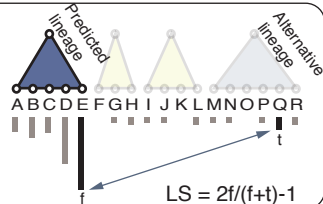


Load the RASE database

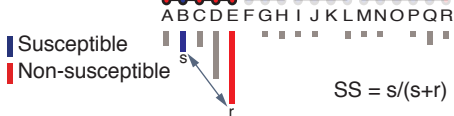


Read next nanopore read

Find nearest neighbor and predict lineage



Predict susceptibility or resistance



Predict and report

Update weights

Match against the database

ProPhyle

