

MASH clustering

sequence_000002.fasta.fasta (2_0)

sequence_000001.fasta.fasta (1_0)

sequence_000000.fasta.fasta (3_0)

100.0

97.5

95.0

92.5

90.0

87.5

85.0

82.5

MASH Average Nucleotide Identity (ANI)

