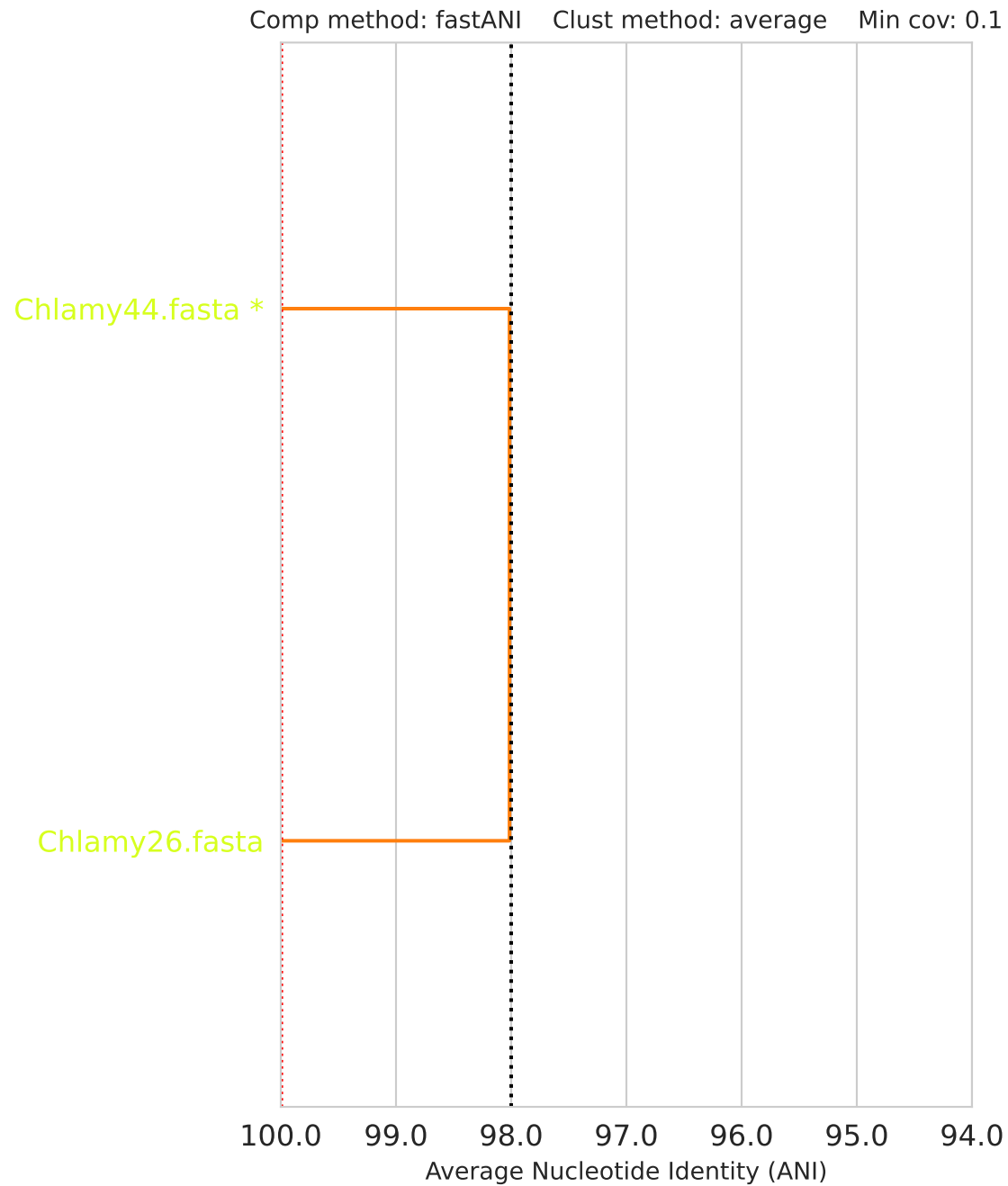
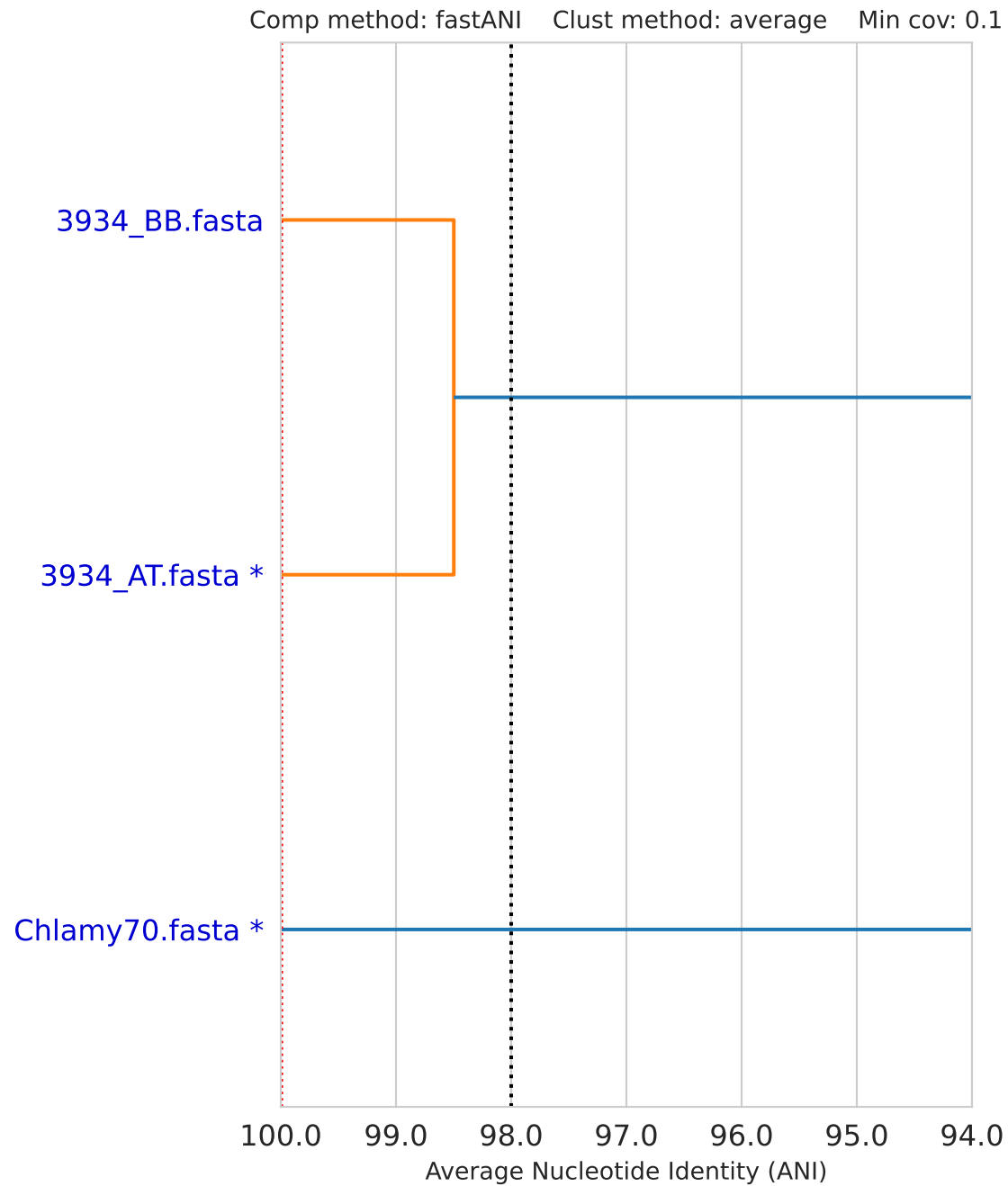


