Primary cluster 1

Comp method: ANImf   Clust method: average   Min cov: 0.1

37-2-1__SURF-B.MEGAHIT.metabat.CheckM__Bin.044.fasta

7-16-1__SURF-B.MEGAHIT.maxbin.CheckM__Bin.009.fasta *

Average Nucleotide Identity (ANI)

100.0   99.5   99.0   98.5   98.0   97.5   97.0

Primary cluster 4

Comp method: ANImf   Clust method: average   Min cov: 0.1

37-2-1__SURF-B.MEGAHIT.metabat.CheckM__Bin.018.fasta

7-16-1__SURF-B.MEGAHIT.maxbin.CheckM__Bin.012.fasta *

100.0   99.5   99.0   98.5   98.0   97.5   97.0

Average Nucleotide Identity (ANI)

Primary cluster 5

Comp method: ANImf   Clust method: average   Min cov: 0.1

37-2-1__SURF-B.MEGAHIT.metabat.CheckM__Bin.039.fasta

7-16-1__SURF-B.MEGAHIT.maxbin.CheckM__Bin.008.fasta *

100.0   99.5   99.0   98.5   98.0   97.5   97.0

Average Nucleotide Identity (ANI)