

MASH clustering

sequence_000001.fasta.fasta (2_0)

sequence_000000.fasta.fasta (1_0)

sequence_000002.fasta.fasta (3_0)

sequence_000003.fasta.fasta (4_0)

100.0

80.0

60.0

40.0

20.0

0.0

MASH Average Nucleotide Identity (ANI)

