

Primary cluster 1

Comp method: fastANI

Clust method: average

Min cov: 0.1

Lactobacillus_fermentum_complete_genome.fa.gz *

2M_ss_metabat2_bin.9.fa.gz

SRR12324251_bin.3.fa.gz

FULL_concoct_bin.2.fa.gz

1M_ss_maxbin2_bin.0.fa.gz

1M_ss_concoct_bin.0.fa.gz

1M_ss_metabat2_bin.4.fa.gz

SRR12324251-1Mss_bin.3.fa.gz

5M_ss_maxbin2_bin.0.fa.gz

5M_ss_concoct_bin.6.fa.gz

SRR12324251-5Mss_bin.5.fa.gz

2M_ss_maxbin2_bin.0.fa.gz

2M_ss_concoct_bin.15.fa.gz

SRR12324251-2Mss_bin.2.fa.gz

FULL_maxbin2_bin.2.fa.gz

100.0

99.0

98.0

97.0

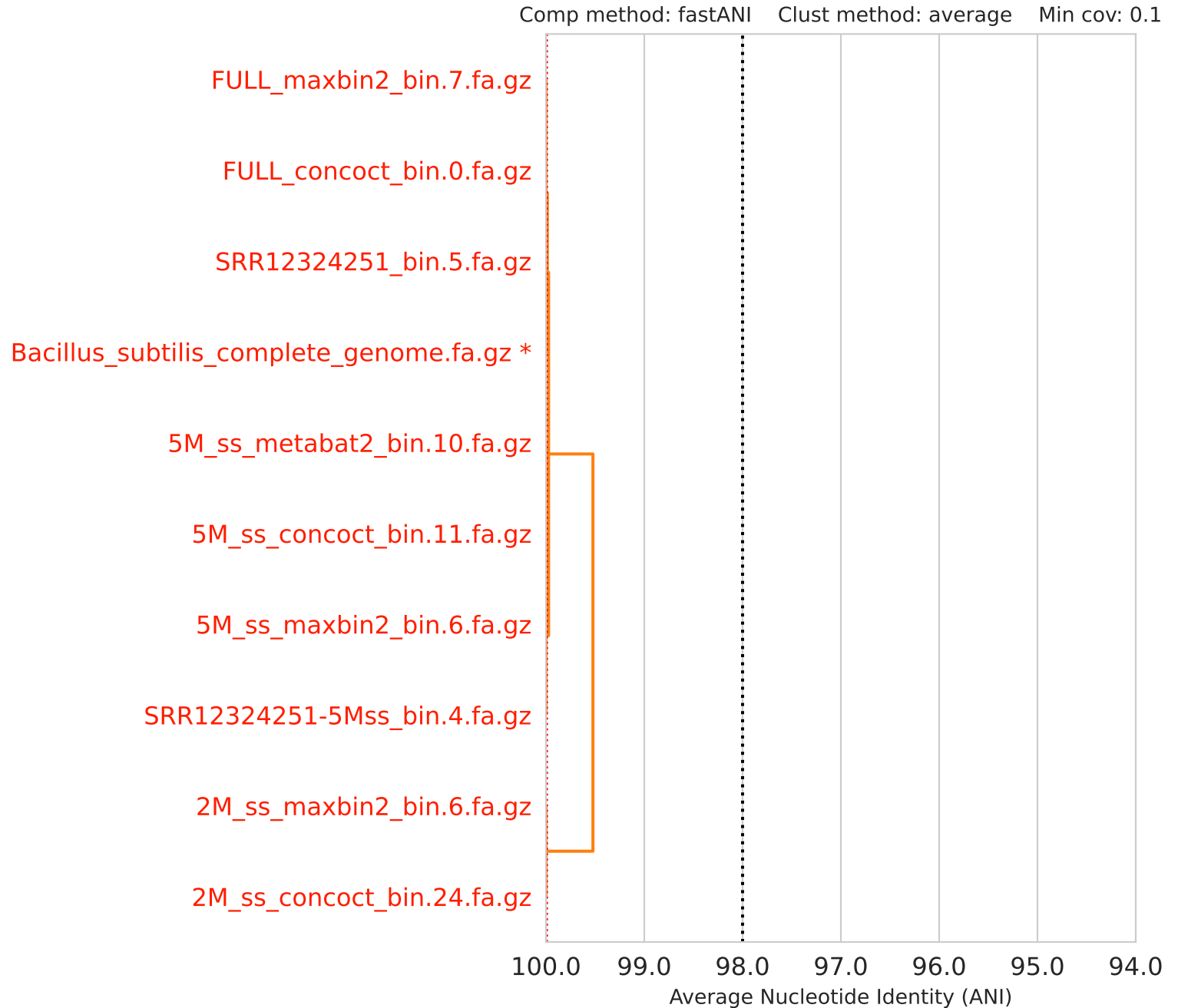
96.0

95.0

94.0

Average Nucleotide Identity (ANI)

