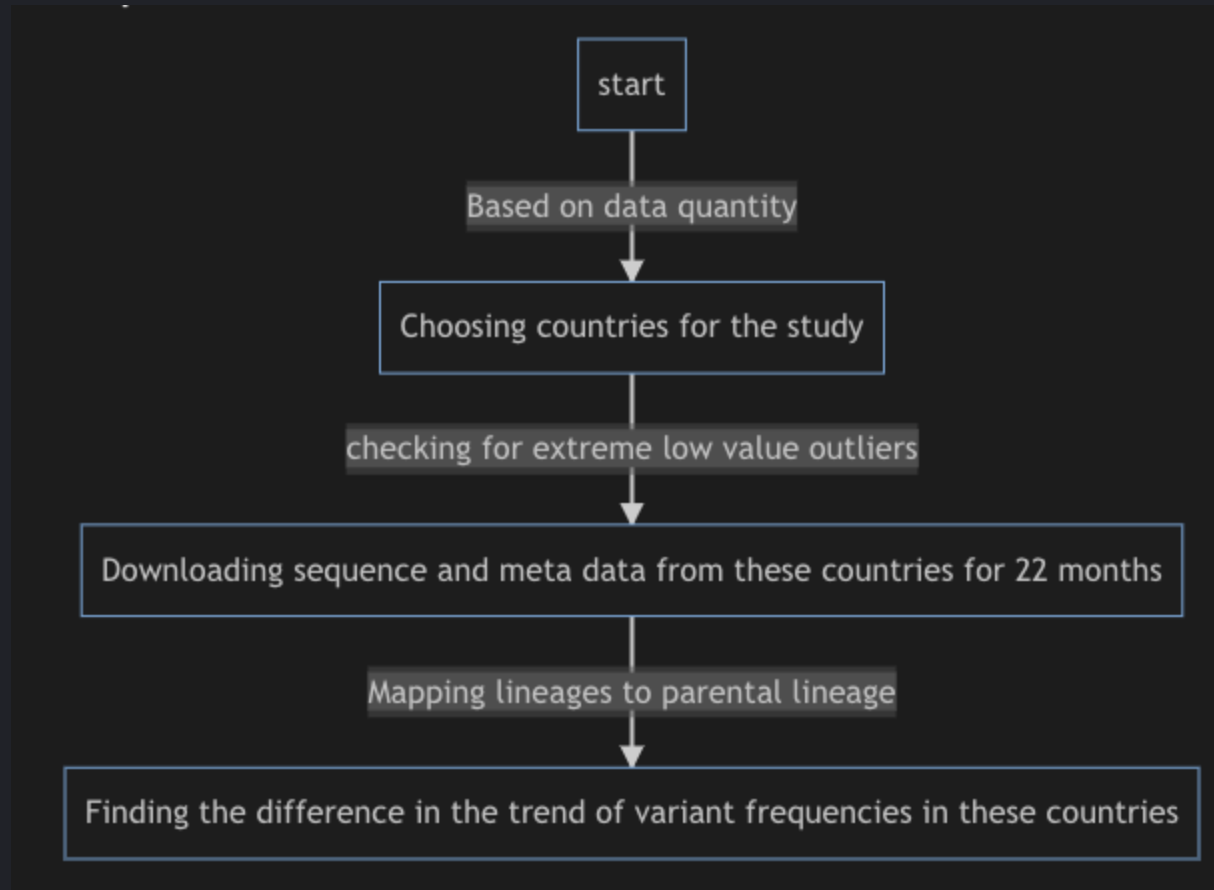


Objective

Studying the influence of the population genetics in the
SARS CoV2 variant susceptibility



