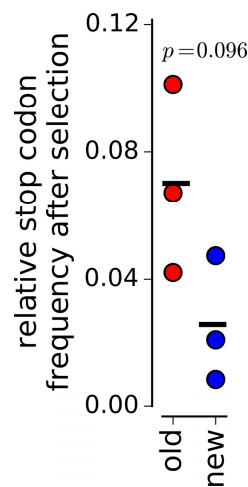
**Figure S1.** An HA-deficient helper virus can replicate in cells constitutively expressing HA protein.



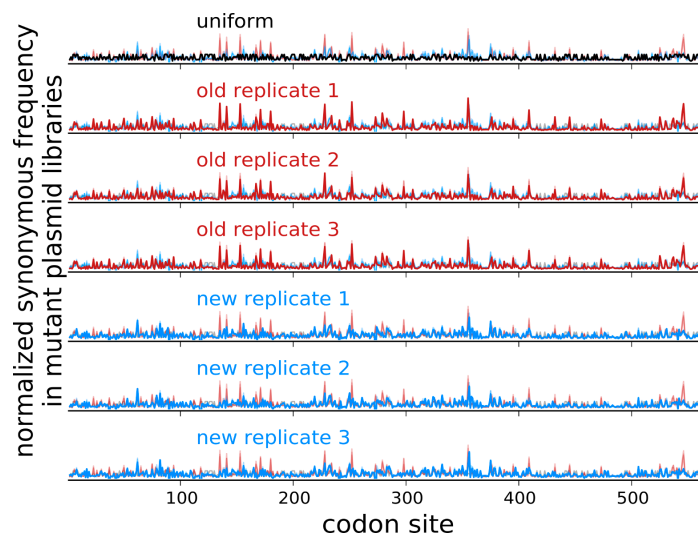
**Figure S2.** The mutant plasmid DNA library used in this study (“new”) has a lower mutation rate than the library used by [4] (“old”).



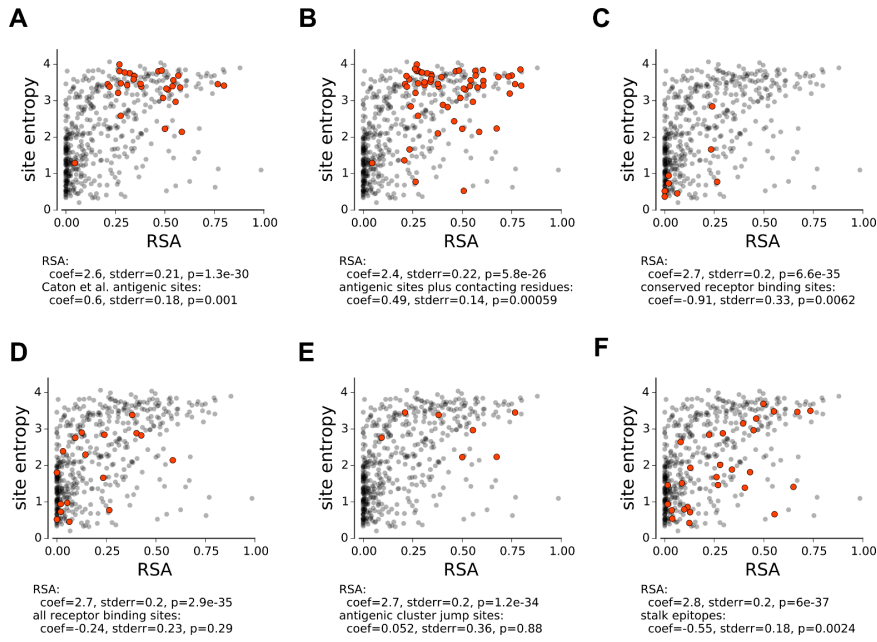
**Figure S3.** Mutant virus library generation is more efficient when HA is encoded on the pHH21 plasmid.



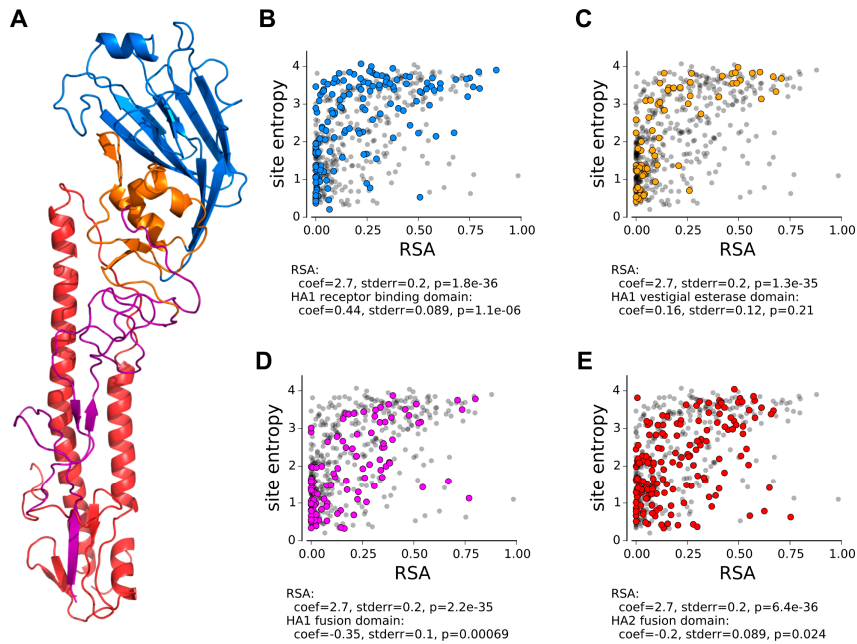
**Figure S4.** Purging of stop codons is more complete in our new experiment than in the previous one.



**Figure S5.** Synonymous frequency peaks observed in bottlenecked virus libraries are not due to the composition of plasmid mutant libraries.



**Figure S6.** Statistical analyses of whether sets of sites have higher or lower mutational tolerance than expected given their solvent accessibility.



**Figure S7.** The mutational tolerance of HA's domains.



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