Primary cluster 2



Primary cluster 3

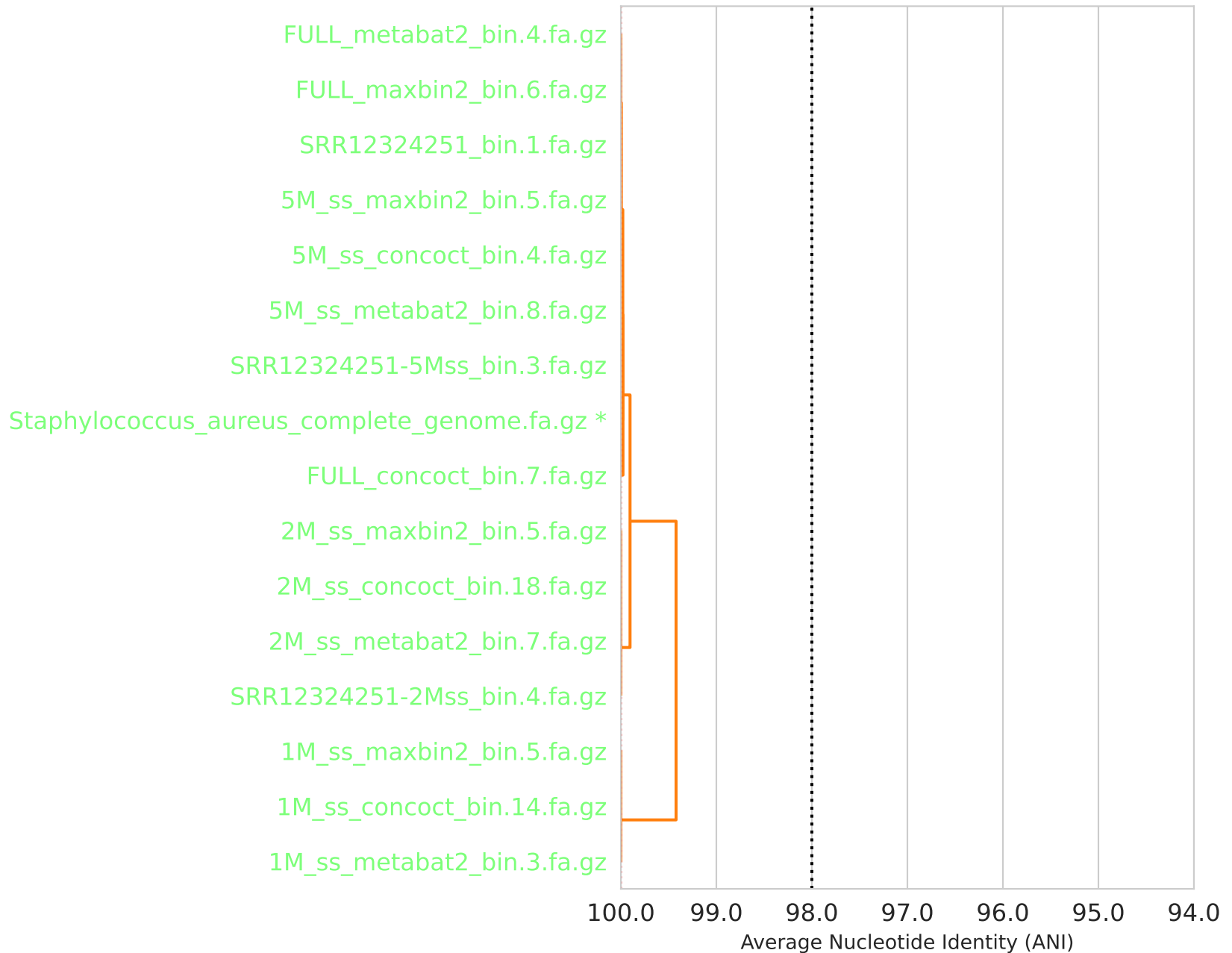


Primary cluster 4

Comp method: fastANI

