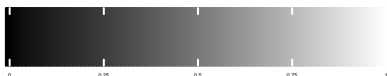


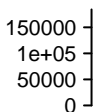
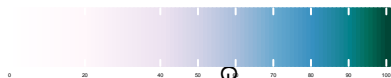
Aligned genome fraction



Genome length ratio



Intergenomic similarity



Genome\_length

100.0 99.8

GU071106

1.0

1.0

1.0

100.0

HQ634190

GU071106

HQ634190