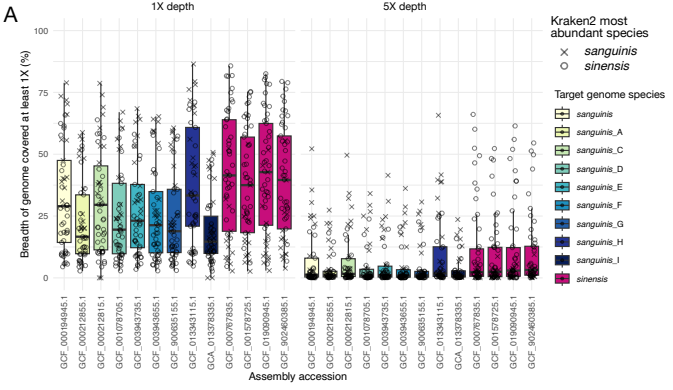
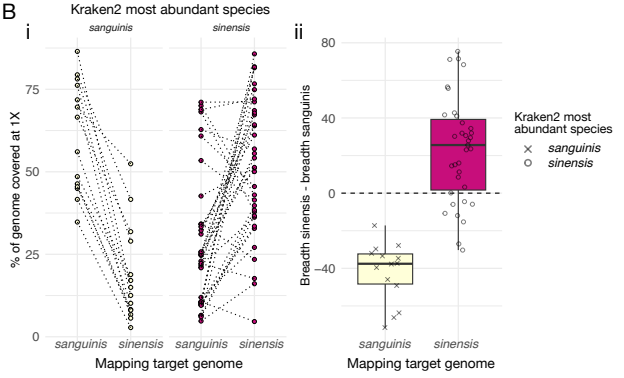
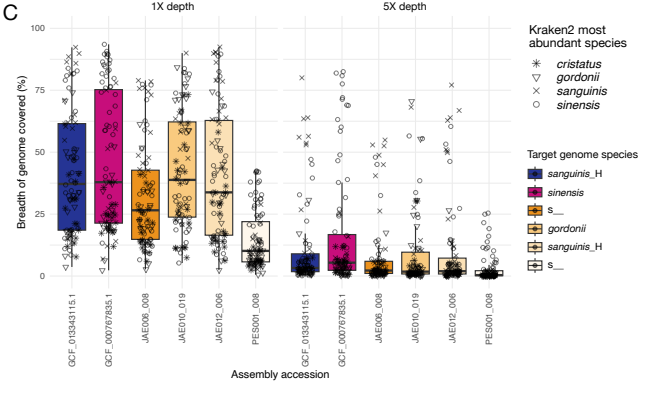

A
1X depth
5X depth
Breadth of genome covered at least 1X (%)
Assembly accession
Kraken2 most abundant species
sanguinis
sinensis
Target genome species
sanguinis
sanguinis_A
sanguinis_C
sanguinis_D
sanguinis_E
sanguinis_F
sanguinis_G
sanguinis_H
sanguinis_I
sinensis


B
Kraken2 most abundant species
i
sanguinis
sinensis
% of genome covered at 1X
Mapping target genome
ii
Breadth sinensis - breadth sanguinis
Kraken2 most abundant species
sanguinis
sinensis
Mapping target genome


C
1X depth
5X depth
Breadth of genome covered (%)
Assembly accession
Kraken2 most abundant species
cristatus
gordonii
sanguinis
sinensis
Target genome species
sanguinis_H
sinensis
s__
gordonii
sanguinis_H
s__