Clust method: average

Min cov: 0.1

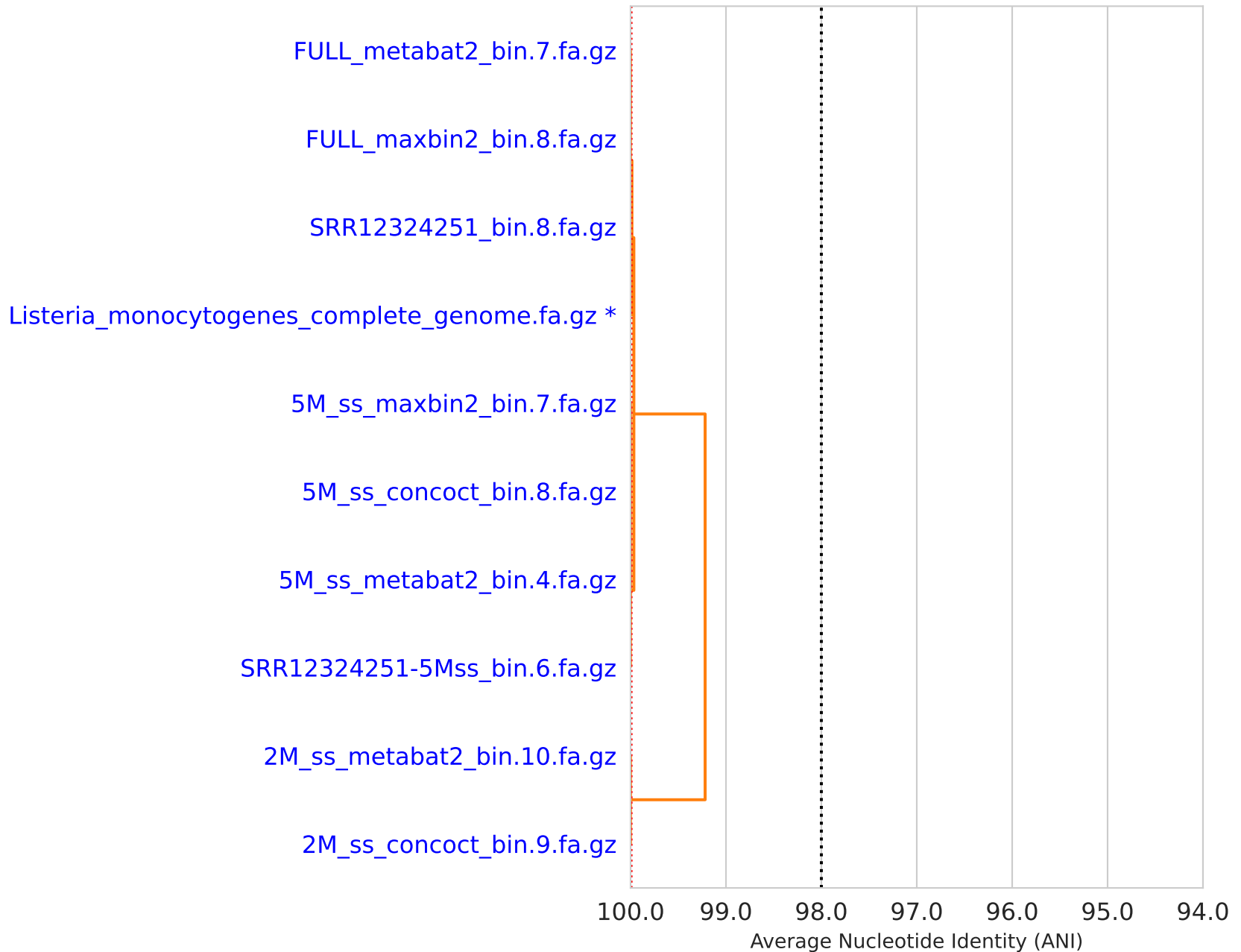


Primary cluster 5

Comp method: fastANI

