

Table S1. Partial homopurine and homopyrimidine sites within the SARS-CoV-2 genome.

Sequence Type (base #)	Target Sequence (5'-3')	Sequence Length	Number of Mutations
Homopurine (1801-1821)	AAAAGGAAAAG <u>CT</u> AAAAAAGG	19 nt	2
Homopurine (13610-13625)	AAGAAAAGGAC <u>G</u> GAAGA	15 nt	1
Homopurine (29378-29399)	AAAAAGGAC <u>A</u> AAAAAGAAGAAGG	21 nt	1
Homopyrimidine (21725-21742)	TTCTT <u>A</u> CCTTTCTTTTCC	17 nt	1
Homopyrimidine (25700-25717)	CCCCTTTTCTCT <u>A</u> TCTTT	17 nt	1
Homopyrimidine (26296-26313)	CTTCTTTTCTT <u>G</u> CTTTC	17 nt	1