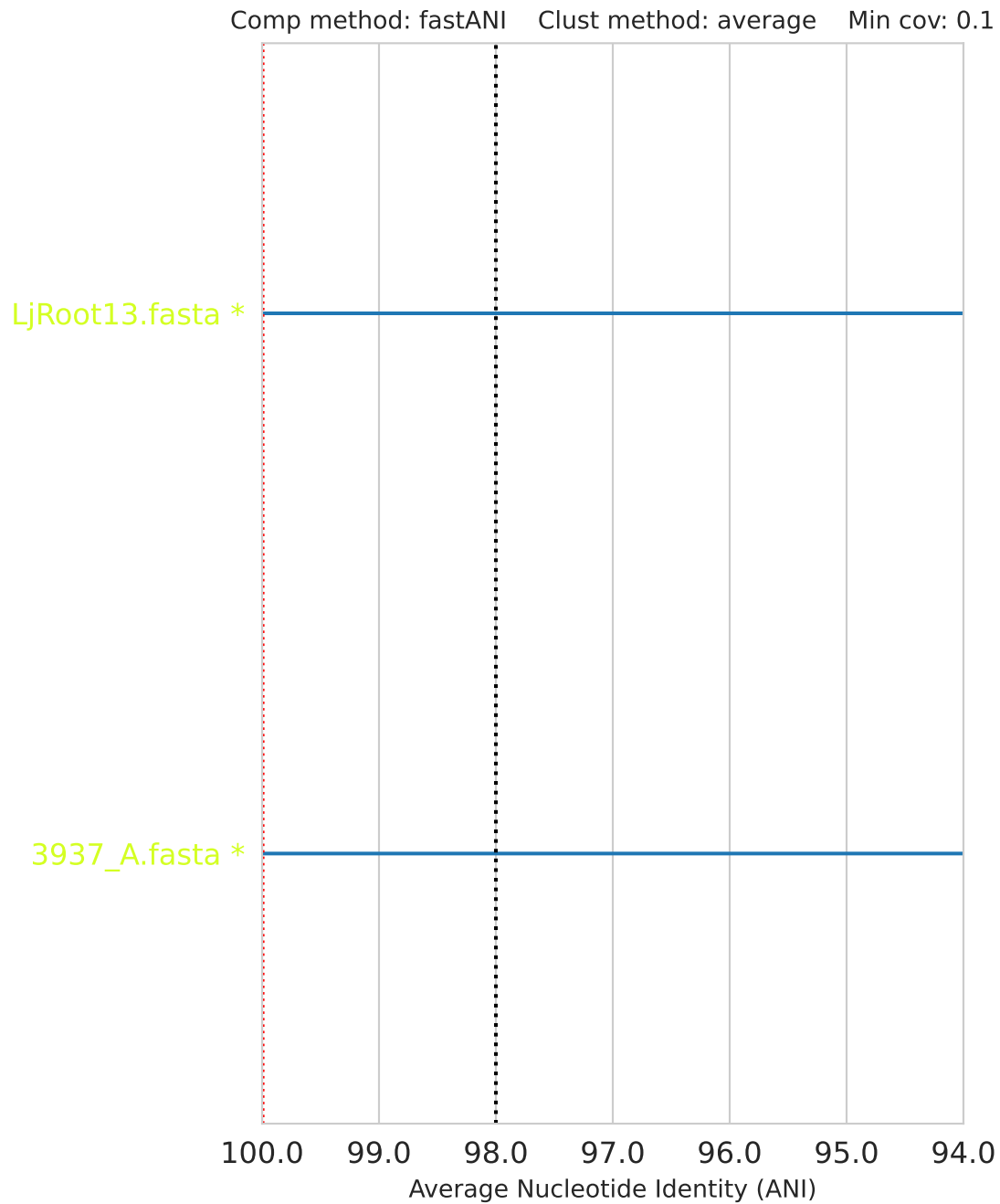
Primary cluster 45



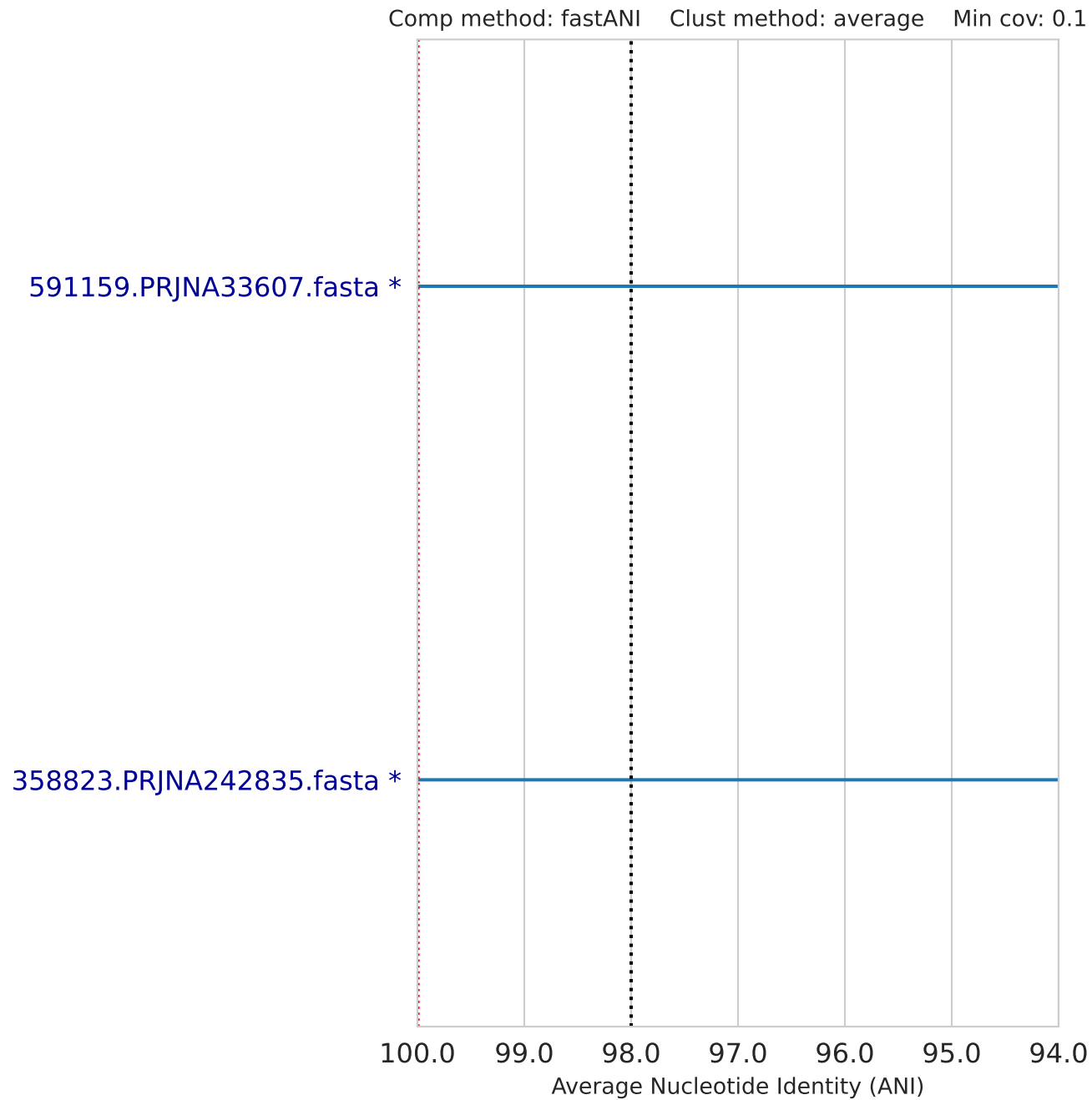
Primary cluster 51



Primary cluster 53



Primary cluster 75



Primary cluster 92

Comp method: fastANI

Clust method: average

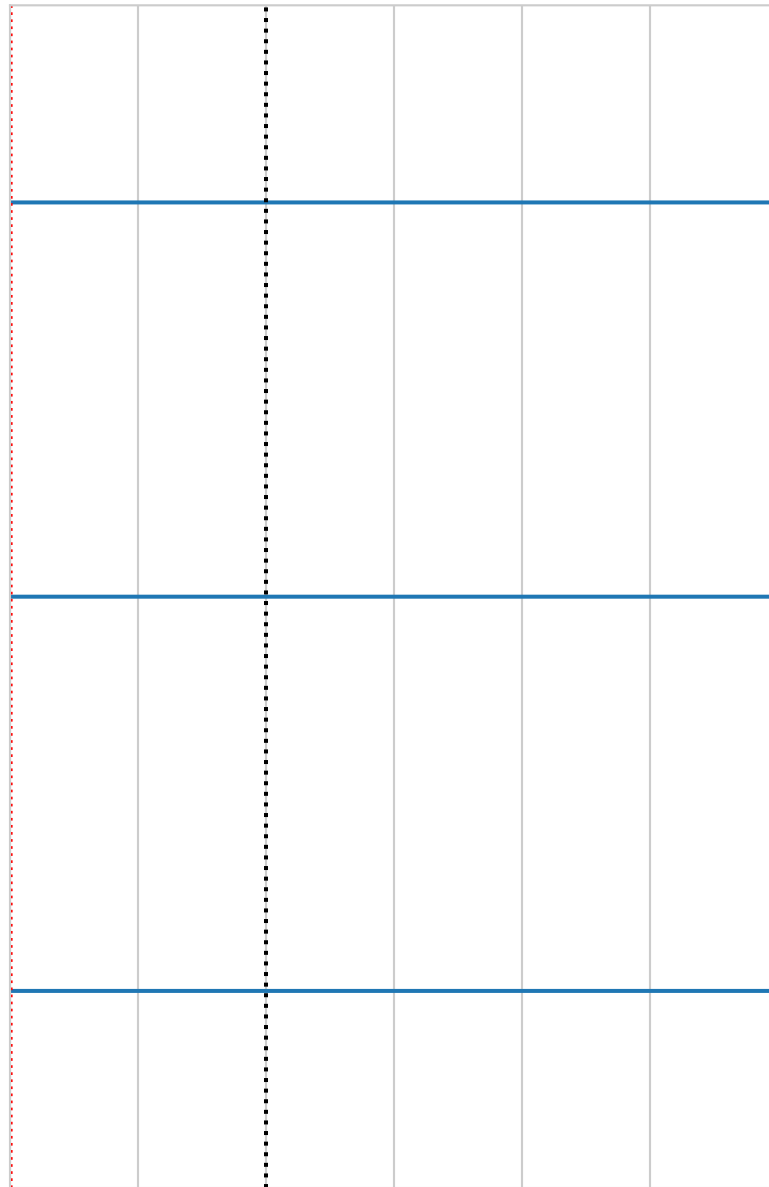
Min cov: 0.1

749927.PRJNA46889.fasta *

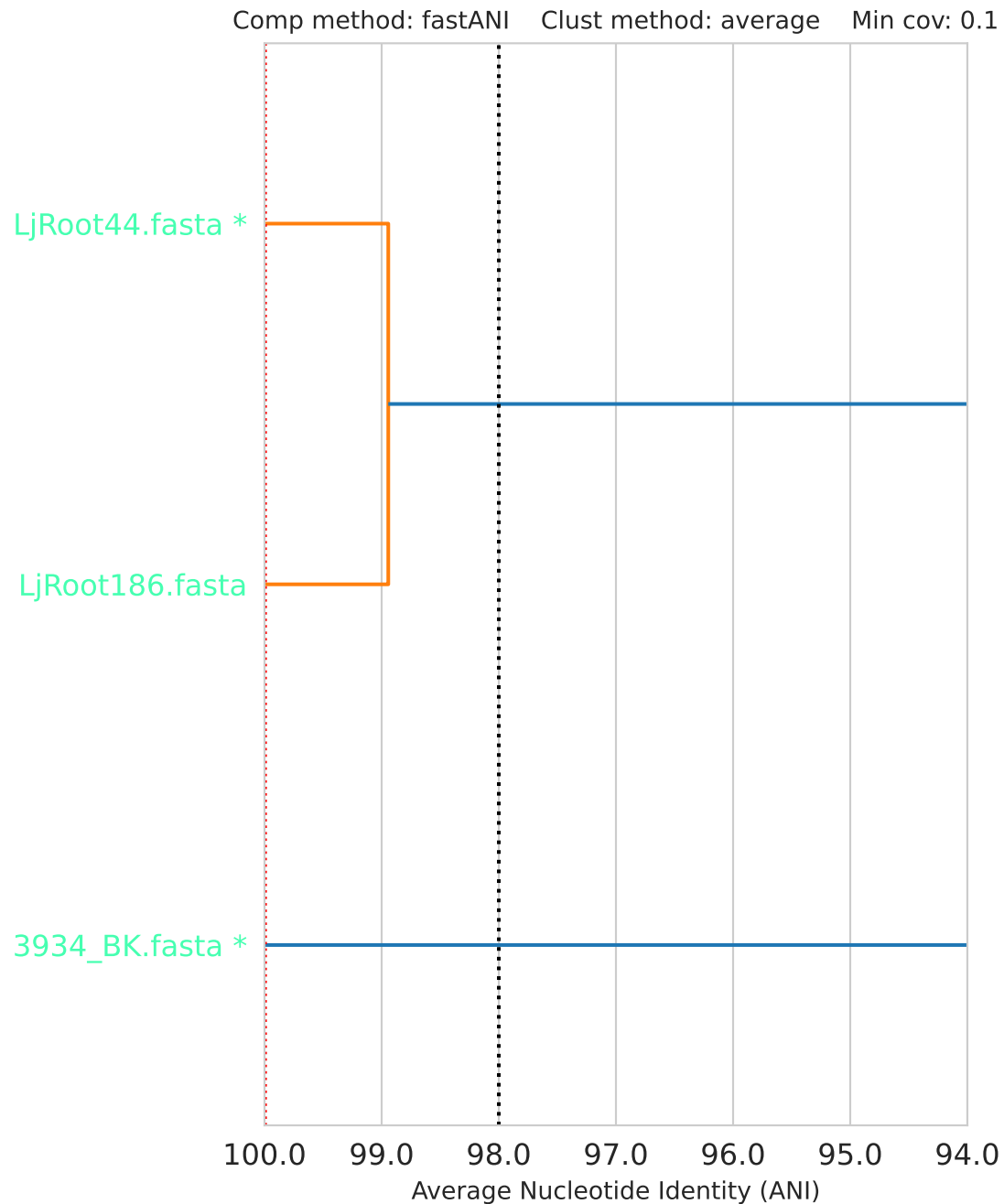
287986.PRJNA244719.fasta *

1089545.PRJNA165347.fasta *

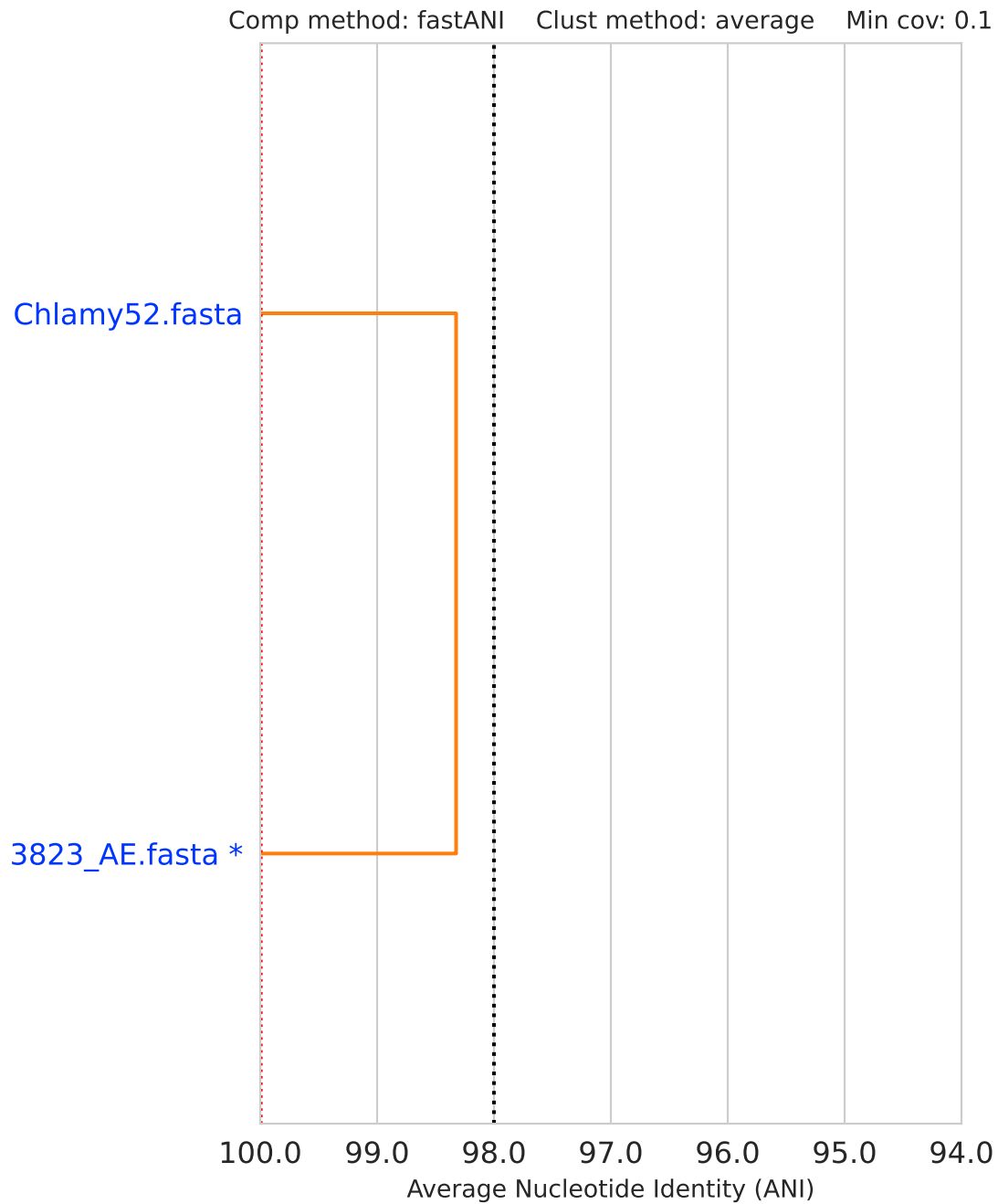
100.0 99.0 98.0 97.0 96.0 95.0 94.0
Average Nucleotide Identity (ANI)



