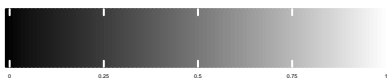


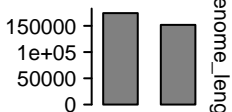
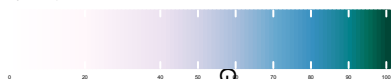
Aligned genome fraction



Genome length ratio



Intergenomic similarity



100.0	93.2
1.0	
0.9	100.0
0.9	

Genome\_length

KY945241

MH107246

KY945241

MH107246