

Breadth of genome covered at least 1X (%)

breadth_1X

100
75
50
25
0

GCF_000194945.1
GCF_000212855.1
GCF_000212815.1
GCF_001078705.1
GCF_003943735.1
GCF_003943655.1
GCF_900635155.1
GCF_013343115.1
GCA_013378335.1
GCF_000767835.1
GCF_001578725.1
GCF_019090945.1
GCF_902460385.1

Assembly accession

breadth_5X

GCF_000194945.1
GCF_000212855.1
GCF_000212815.1
GCF_001078705.1
GCF_003943735.1
GCF_003943655.1
GCF_900635155.1
GCF_013343115.1
GCA_013378335.1
GCF_000767835.1
GCF_001578725.1
GCF_019090945.1
GCF_902460385.1

species

- × sanguinis
- sinensis

GTDB_species

- sanguinis
- sanguinis_A
- sanguinis_C
- sanguinis_D
- sanguinis_E
- sanguinis_F
- sanguinis_G
- sanguinis_H
- sanguinis_I
- sinensis