Clust method: average

Min cov: 0.1

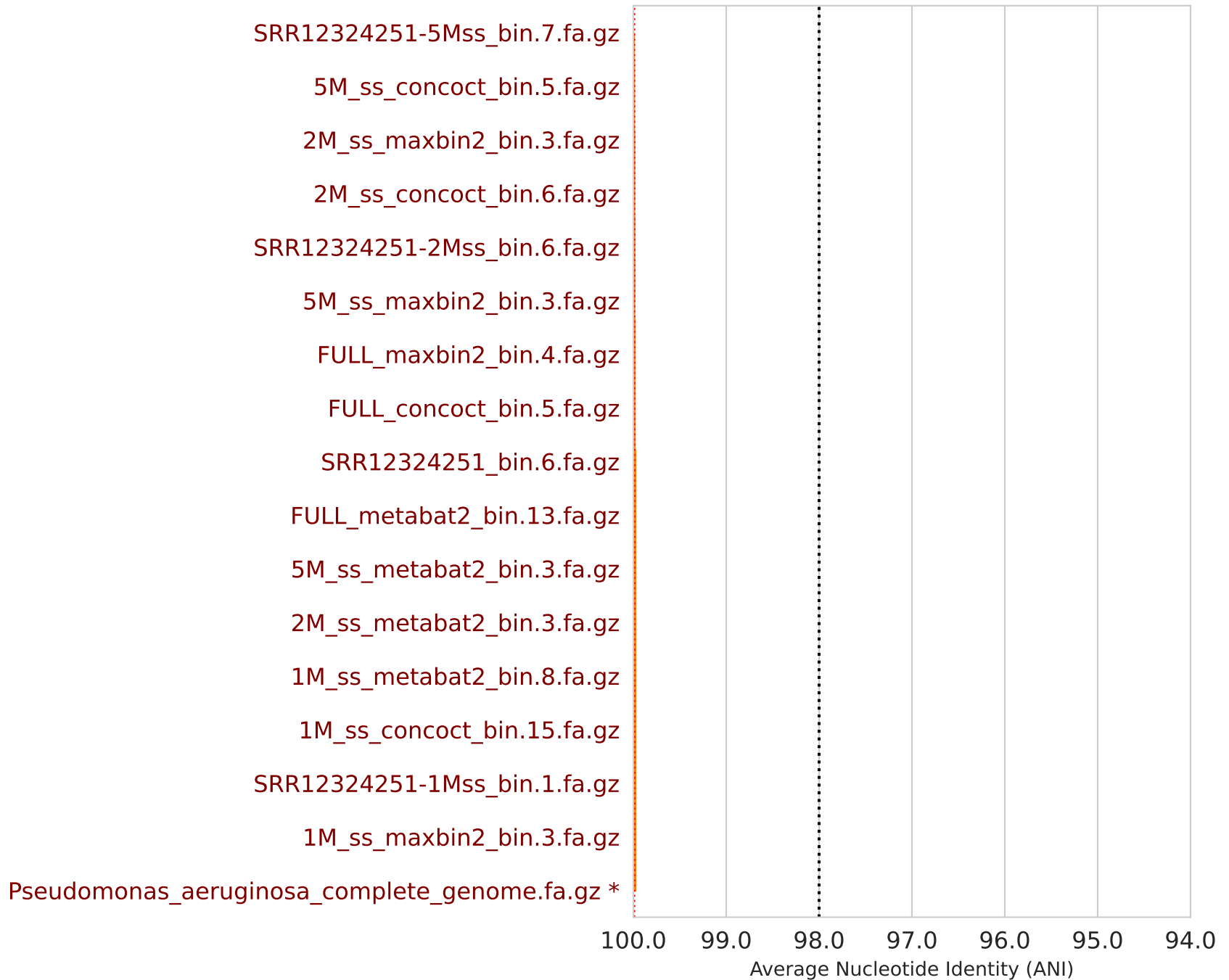


Primary cluster 6