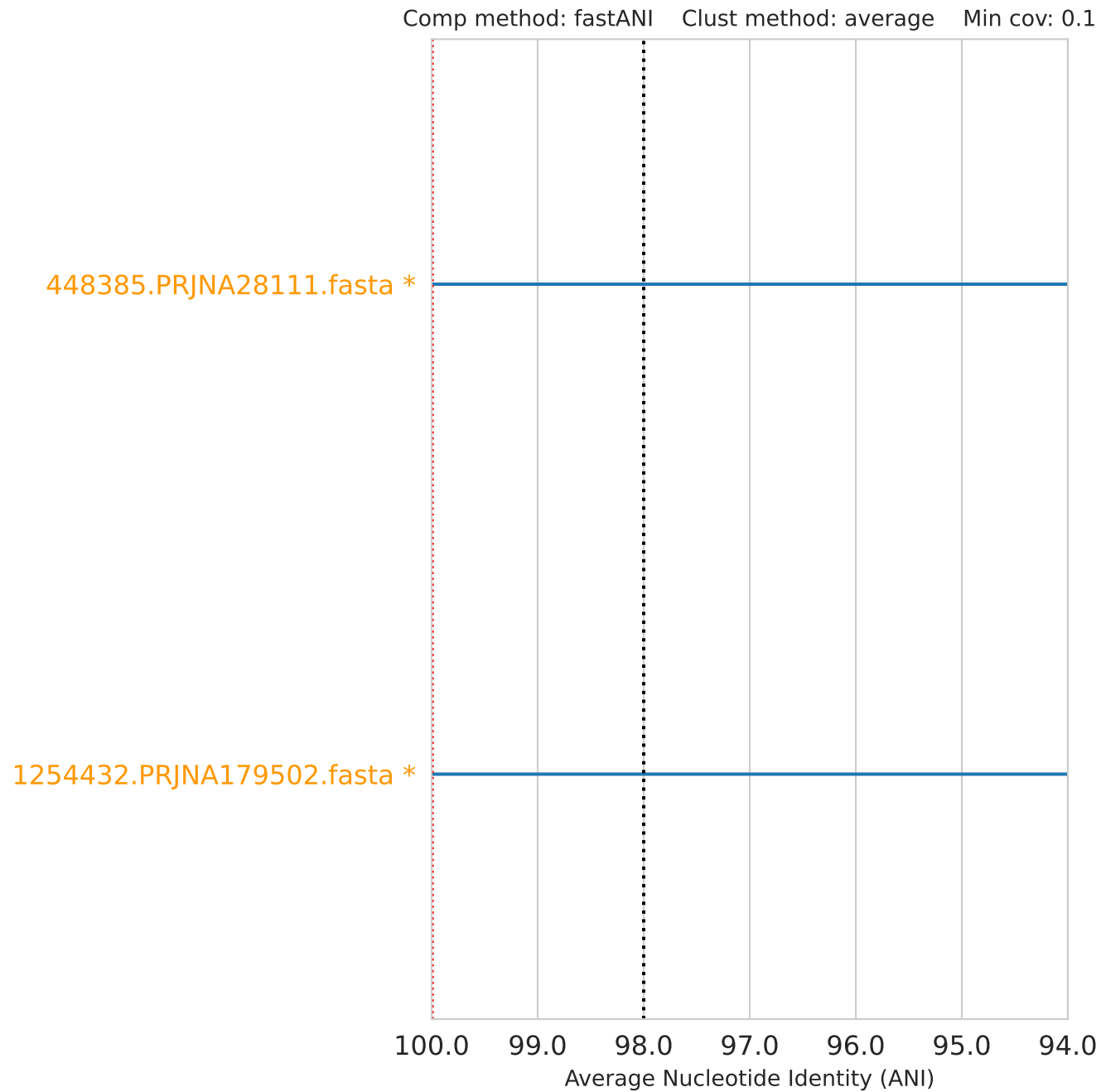
Primary cluster 119



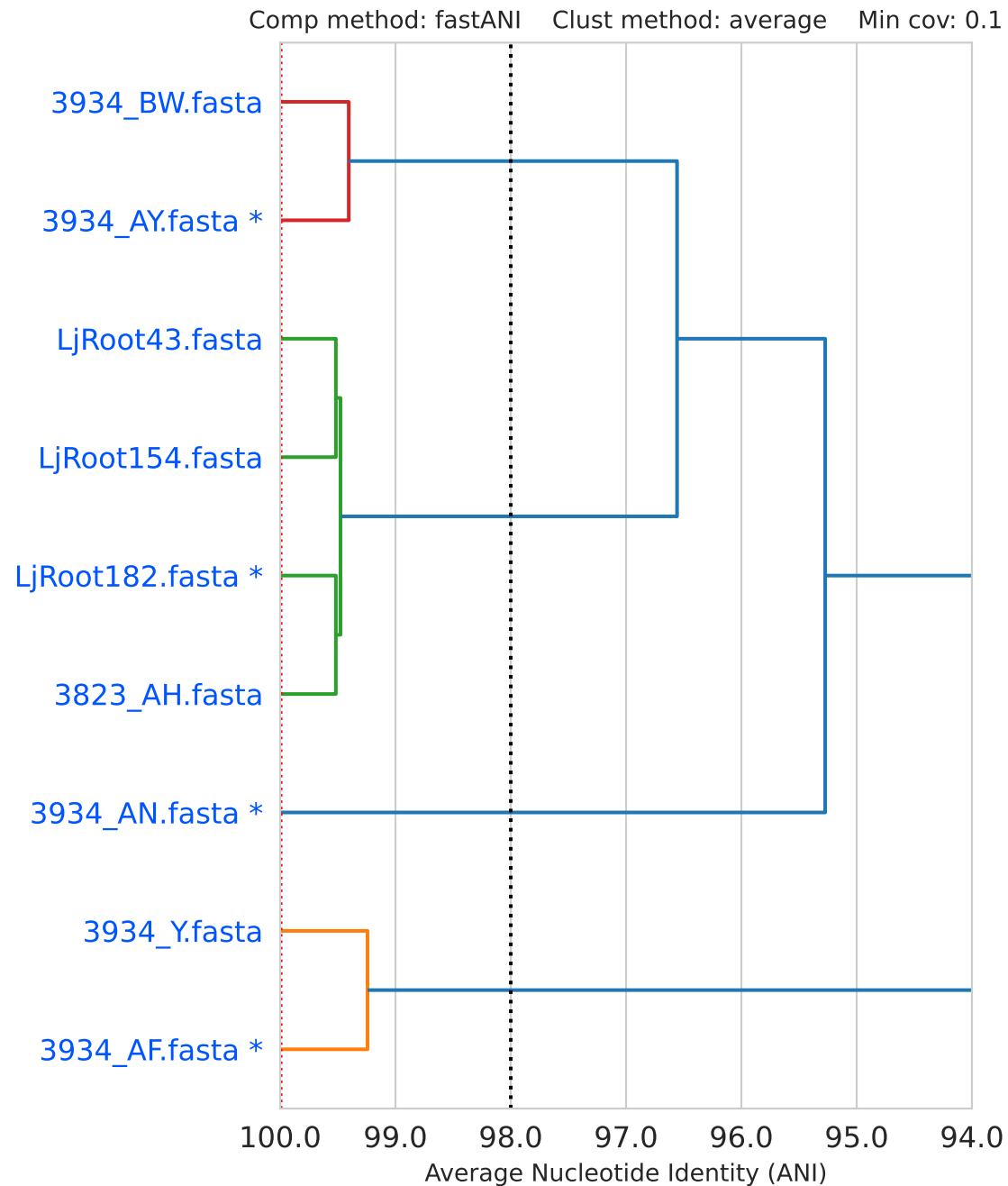
Primary cluster 120



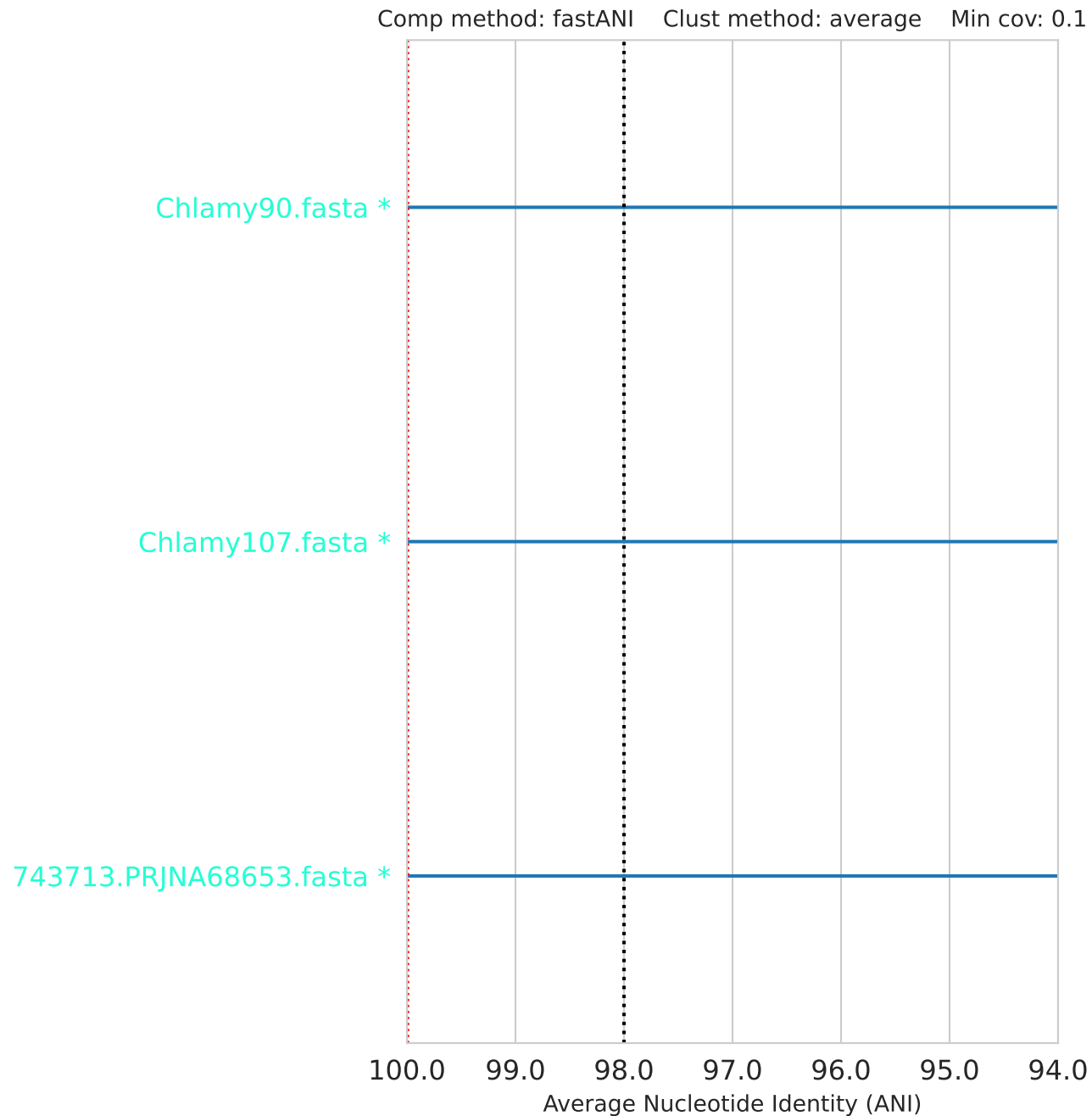
Primary cluster 138



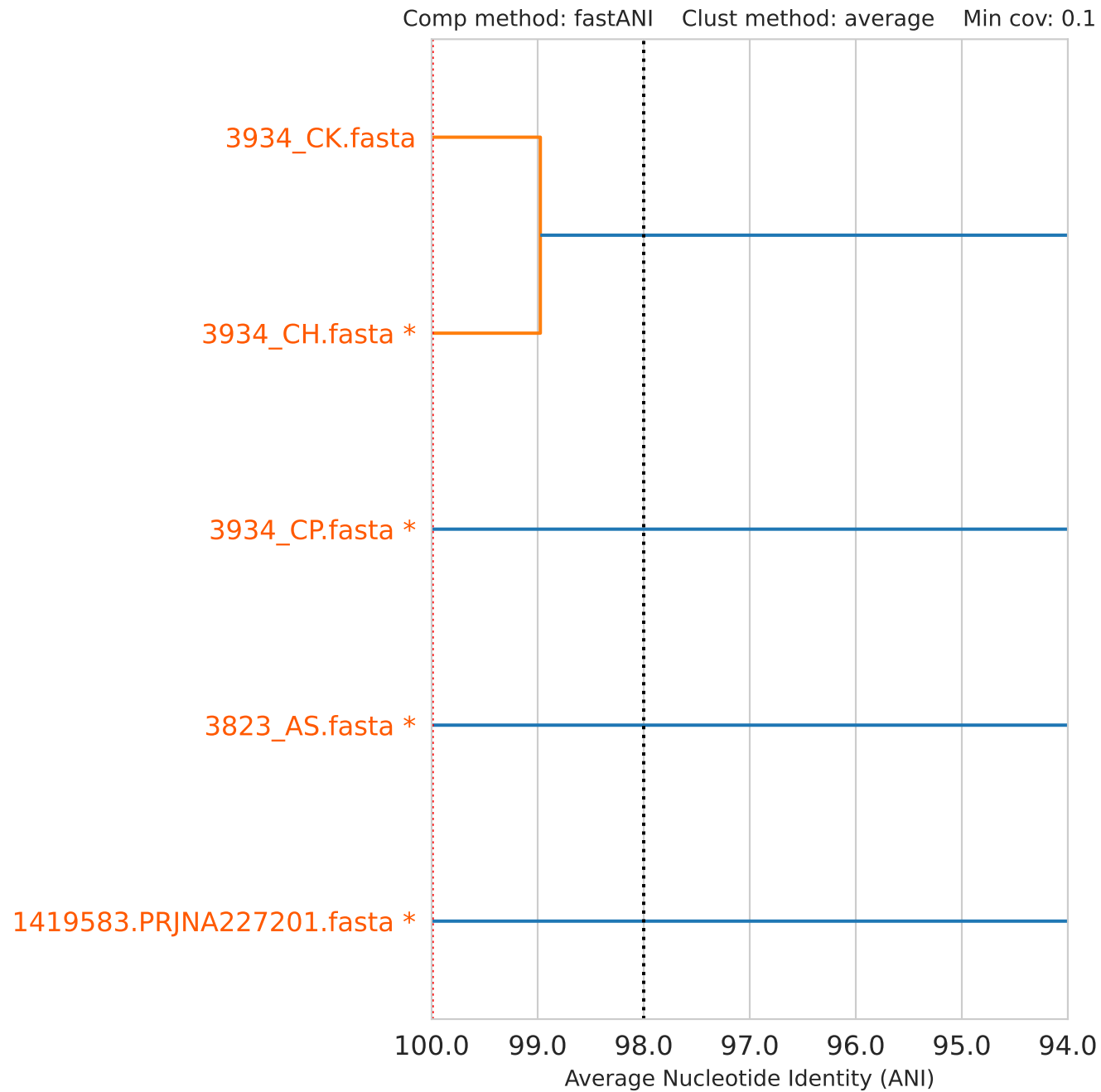
Primary cluster 151



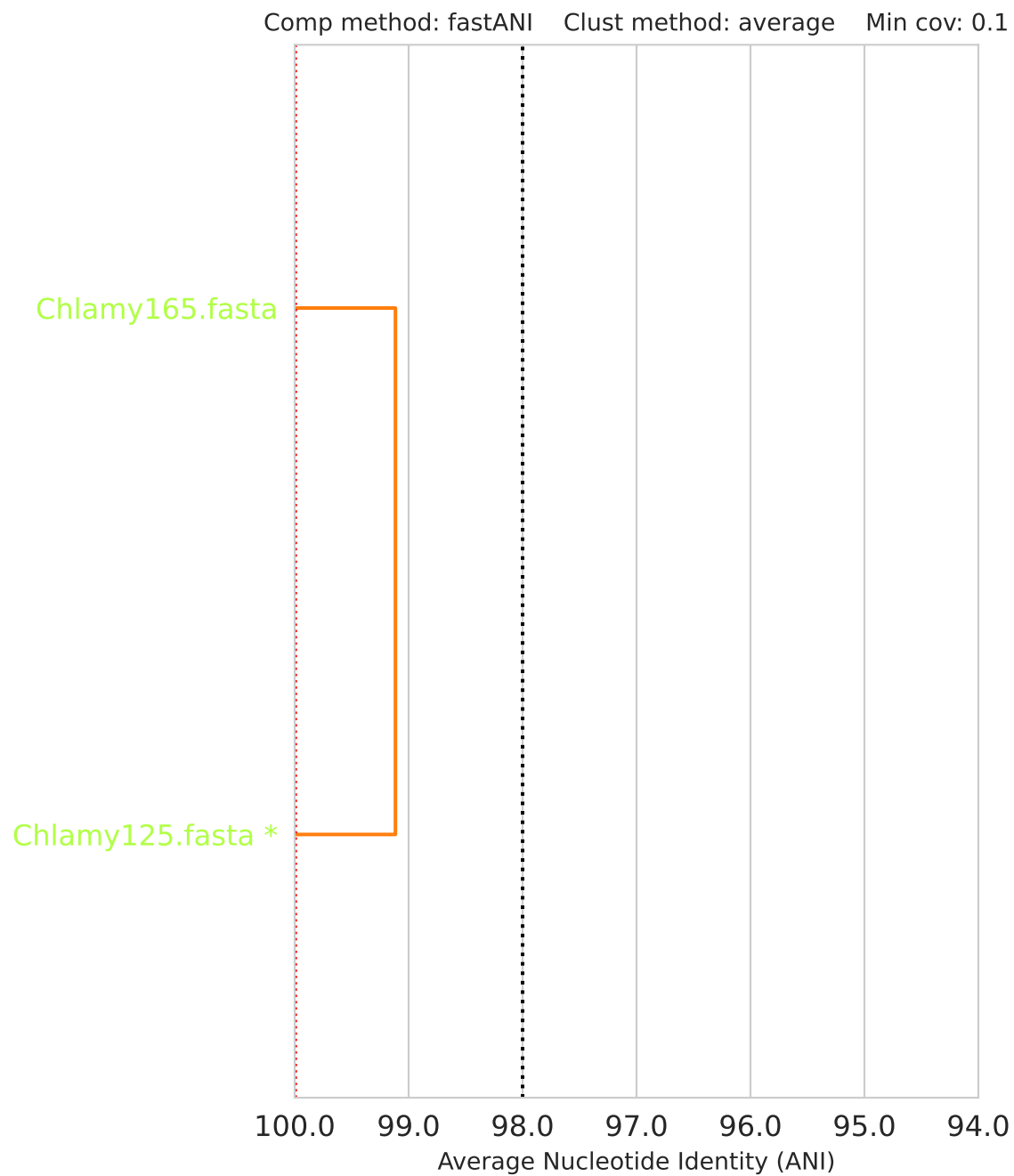
Primary cluster 154



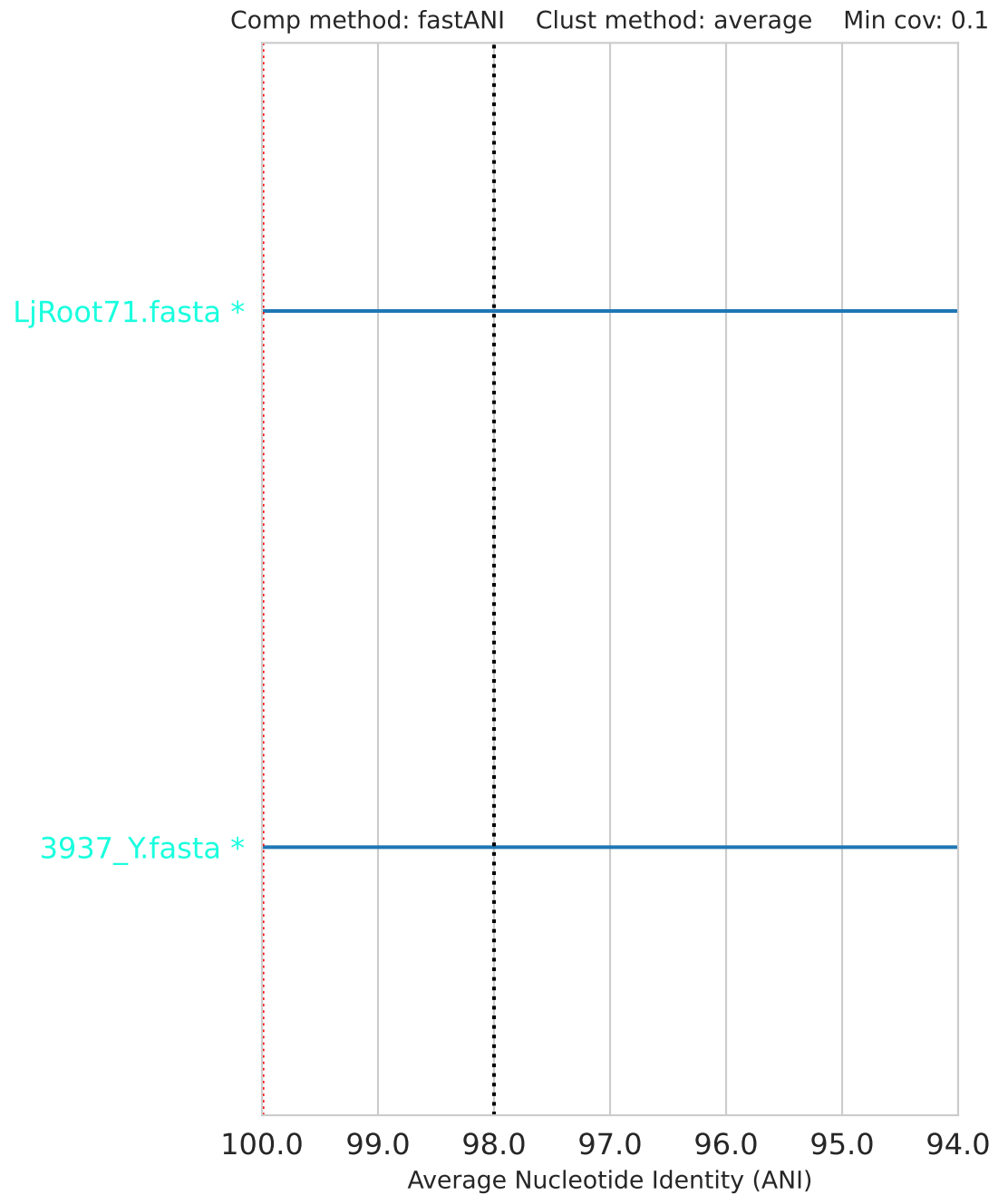
Primary cluster 155



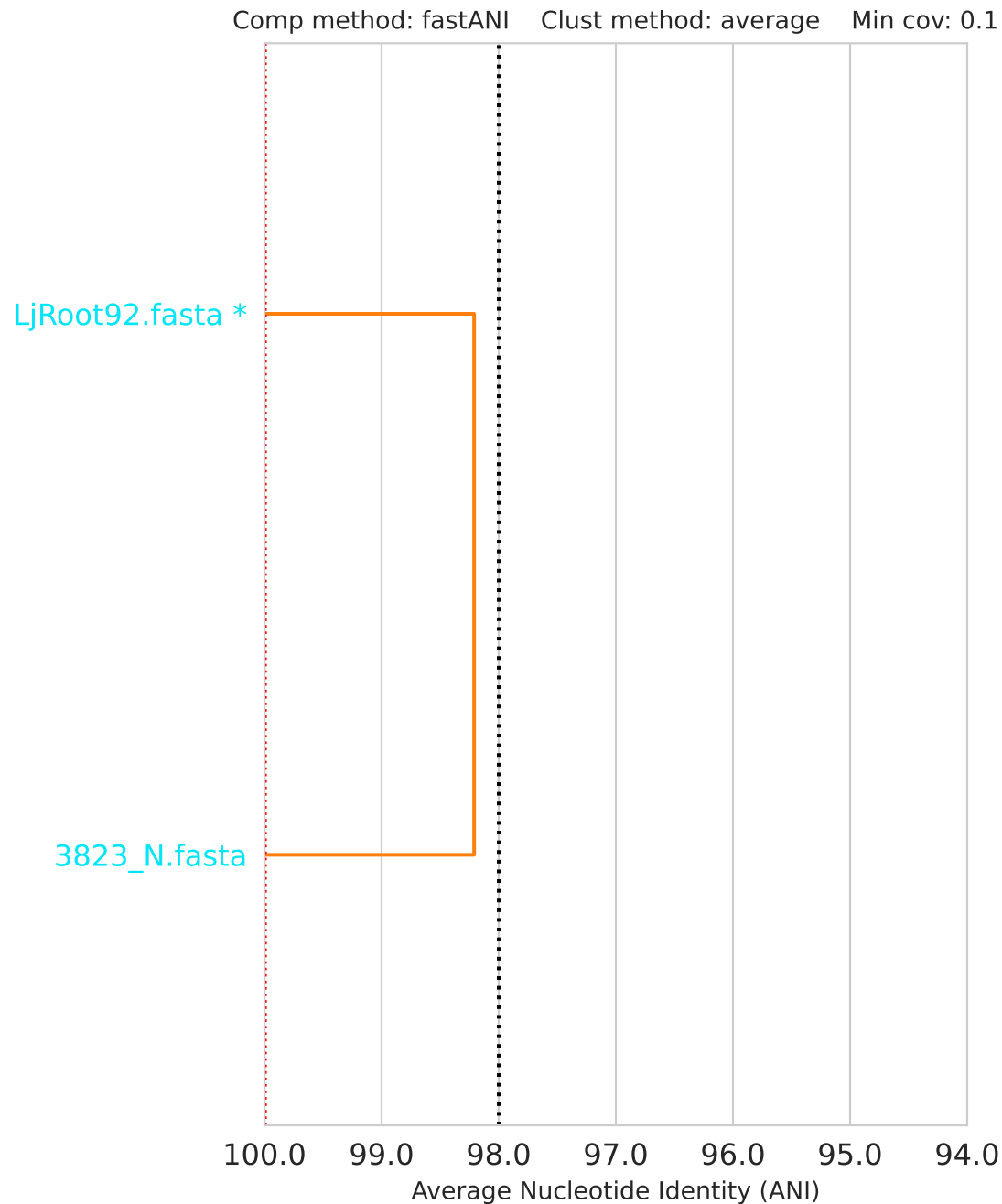
Primary cluster 156



