DATA sheet

Look at the emergence of variants using NextStrain.org & identify mutations

1. Record information from the sample you chose to investigate, in part B, in the table below.

|  |  |
| --- | --- |
| Sample ID\* |  |
| Collection date |  |
| S1 mutations |  |
| Country |  |
| Host |  |
| Clade |  |
| Emerging Lineage |  |
| GenBank accession |  |

\*This is the number at the top of the sample information. It is found above the collection date.

Note: If you choose to enter the data in the table above via copying it and pasting from the pop-up screen, the text might appear white. Change the font color of the pasted text to black and it will become visible again.

1. Enter the Genbank accession number for the spike protein sequence from your sample.