



Figure S2. Comparison of viral infectivity pseudotyped with BA.1 S and BA.2 S

HOS-ACE2-TMPRSS2 cells were infected with pseudoviruses bearing each S protein. The amount of input virus was normalized based on the amount of HIV-1 p24 capsid protein. The percent infectivity compared to that of the virus pseudotyped with BA.1 S is shown. Assays were performed in triplicate. The presented data are expressed as the average \pm SD. Each dot indicates the result of an individual replicate. Statistically significant difference (*, $P < 0.05$) versus parental BA.1 was determined by a two-sided Student's *t* test. The red number indicates the fold change between BA.1 and BA.2.