| Mutation    | Mutation type | Gene      | Substitution | Frequency in S1-mut-<br>enriched clades |  |
|-------------|---------------|-----------|--------------|---|--|
| C3037T      | syn           | ORF1a     |              | 1                                       | Also arise independently   |
| C14408T     | nonsyn        | RdRp      | P323S        | 1                                       | Occur in in a subclade the of 19A  |
| C241T       | syn           | noncoding |              | 1                                       | ancestor of all "20" clades  |
| A23403G     | nonsyn        | S1        | D614G        | 1                                       | Siddes   |
| 11288-11296 | deletion      | ORF1a     |              | 0.96707819                              | Nsp6 deletion. Occurs several times independently in mid-late 2020   |
| A23063T     | nonsyn        | S1        | N501Y        | 0.93415638                              | N501Y and Nsp6 deletion have similar distribution, but both also occur independently                                     |
| G28881A     | nonsyn        | N         | R203K        | 0.739369                                | Basal to 20B (so this mut is present   |
| G28883C     | nonsyn        | N         | G204R        | 0.739369                                | in 501Y.V1 and 501Y.V2, but not 501Y.V3)   |
| G28882A     | syn           | N         |              | 0.739369                                | Occurs independently in 501Y.V1 and a small subclade of  |
| 21992-21994 | deletion      | S1        |              | 0.69547325                              | 20A which has 7 S1 muts and high logistic growth (~9). Also has Nsp6 del but not 501Y                                    |
| 21765-21771 | deletion      | S1        |              | 0.69410151                              | Same as above, but with the addition of a second independent occurrence in a 20A subclade (3-4 S1 mutations)             |
| A28271-     | deletion      | noncoding |              | 0.68861454                              | Between ORF8 and N. In 501.V1 and lots of small clades with at least 3 S1 mutations                                      |
| C23604A     | nonsyn        | S1        | P681H        | 0.67215364                              | Common S1 mutation. Could have functional effect. Seems to come later (not one of the muts preceding a storm of S1 muts) |