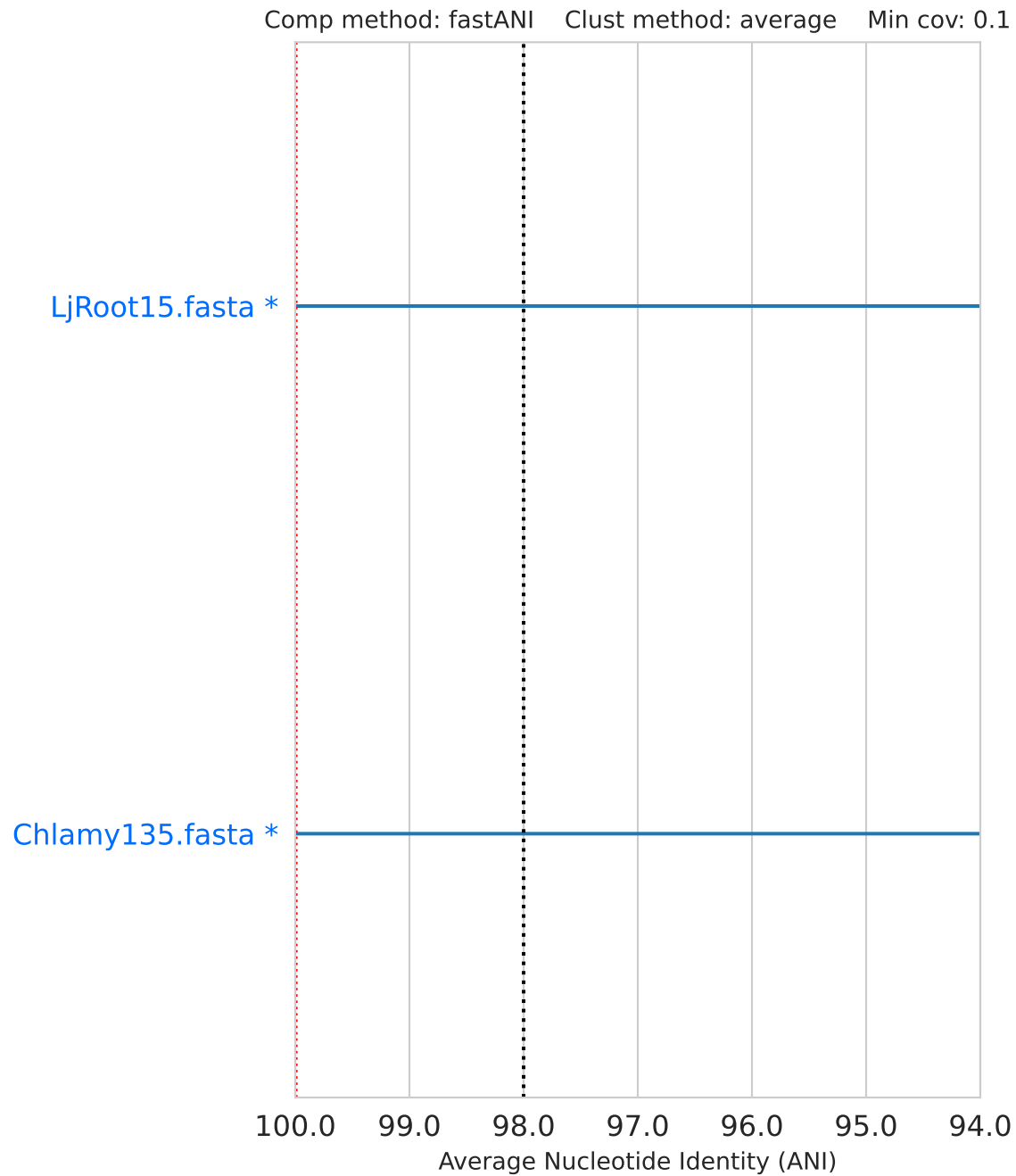
Primary cluster 163



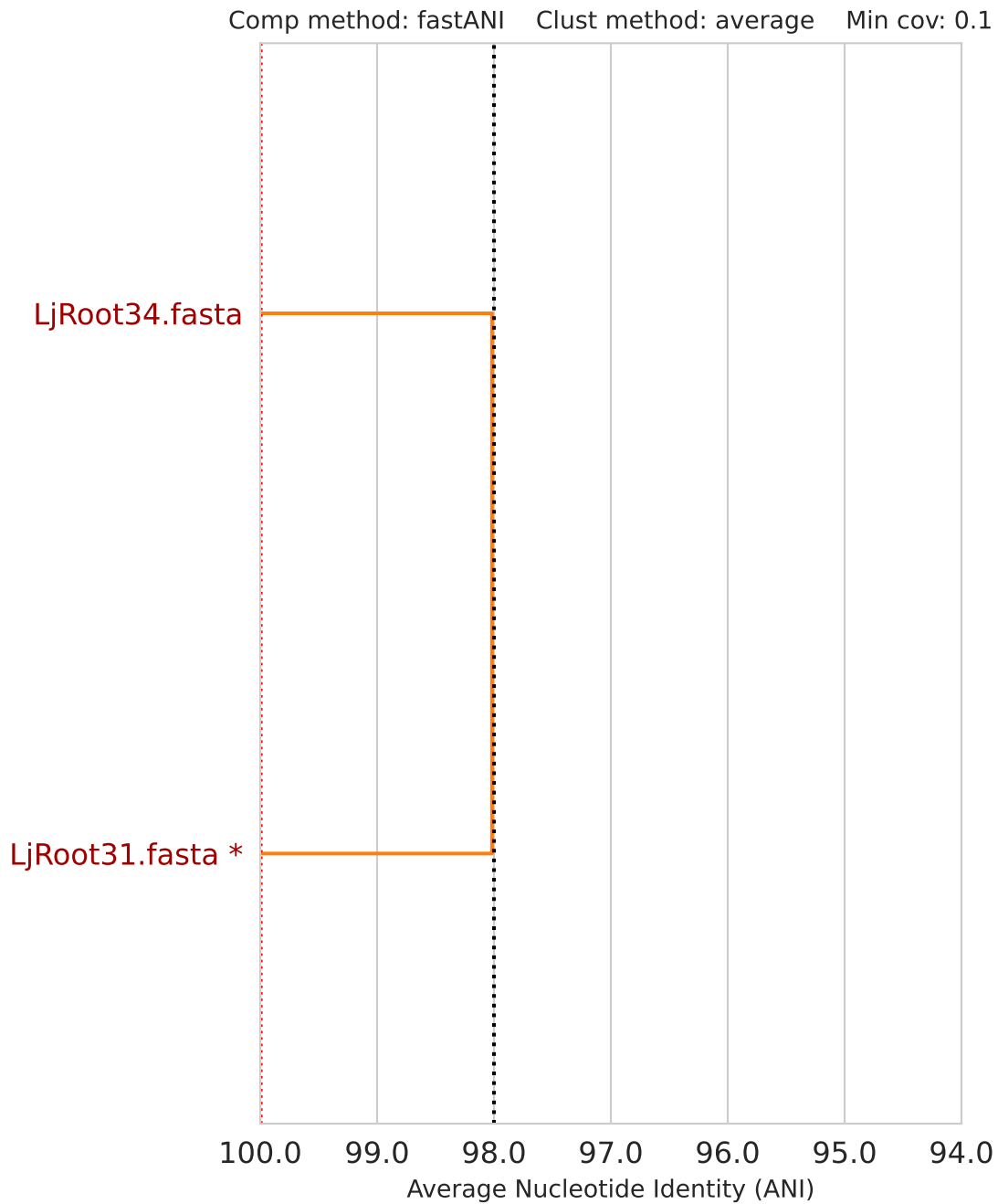
Primary cluster 164



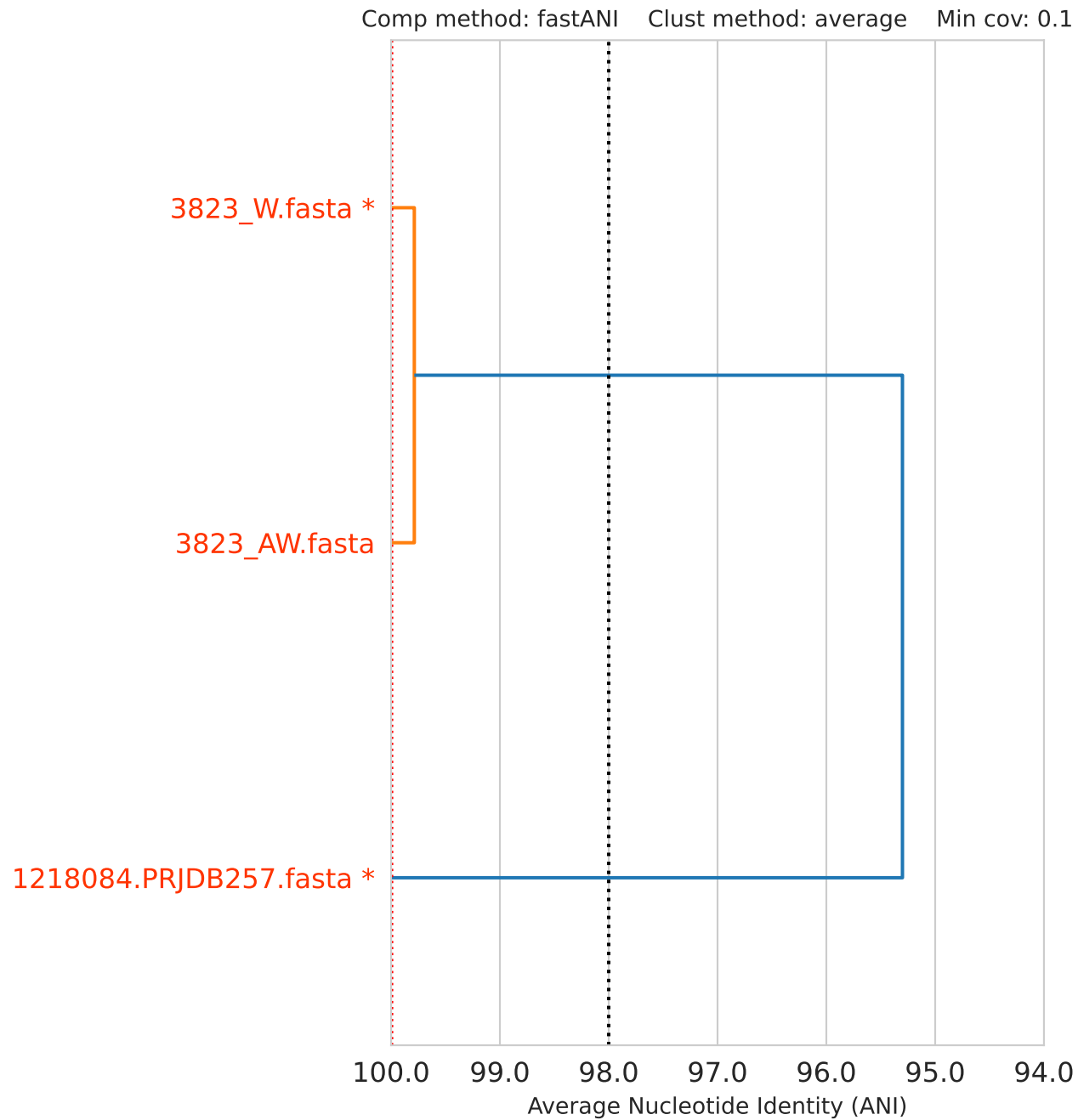
Primary cluster 183



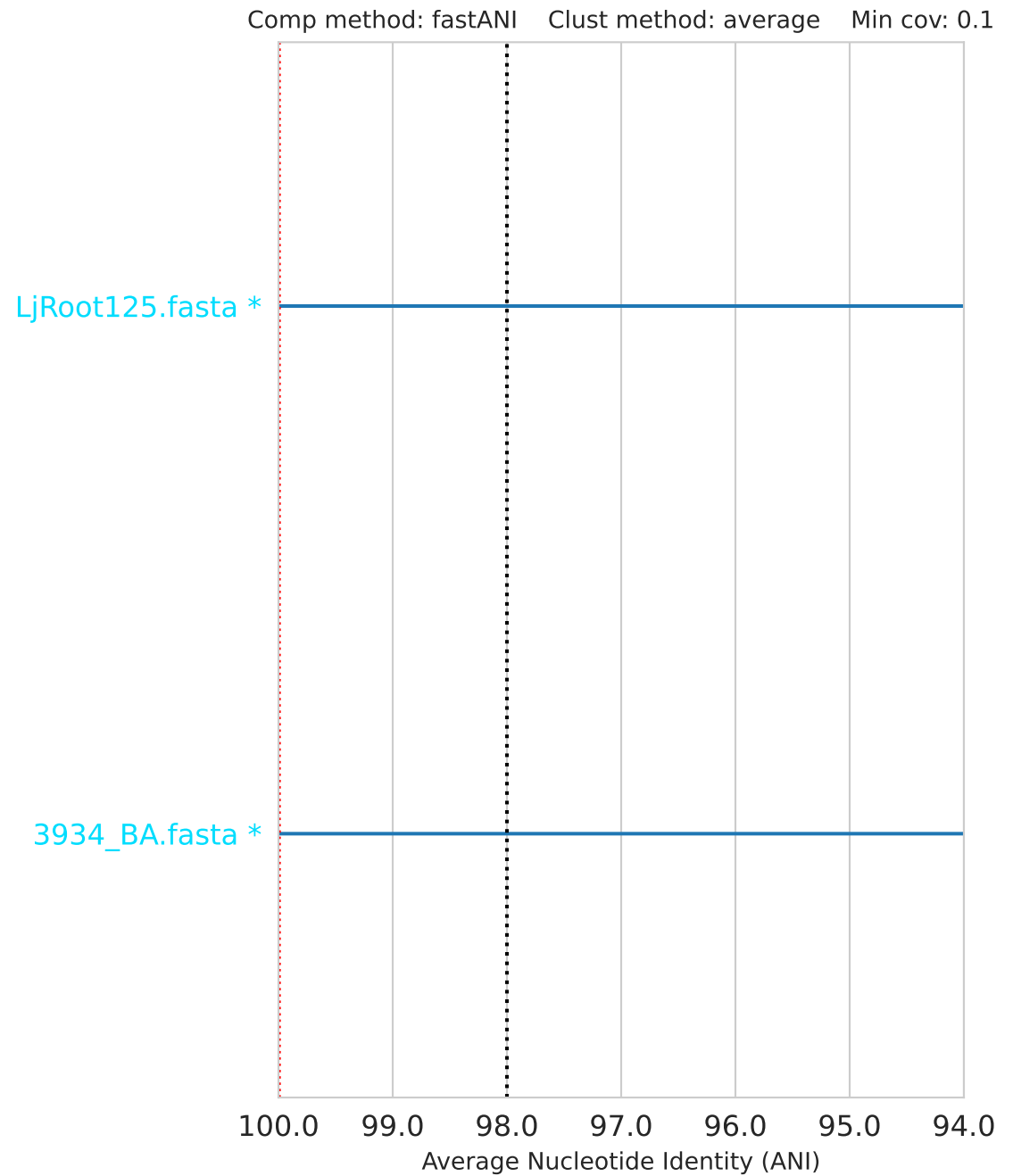
Primary cluster 209



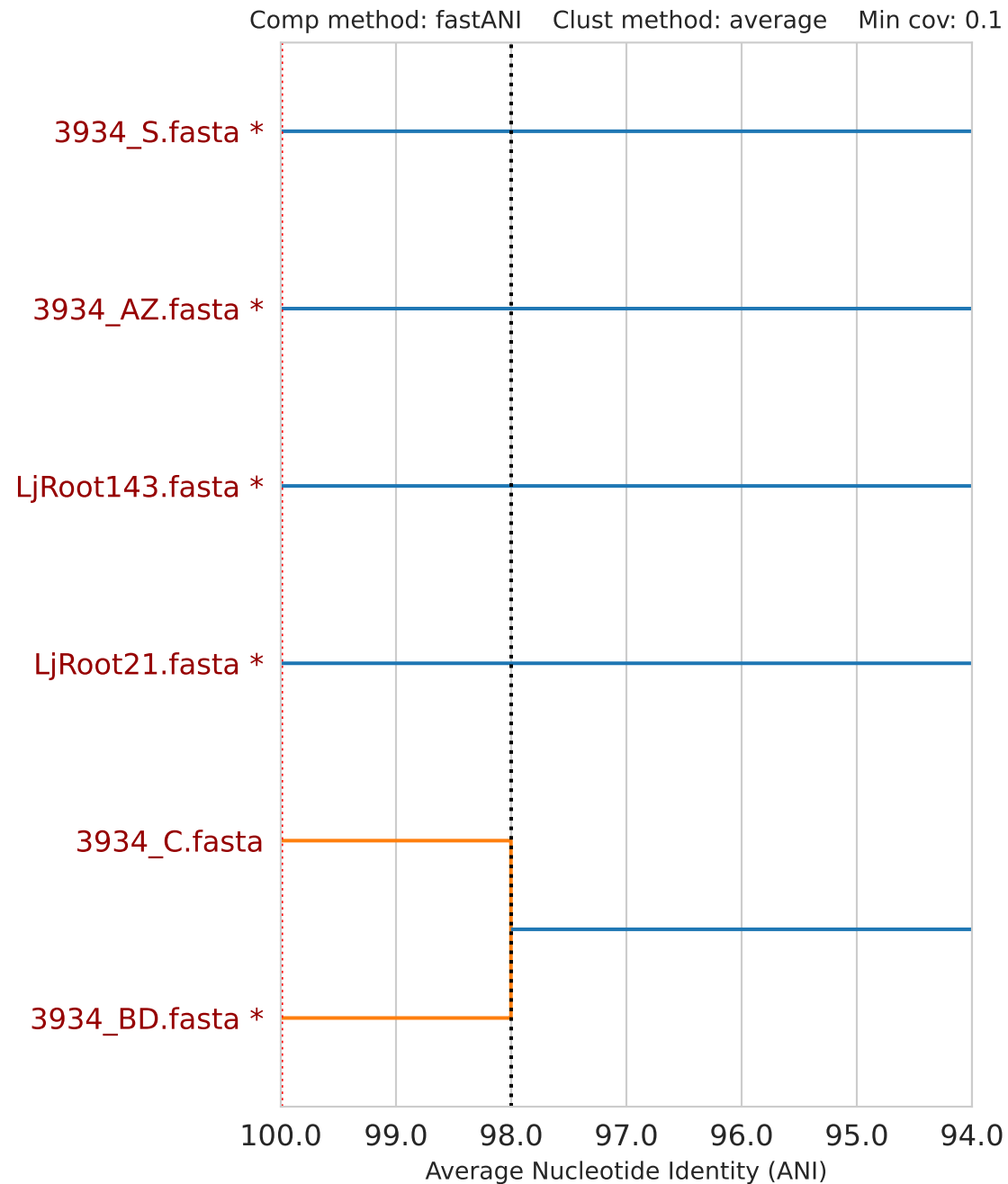
Primary cluster 211



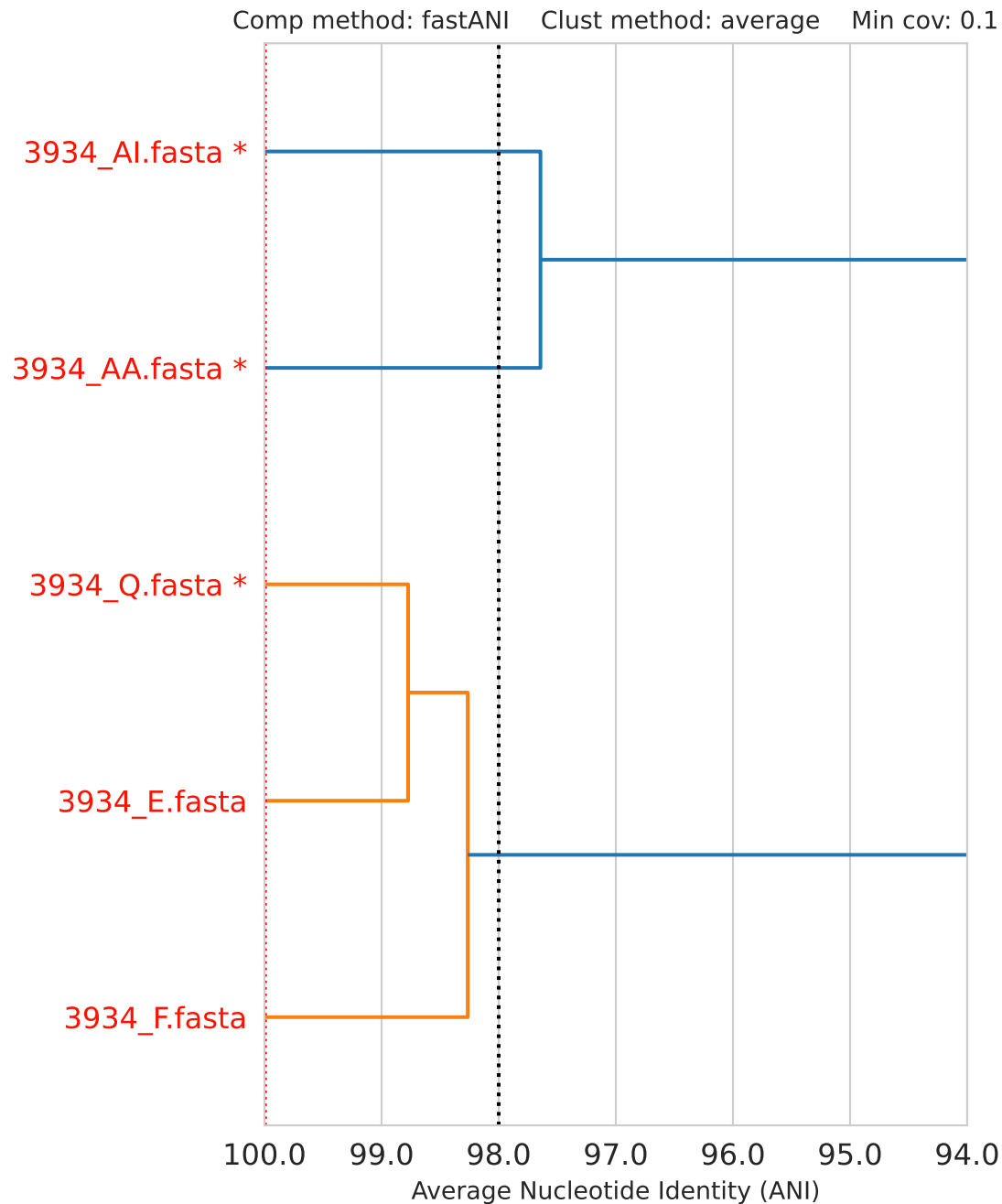
Primary cluster 218



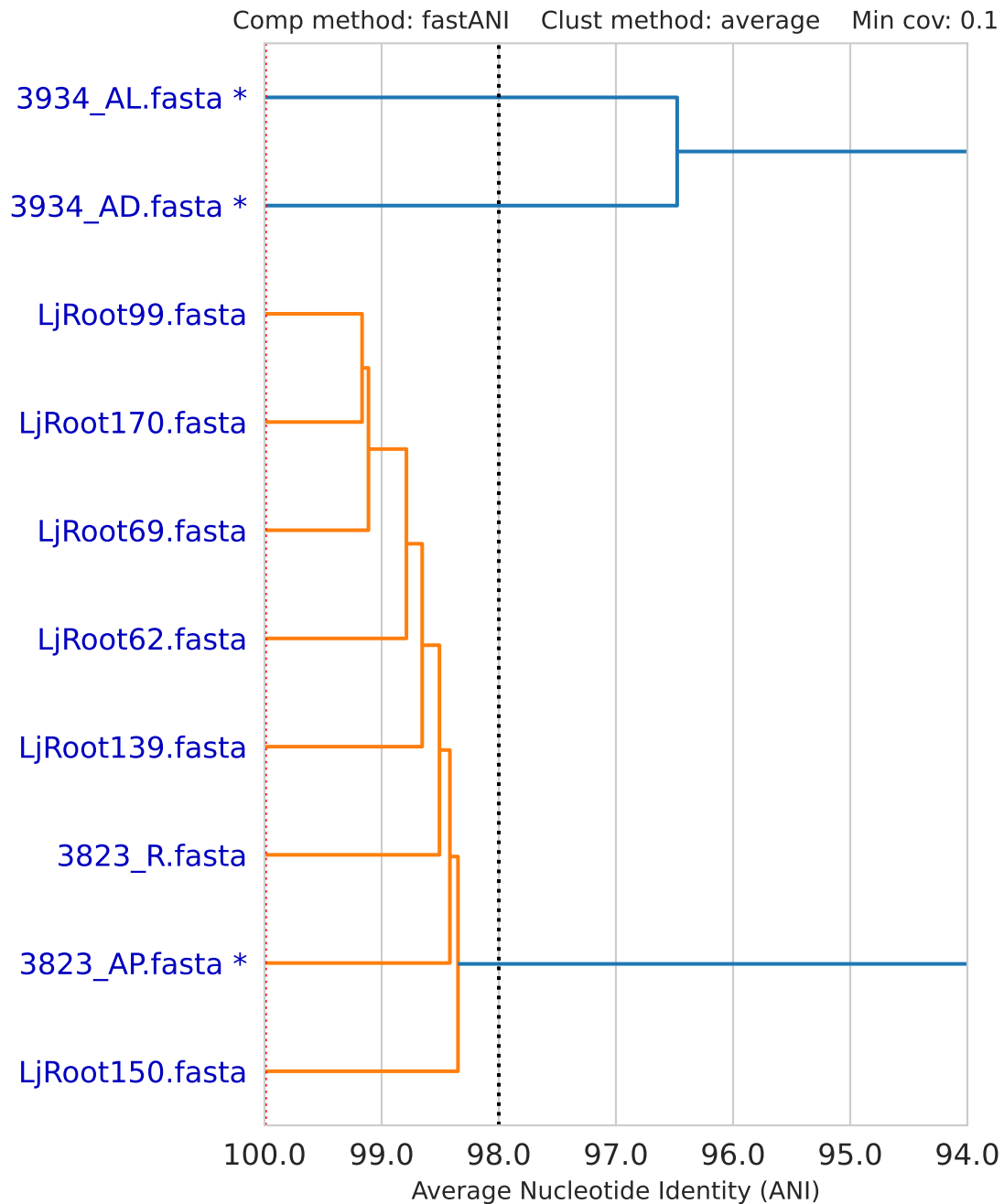
Primary cluster 220



