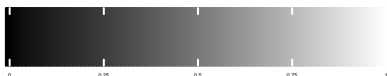


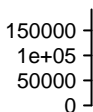
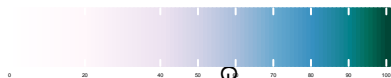
Aligned genome fraction



Genome length ratio



Intergenomic similarity



Genome_length

100.0 100.0

HQ316583

1.0

1.0

1.0

100.0

HQ316603

HQ316583

HQ316603