

Primary cluster 1

Comp method: fastANI

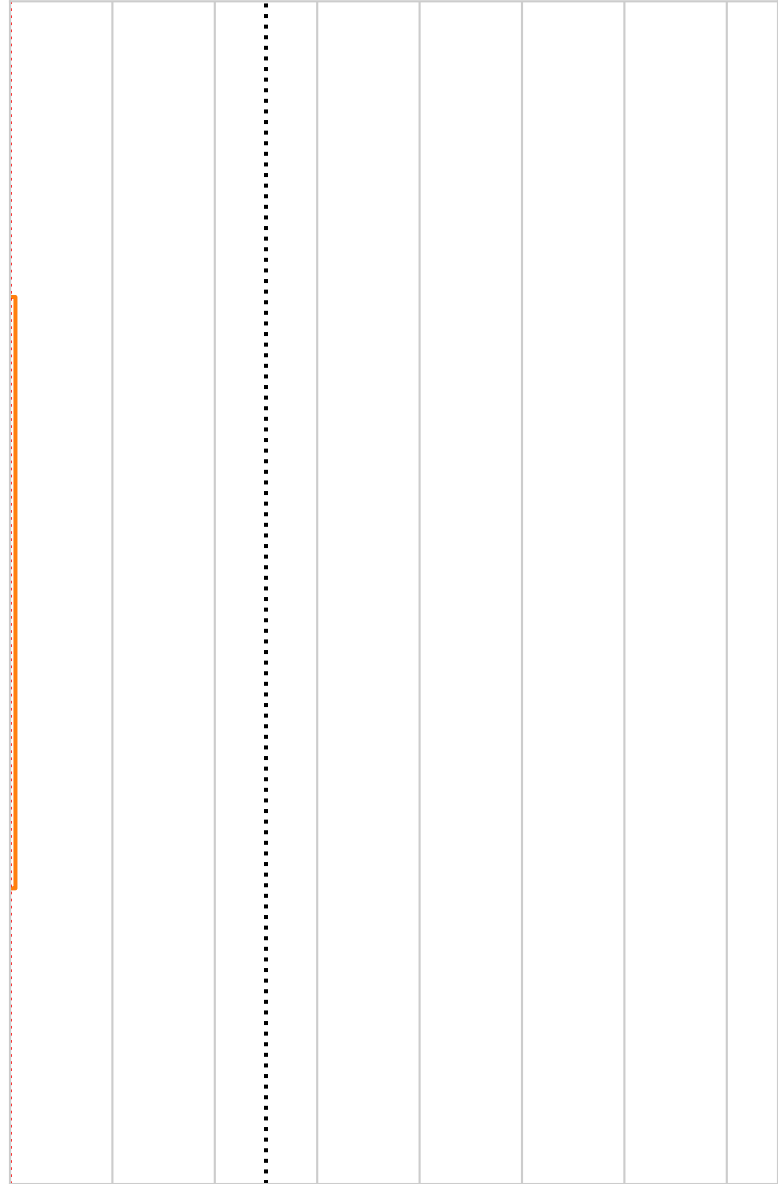
Clust method: average

Min cov: 0.1

target.reordered.fasta

H37Rv.fasta

100.0 98.0 96.0 94.0 92.0 90.0 88.0 86.0
Average Nucleotide Identity (ANI)



Primary cluster 2