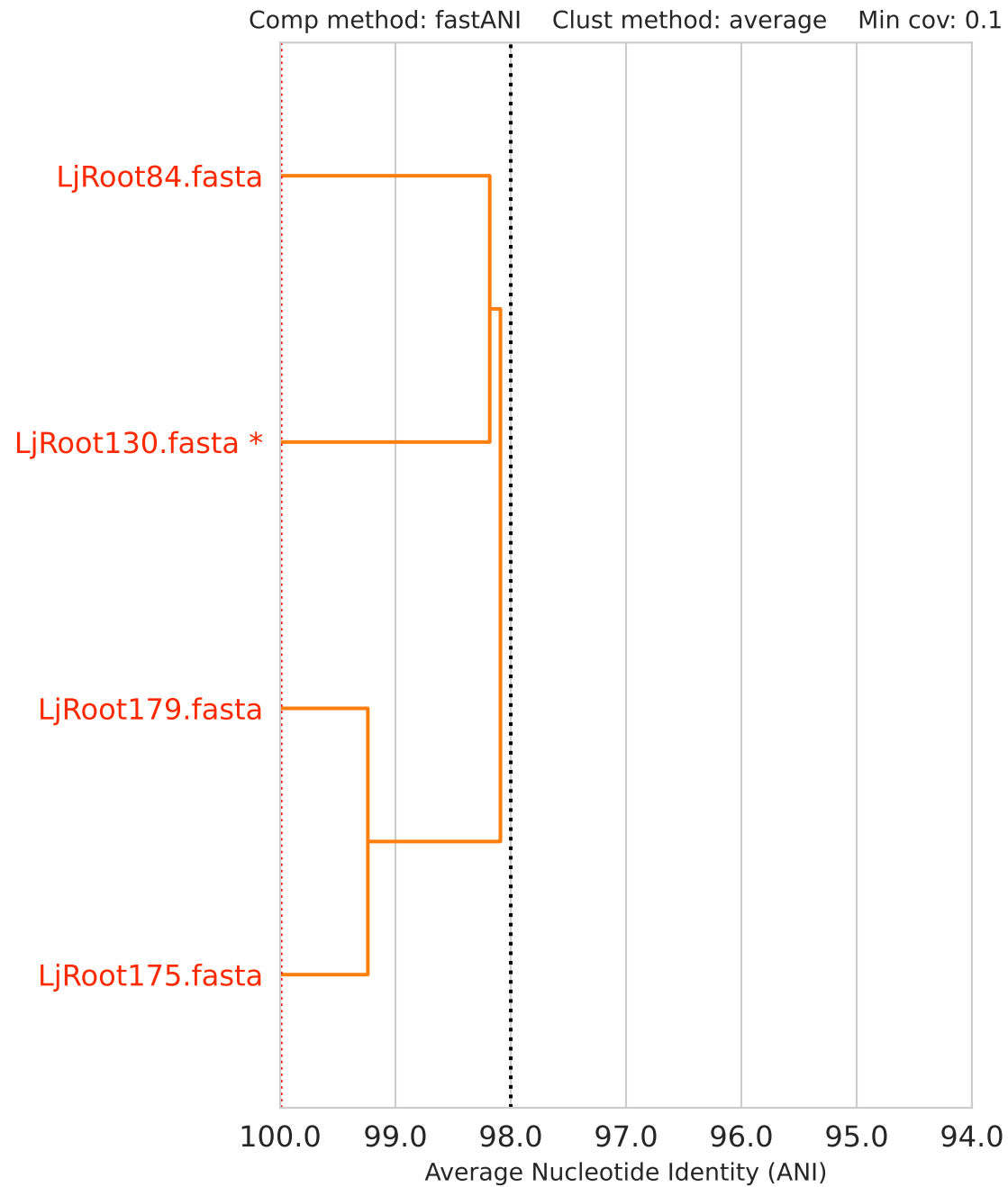
Primary cluster 222



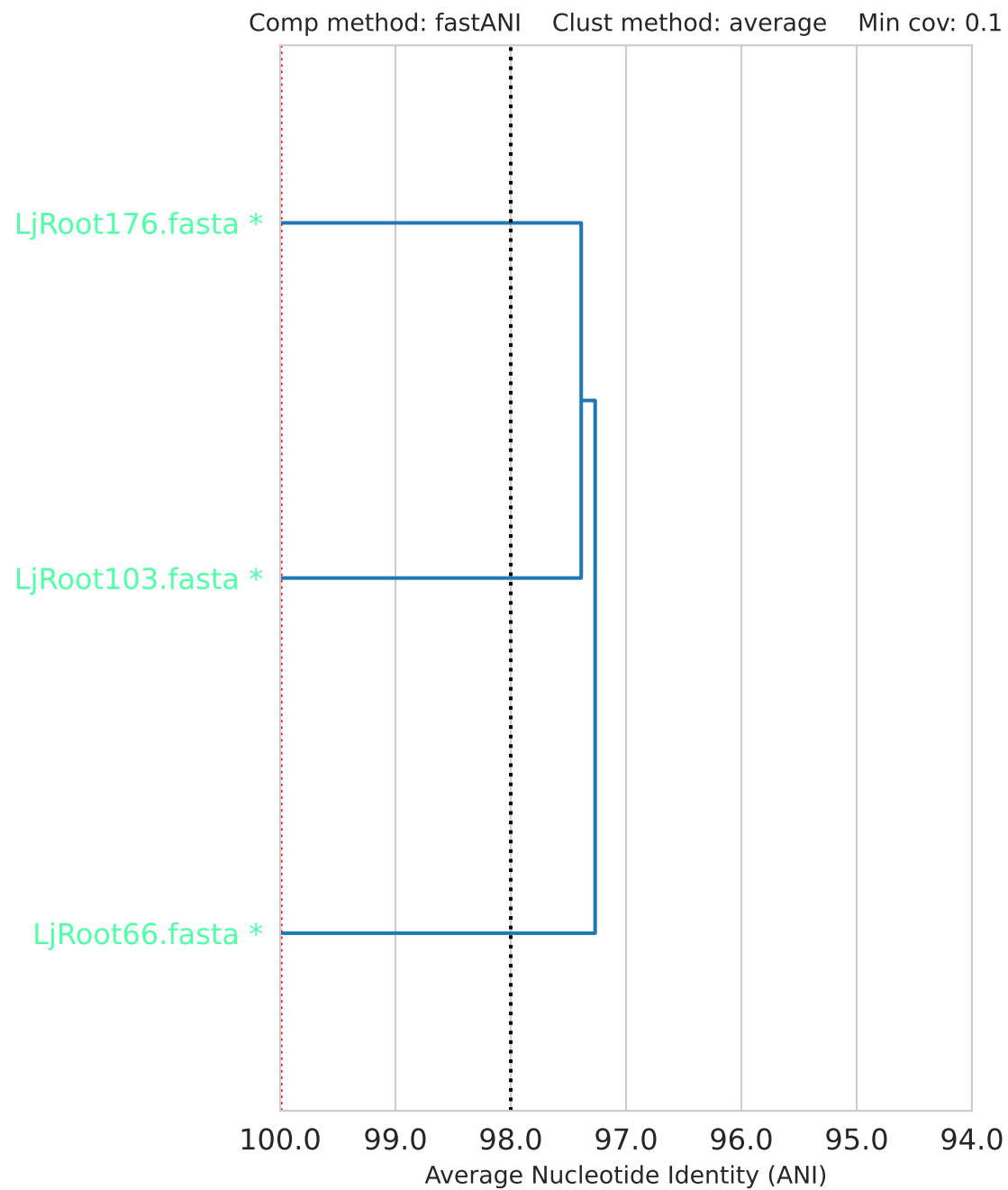
Primary cluster 230



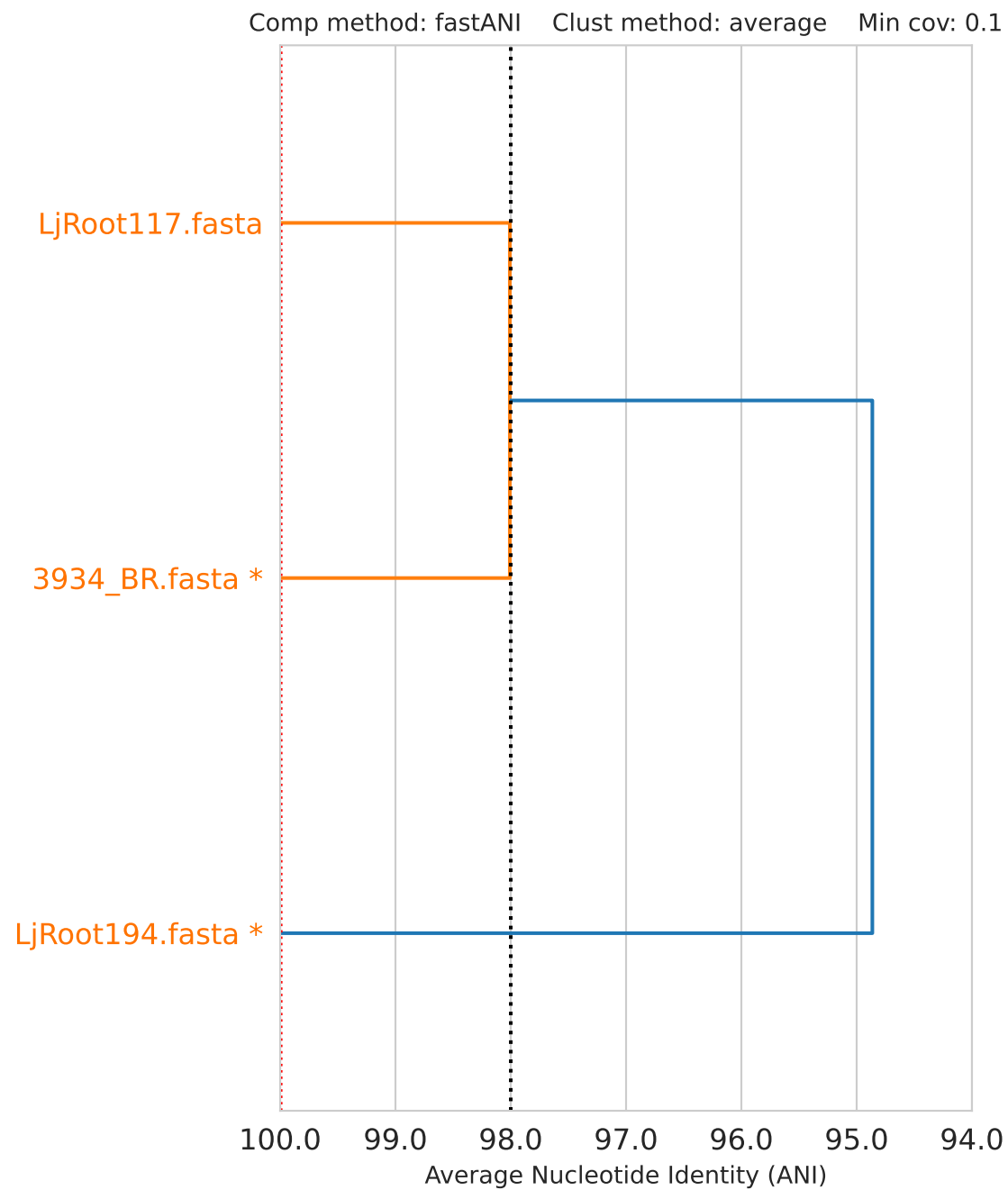
Primary cluster 232



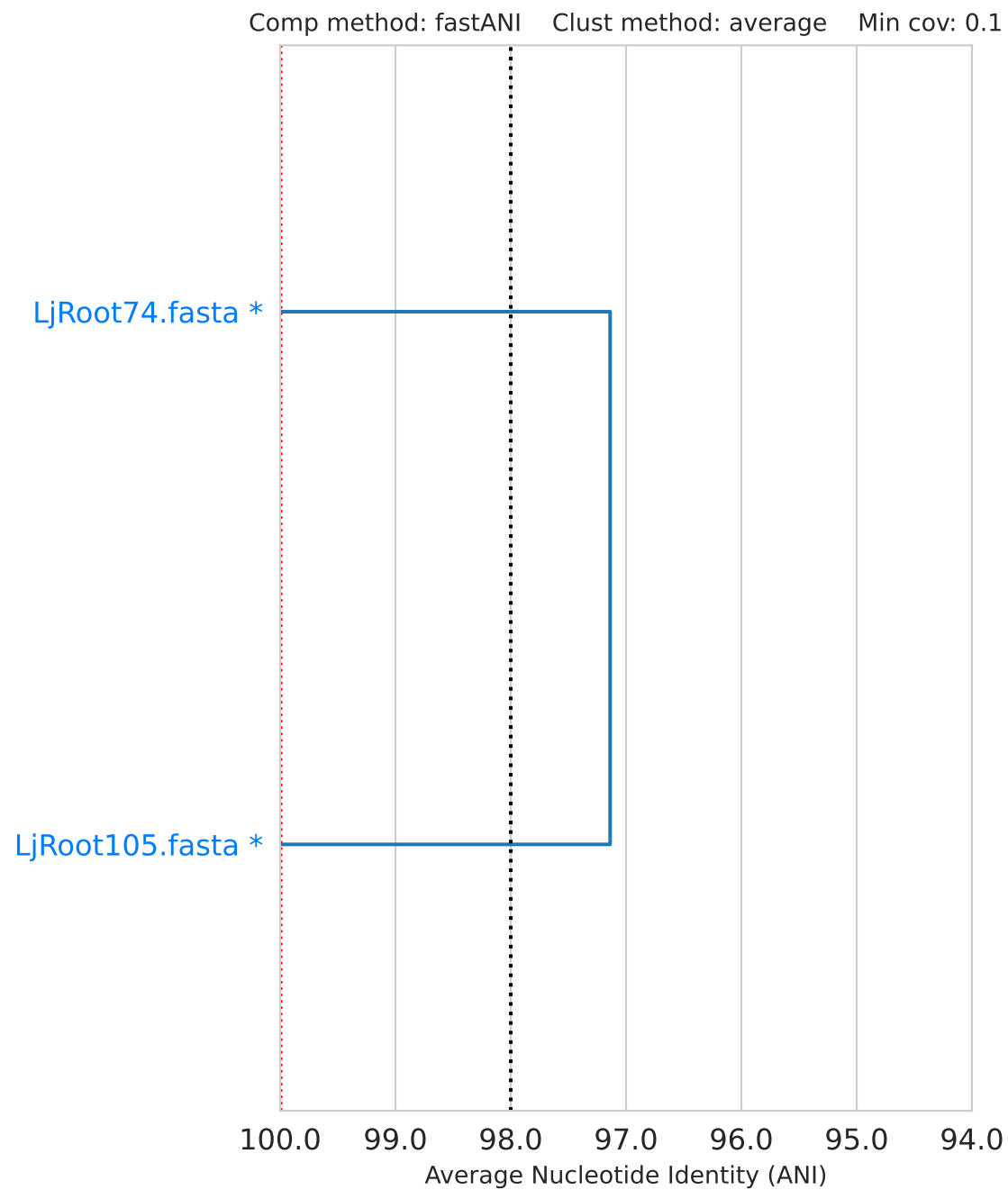
Primary cluster 236



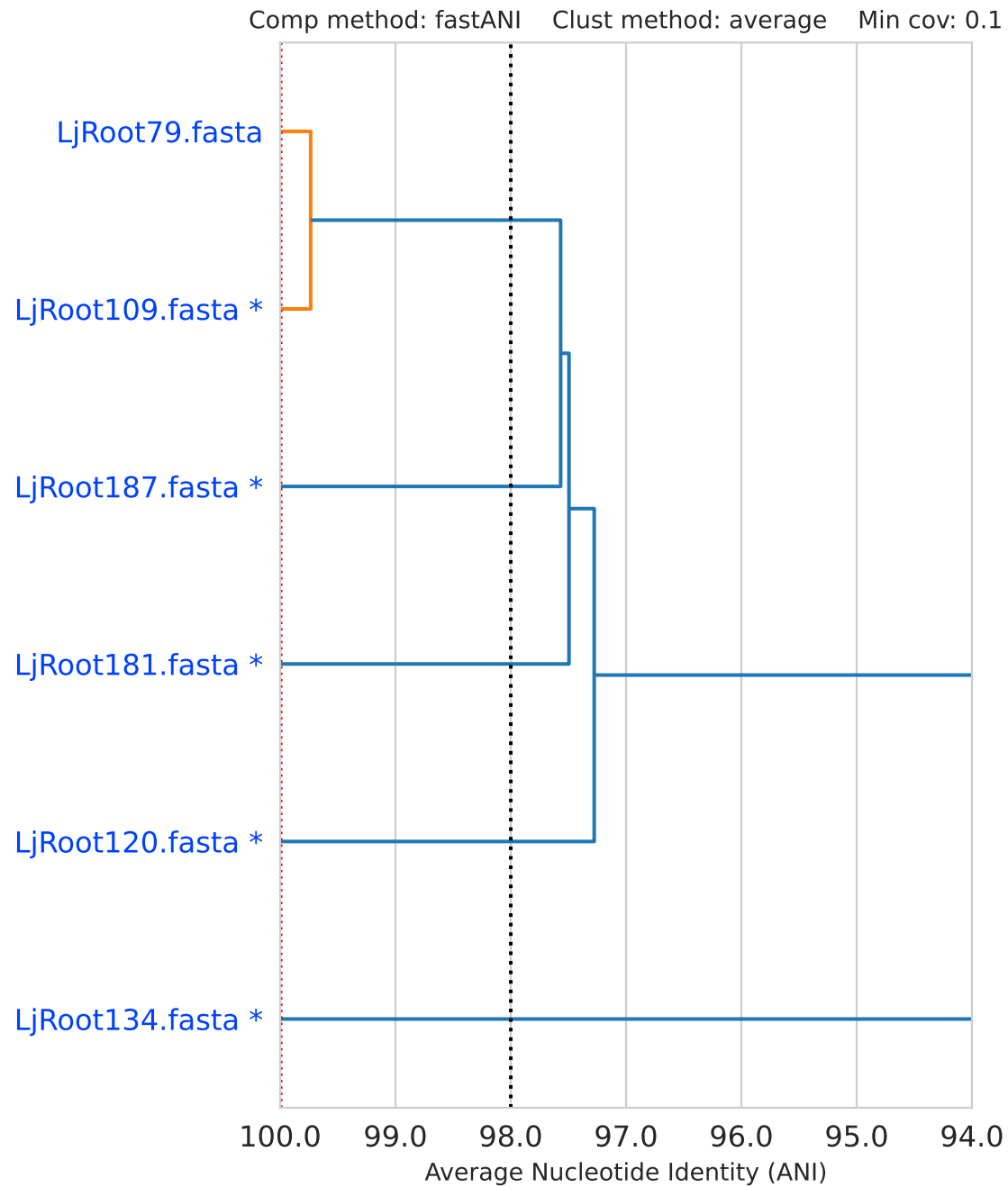
Primary cluster 237



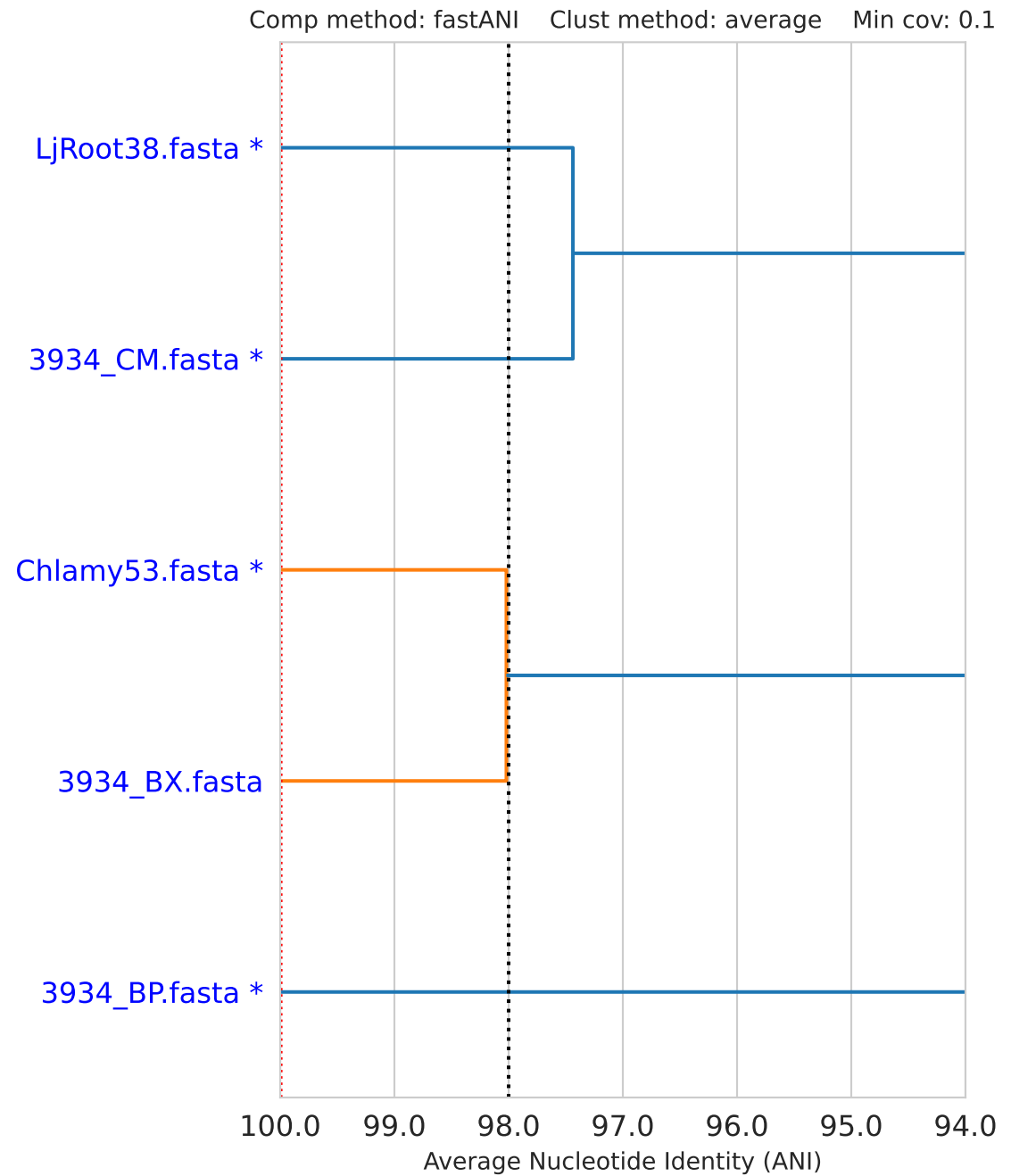
Primary cluster 239



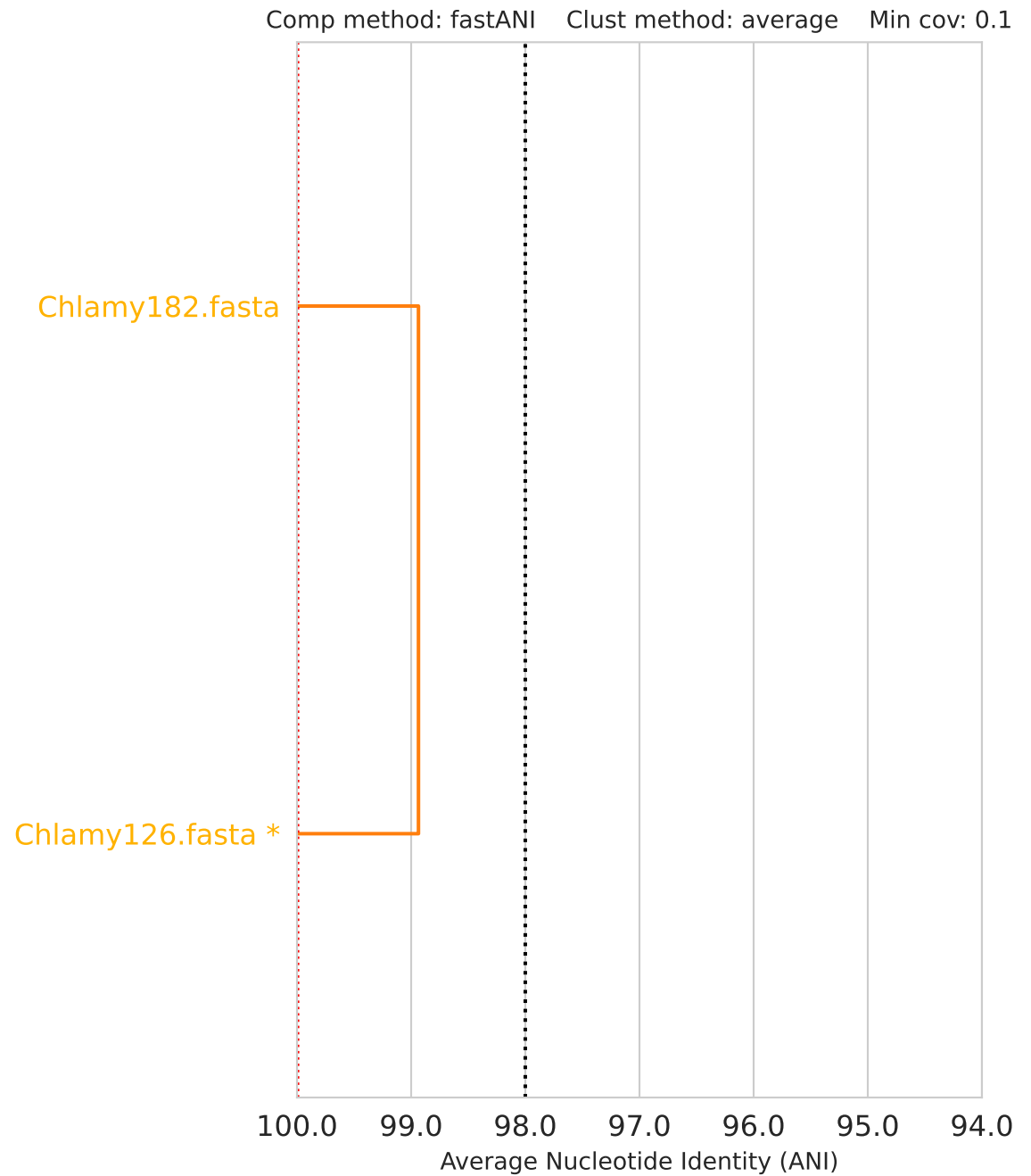
Primary cluster 240