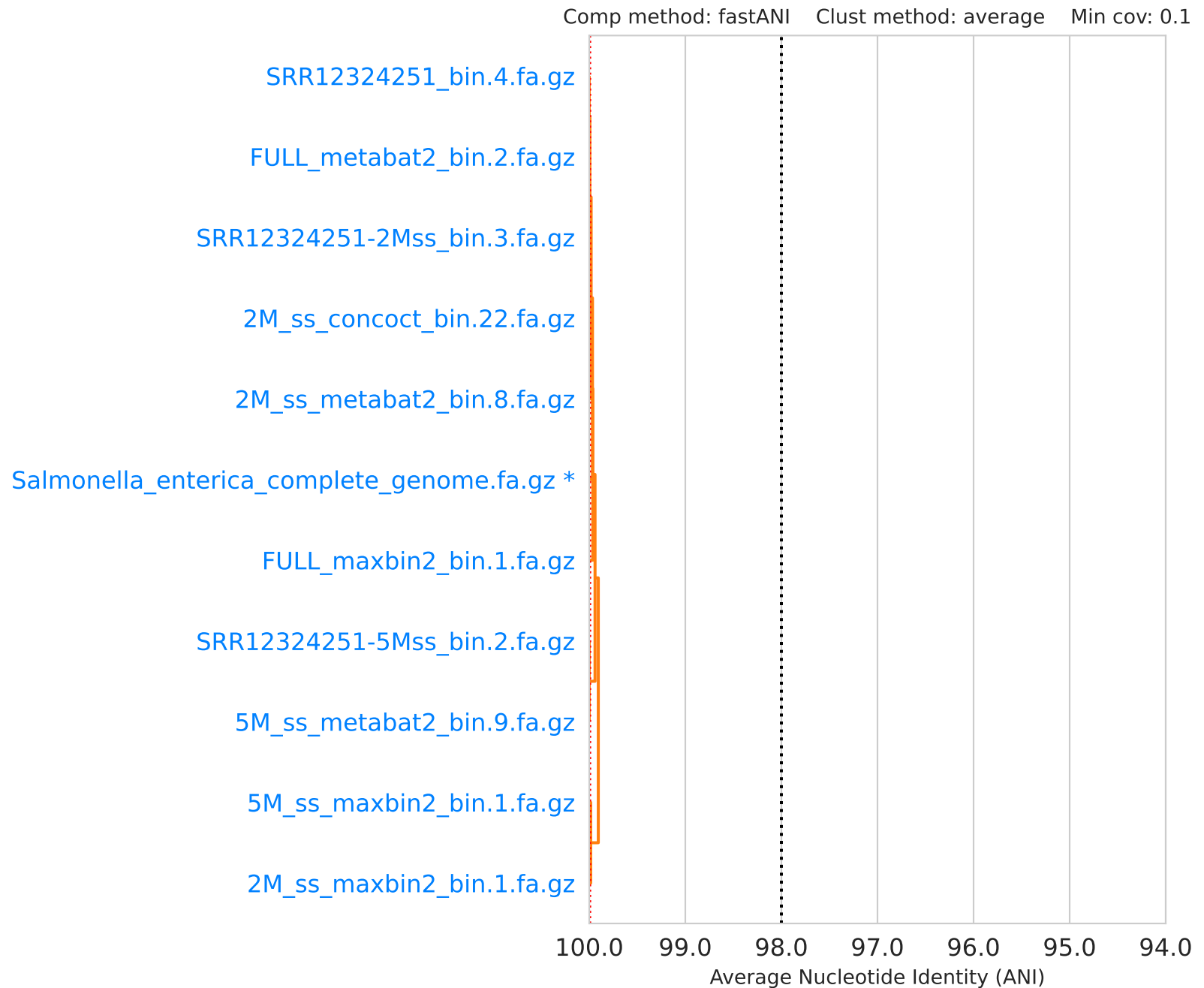
Comp method: fastANI

Clust method: average

Min cov: 0.1



Primary cluster 7



Primary cluster 8