Work done in steps

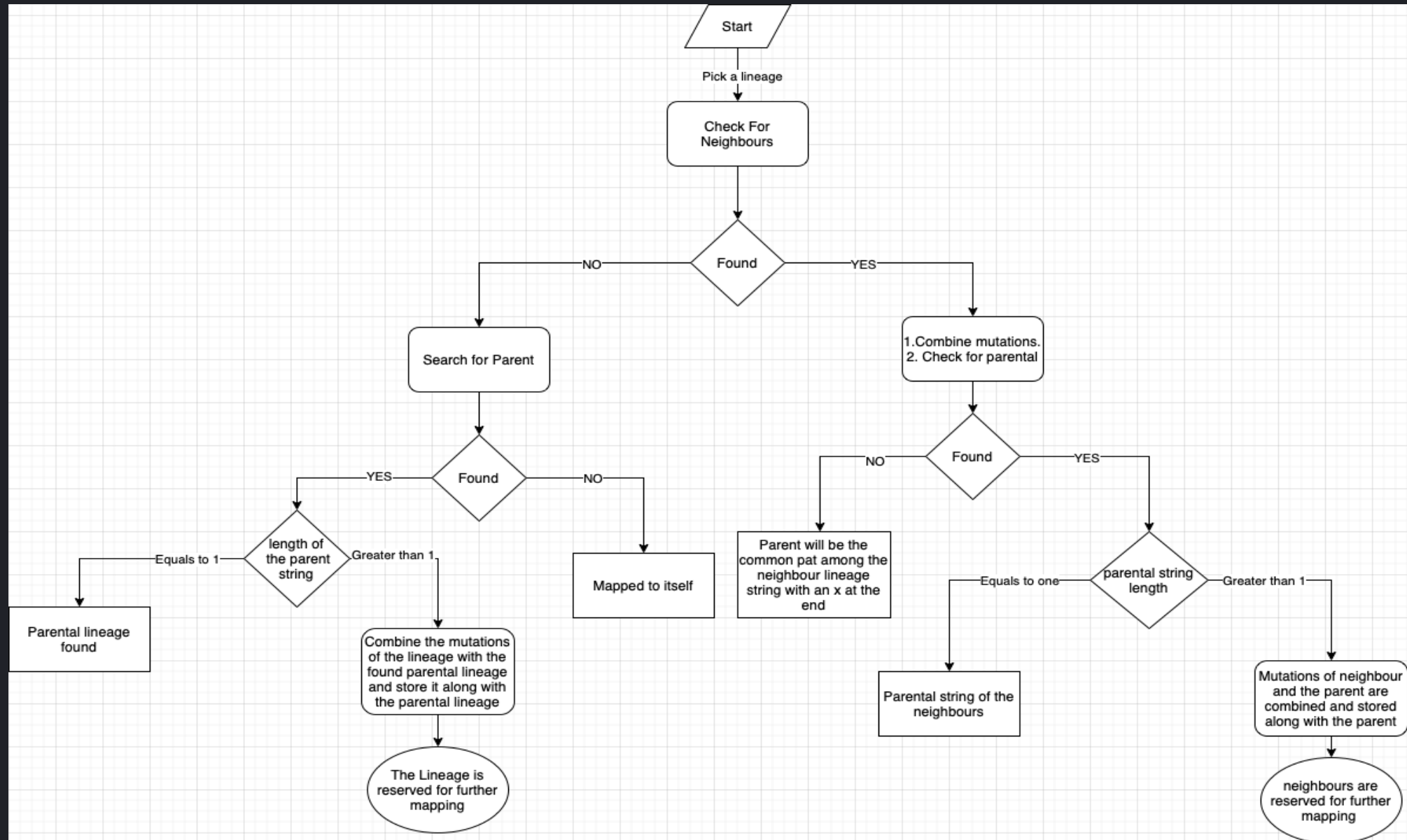


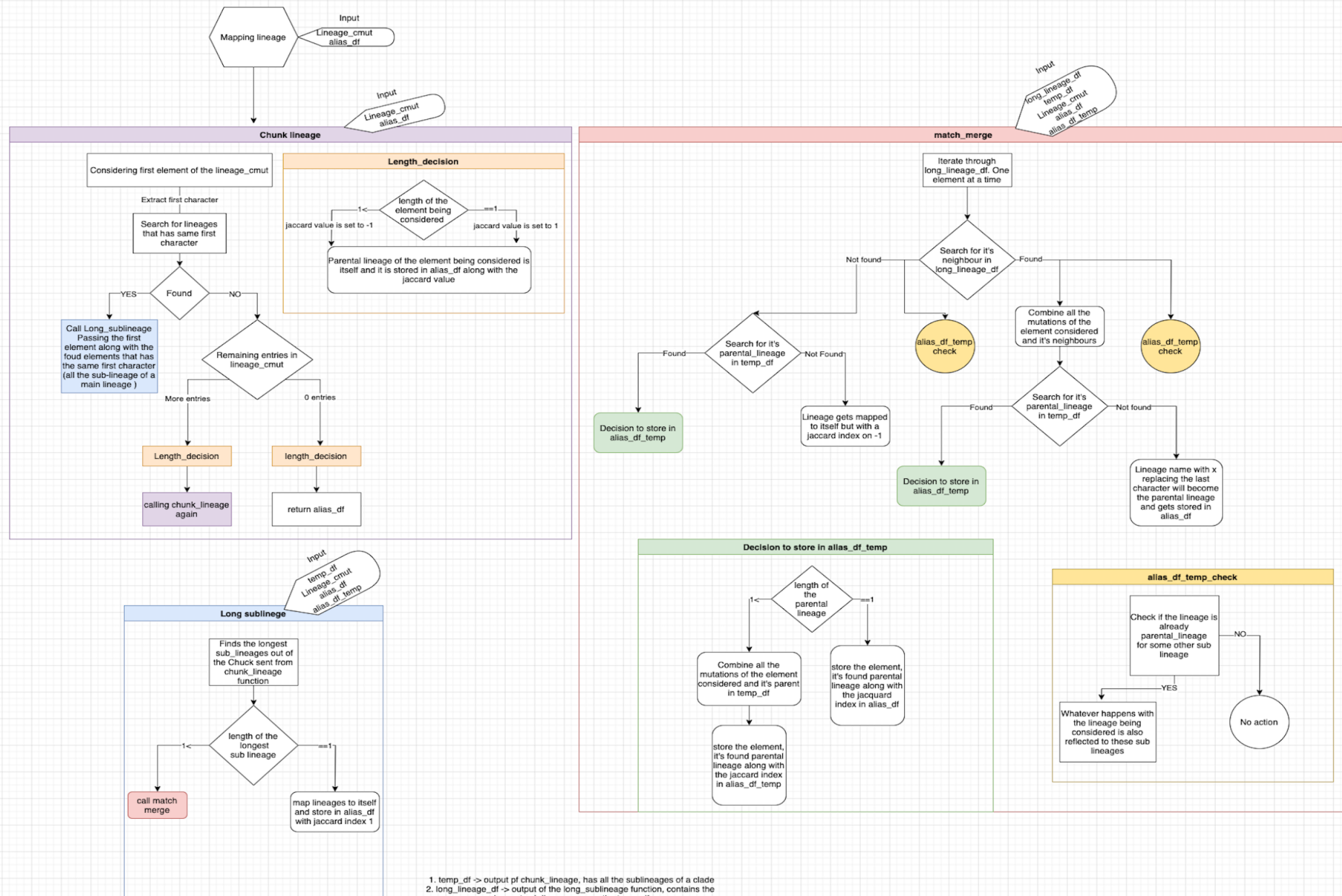
Mapping Lineages

For the variants

- BA.2.86+BA.2.86.*
- B.1.351+B.1.351.2+B.1.351.3
- B.1.617.2+AY.*
- B.1.1.529+BA.*
- XBB.1.16+XBB.1.16.*
- XBB.1.5+XBB.1.5.*
- XBB.1.9.1+XBB.1.9.1.*
- XBB+XBB.*

For other lineages





Inference

Interesting inference