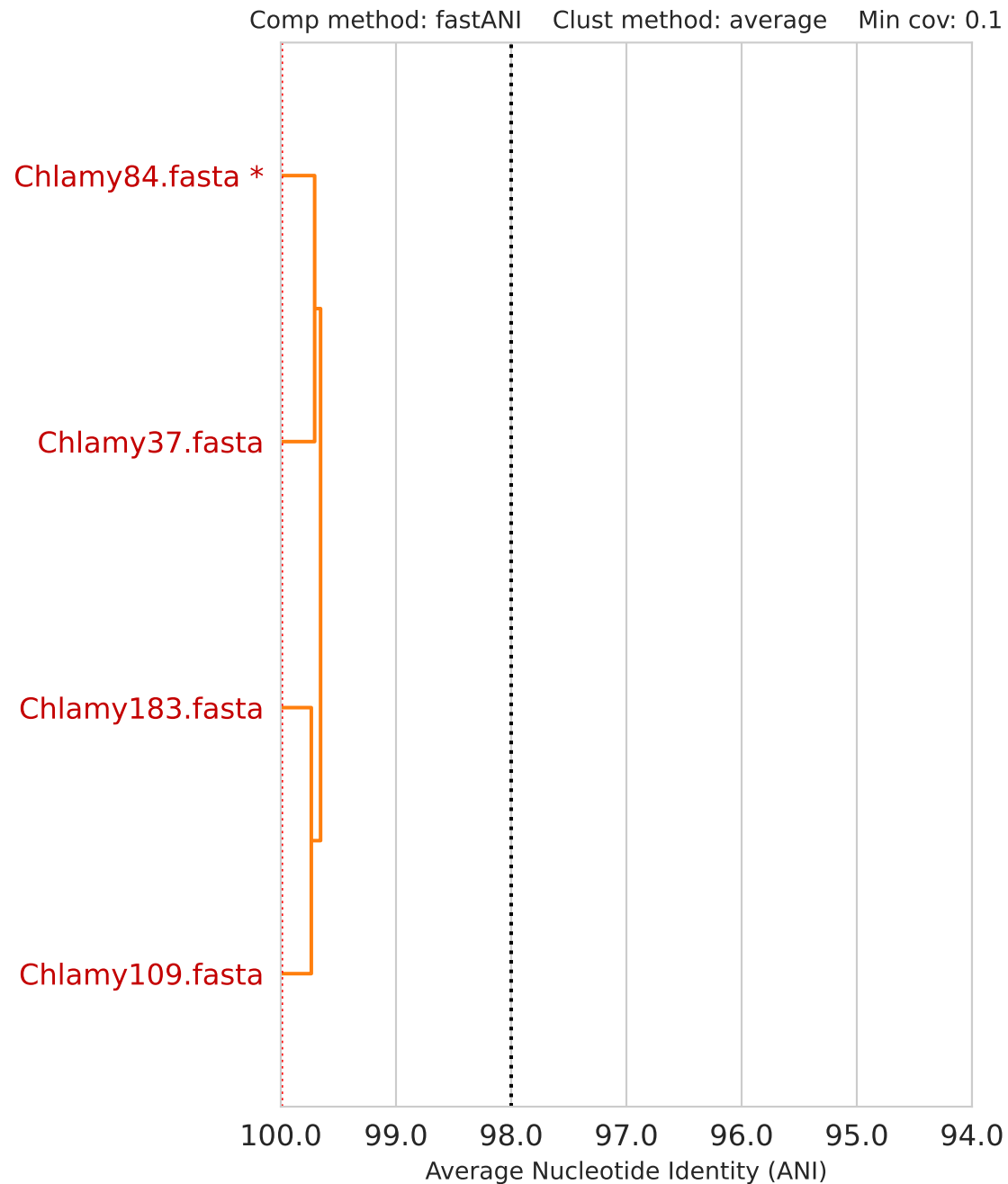
Primary cluster 241



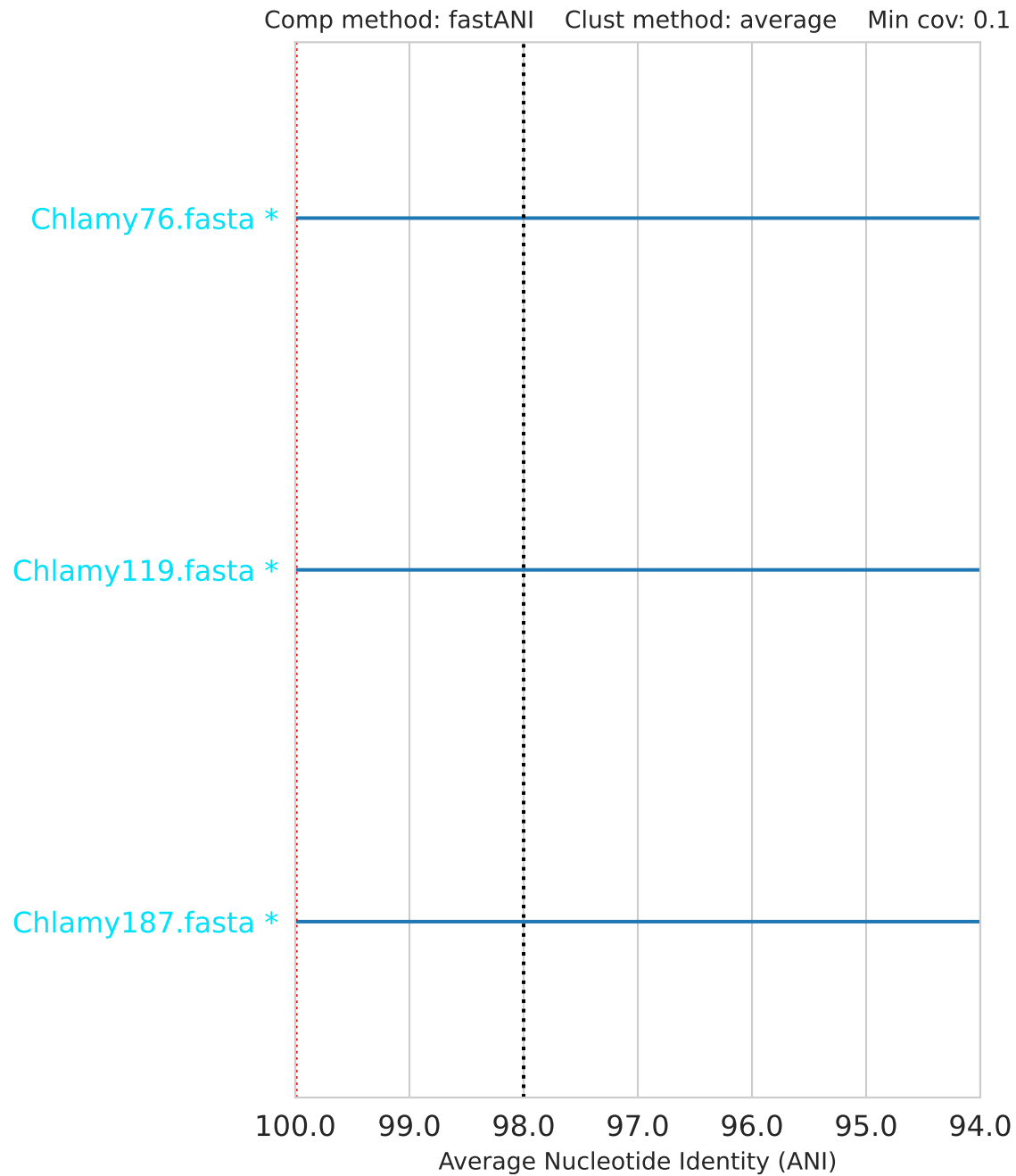
Primary cluster 245



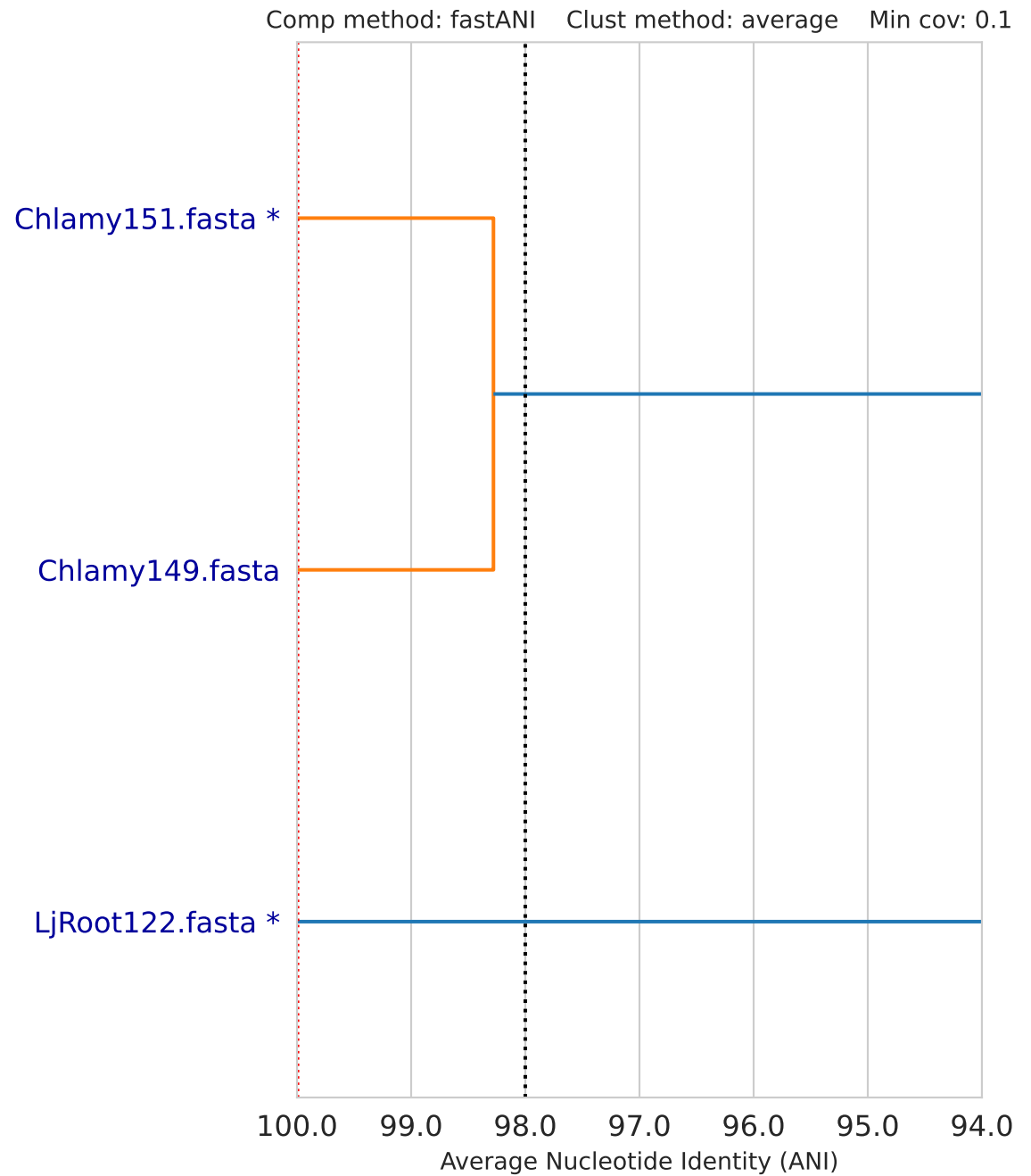
Primary cluster 246



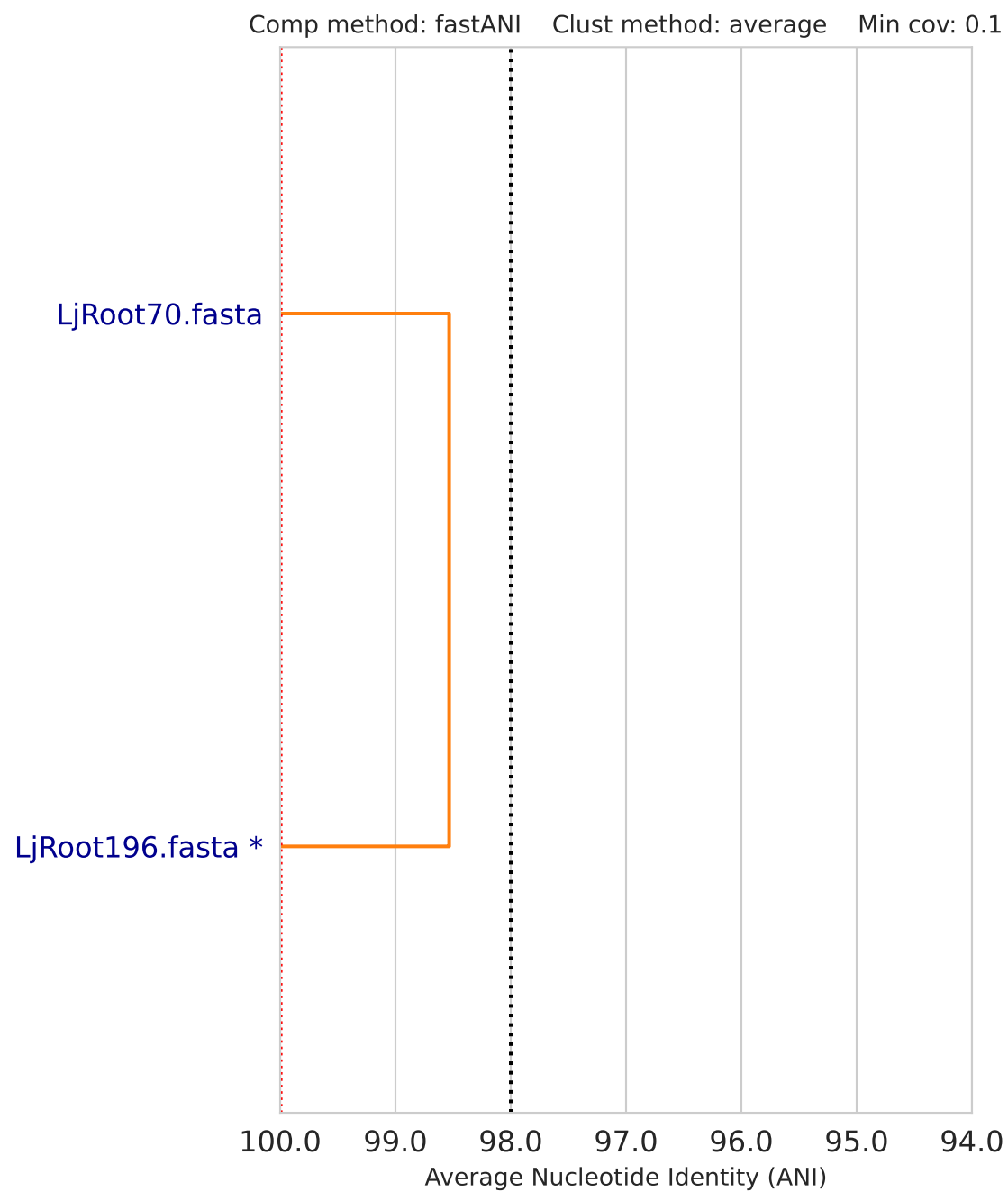
Primary cluster 247



Primary cluster 250



Primary cluster 251

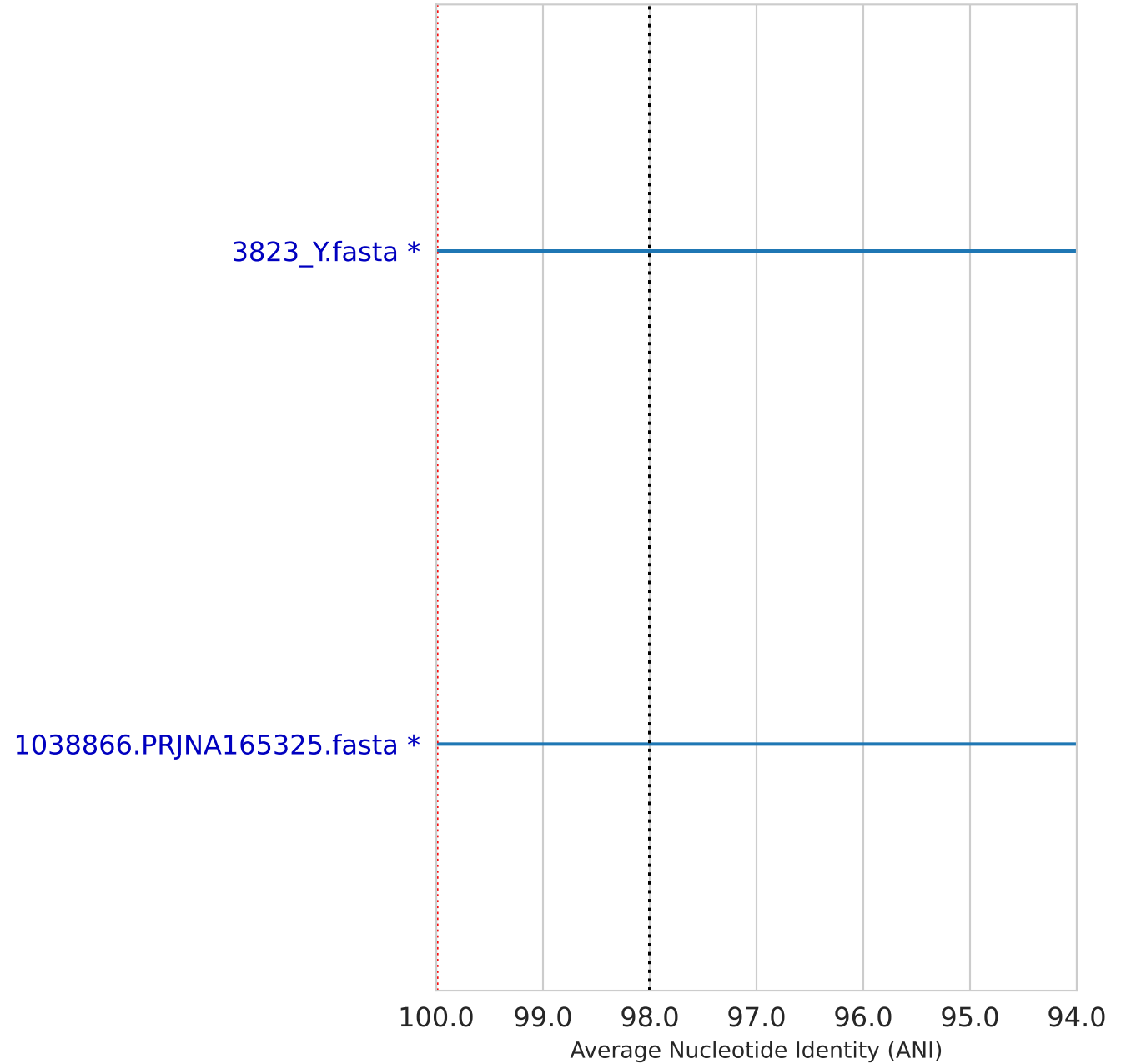


Primary cluster 270

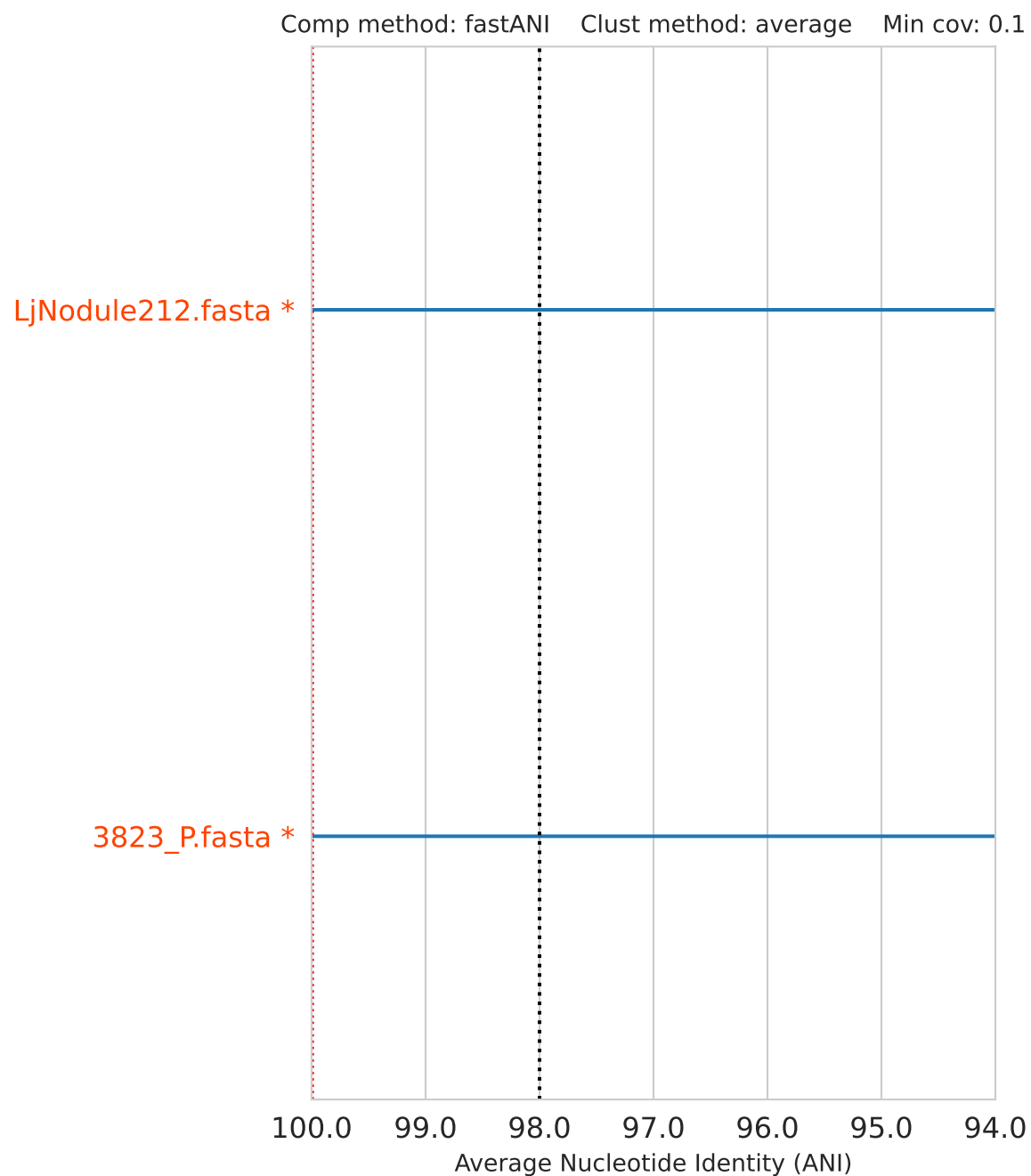
Comp method: fastANI

Clust method: average

