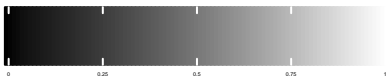


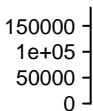
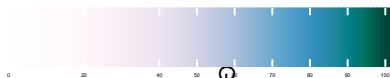
Aligned genome fraction



Genome length ratio



Intergenomic similarity



Genome length	
150000	150000
1e+05	1e+05
50000	50000
0	0
100.0	99.5
1.0	1.0
1.0	100.0
1.0	1.0
GU071099	GU071099
MH512890	MH512890