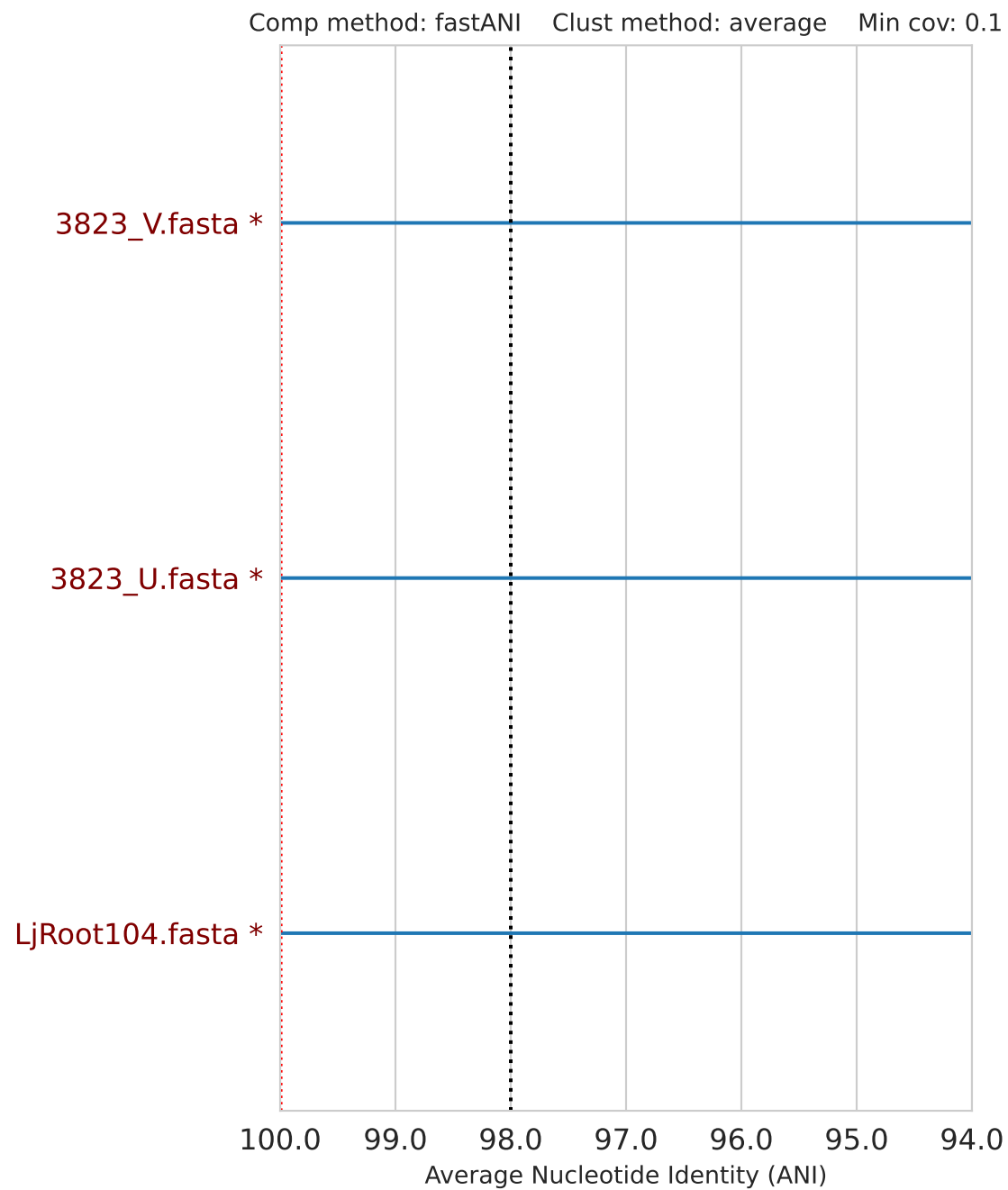
Min cov: 0.1



Primary cluster 284



Primary cluster 287

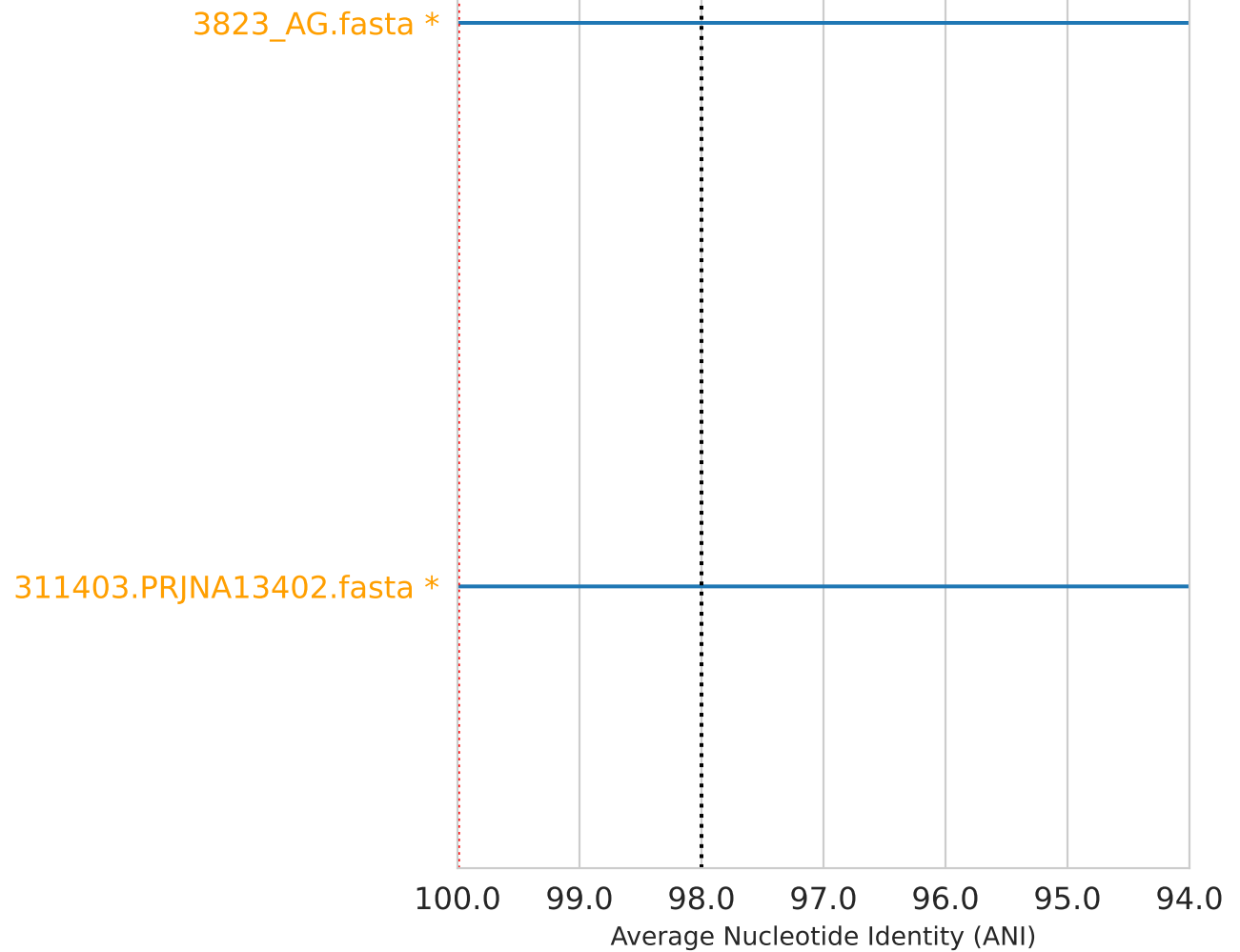


Primary cluster 289

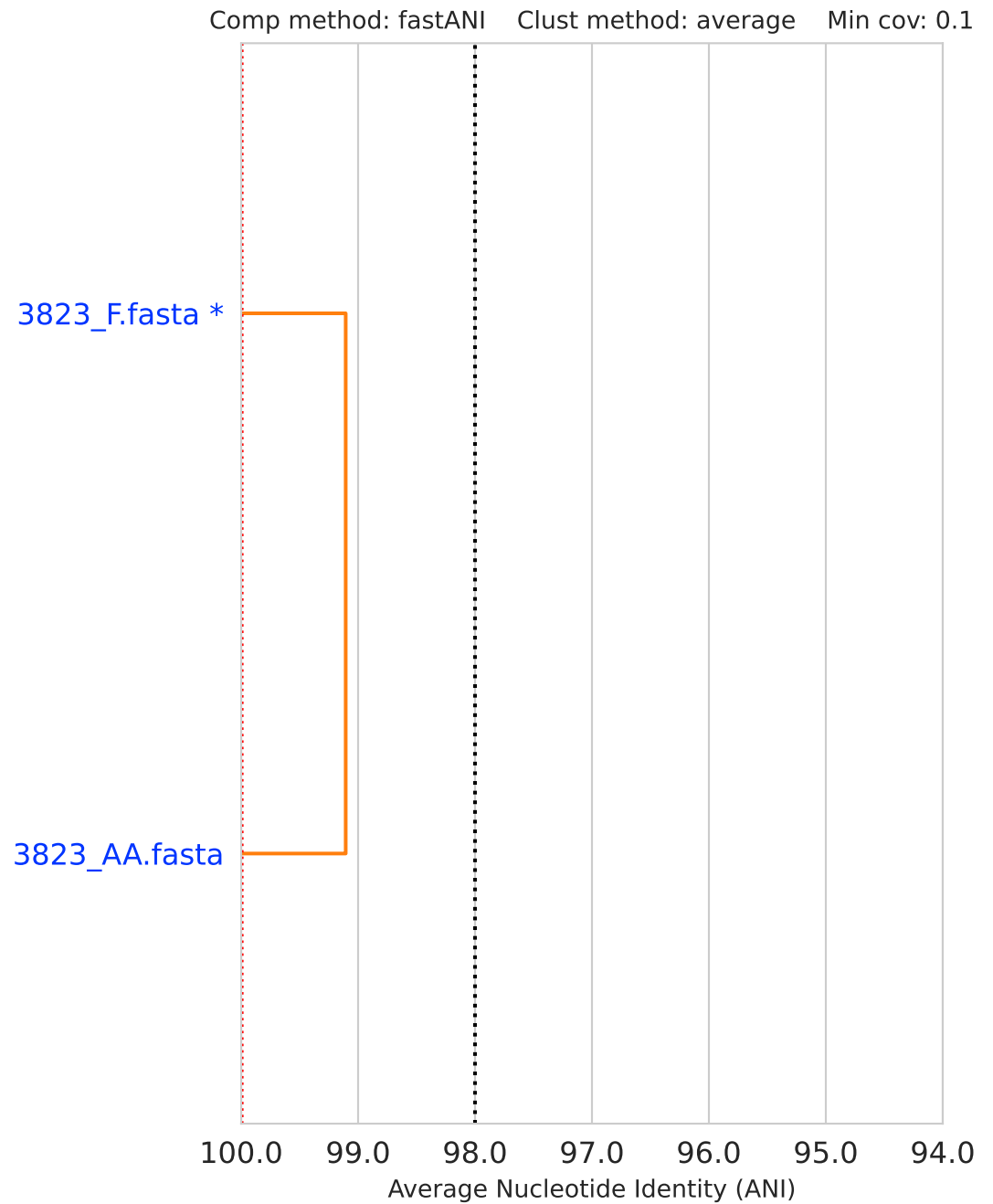
Comp method: fastANI

Clust method: average

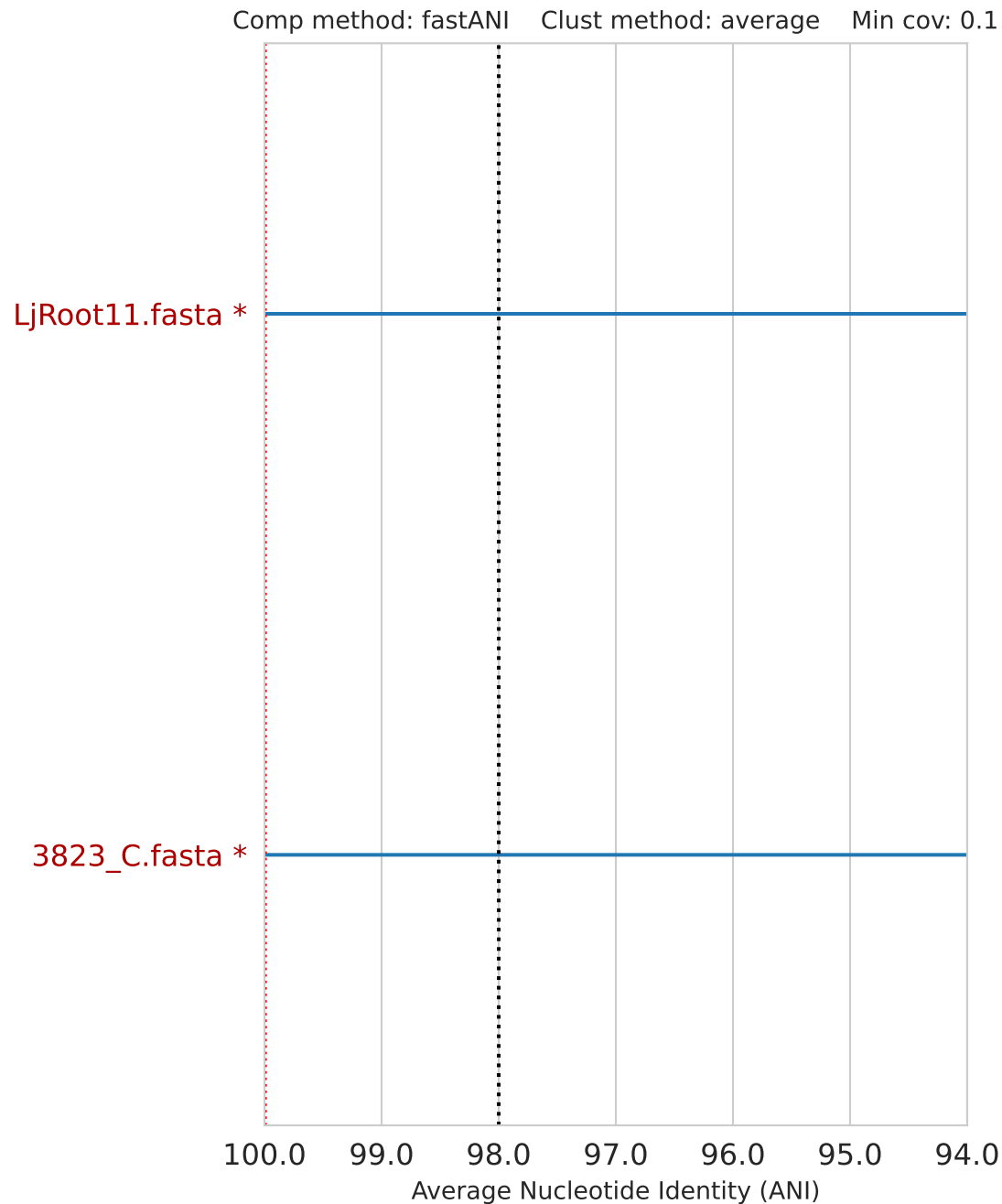
Min cov: 0.1



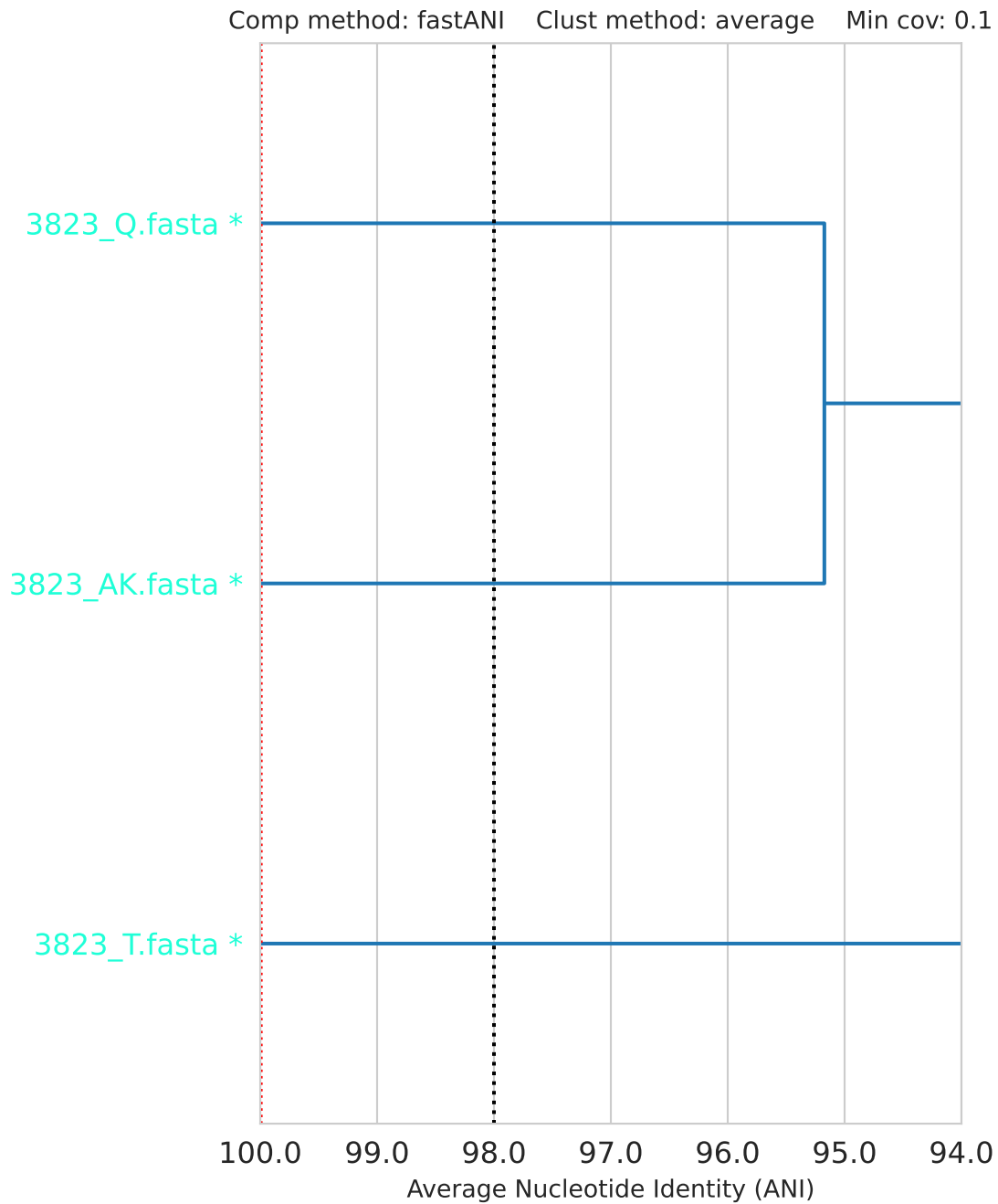
Primary cluster 290



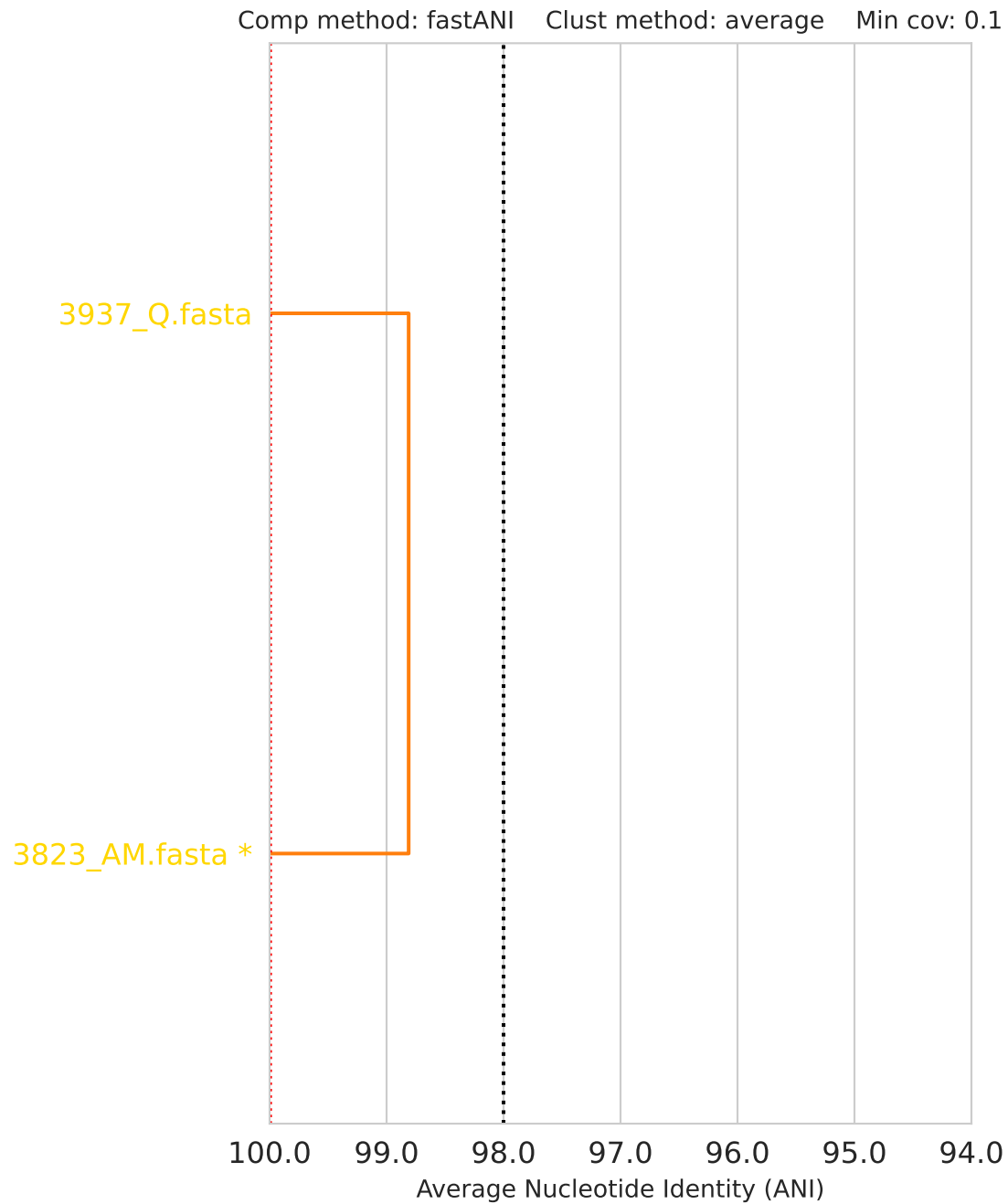
Primary cluster 295



Primary cluster 299



Primary cluster 300

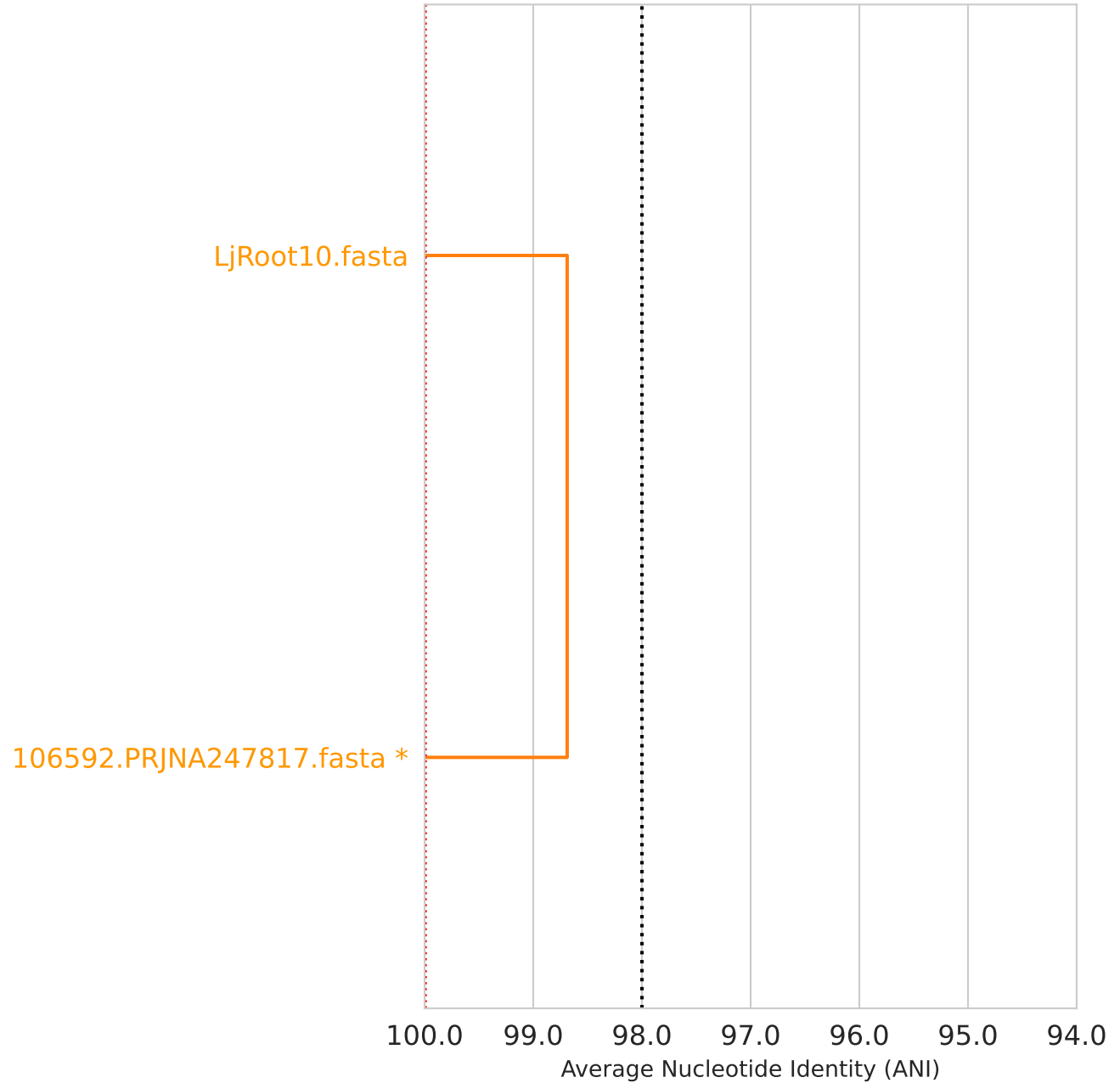


Primary cluster 301

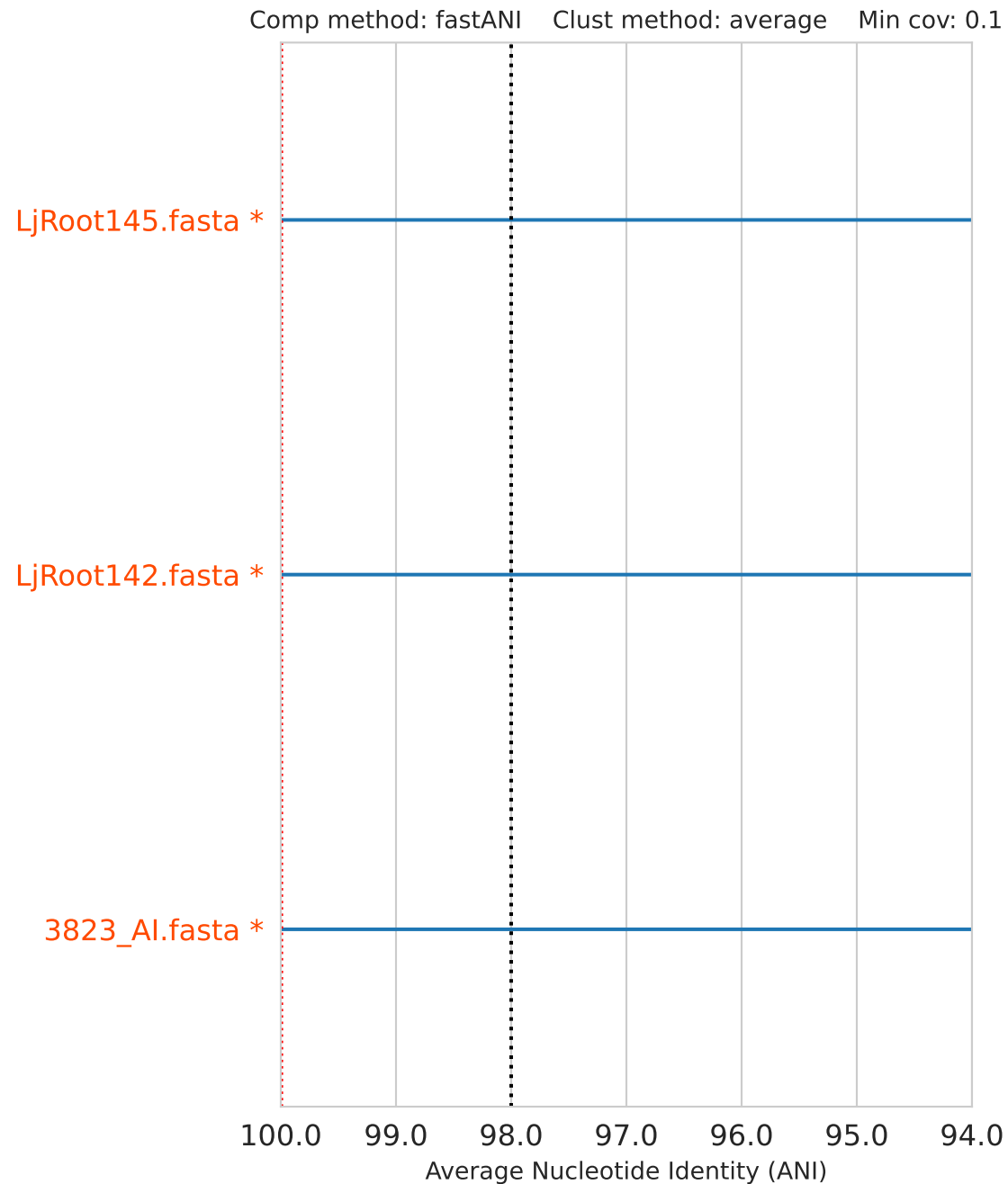
Comp method: fastANI

Clust method: average

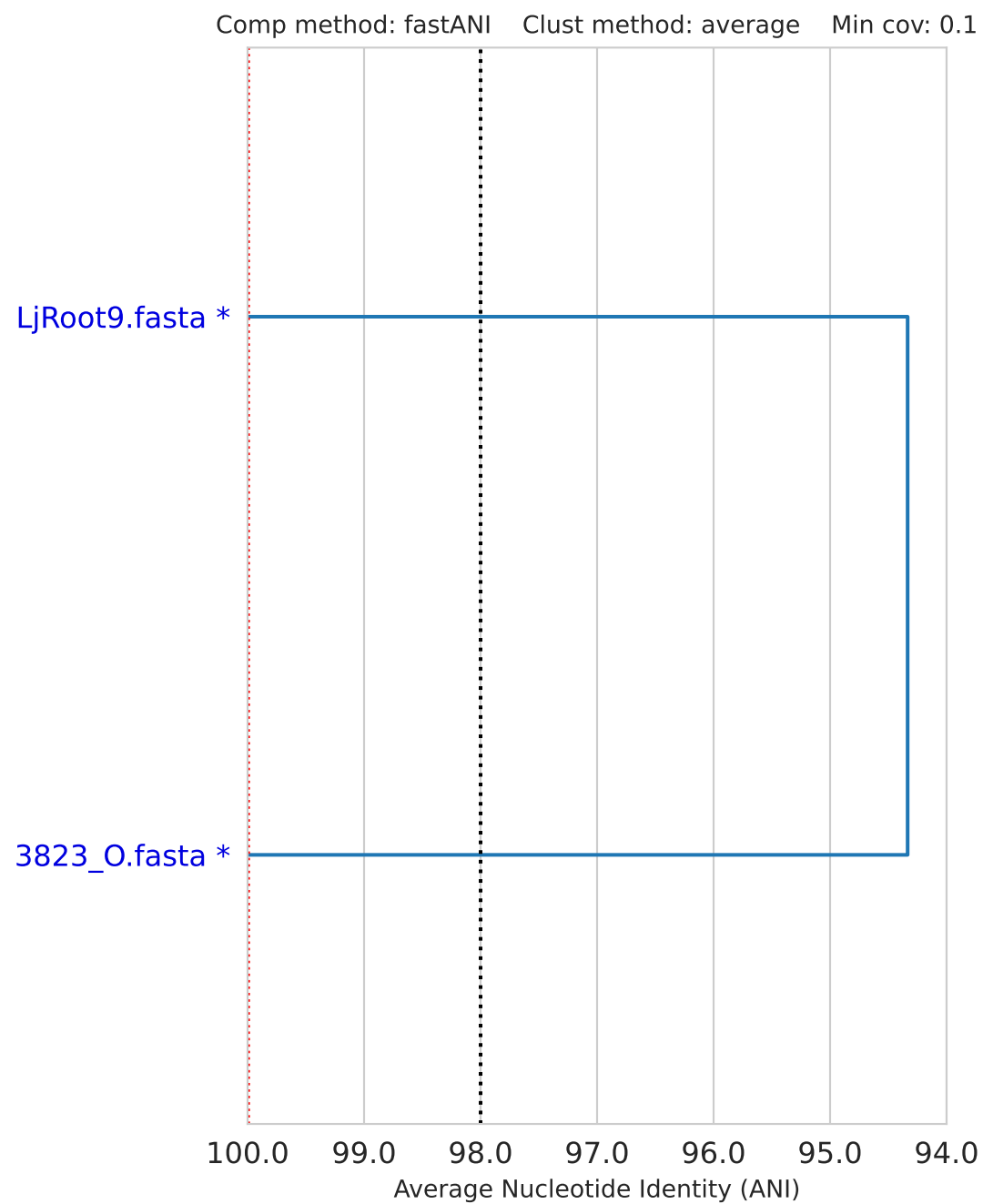
Min cov: 0.1



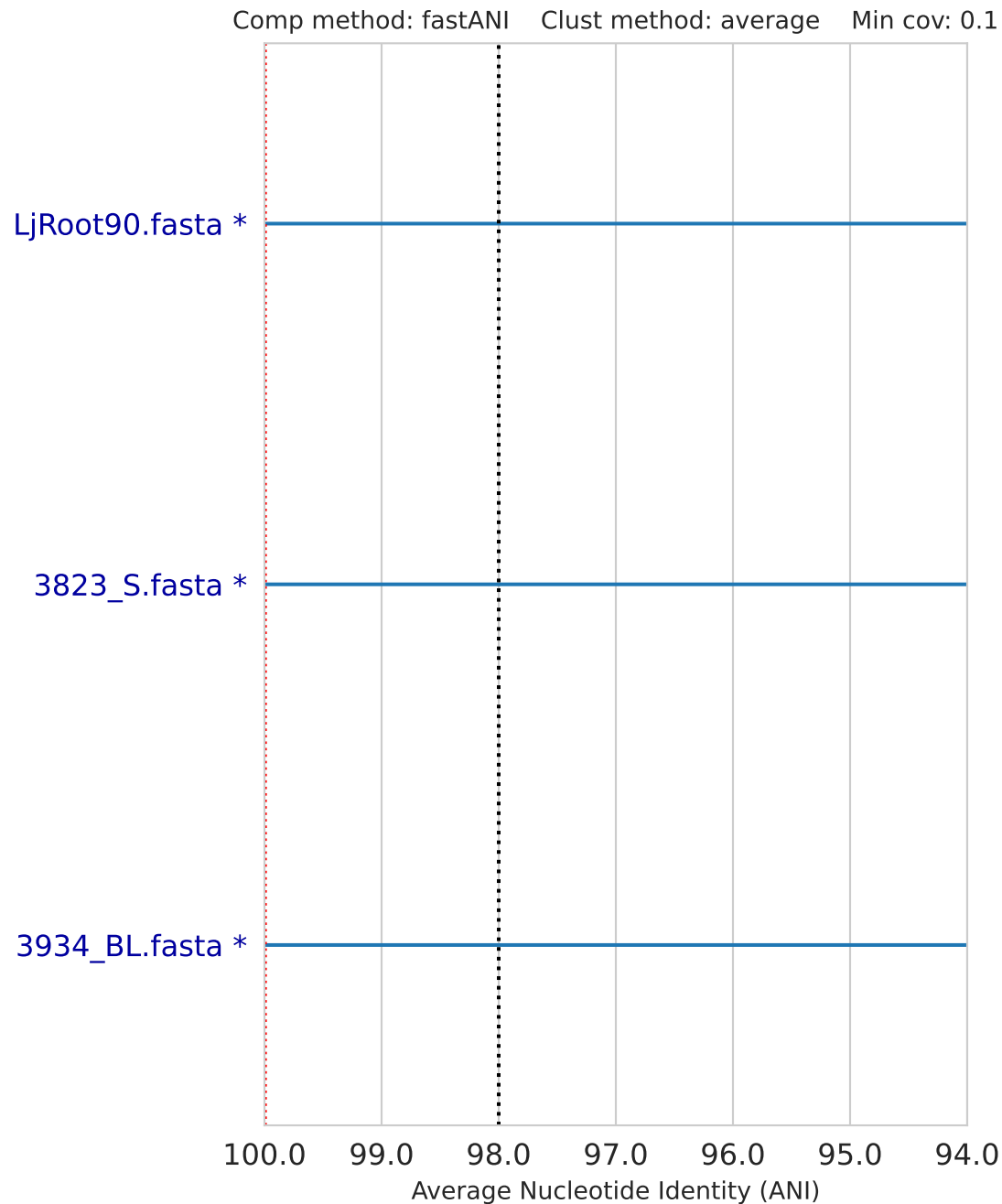
Primary cluster 302



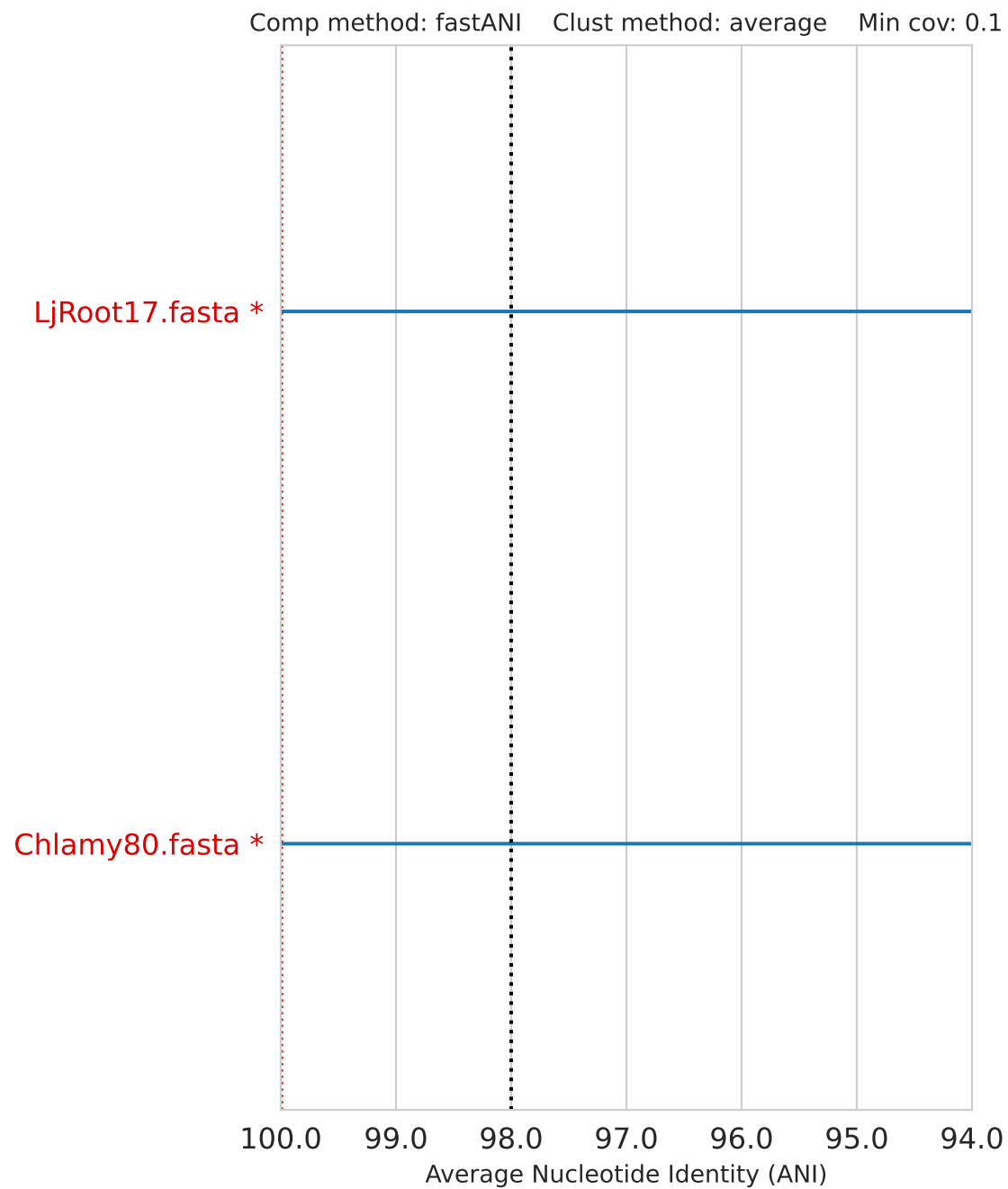
Primary cluster 303



Primary cluster 304



Primary cluster 307



Primary cluster 309

