Aligned genome fraction
0
0.25
0.5
0.75
1
Genome length ratio
0
0.25
0.5
0.75
1
Intergenomic similarity
0
20
40
50
60
70
80
90
100
60000
40000
20000
0
Genome_length
NC_022063.1_Myco_phg_Phelemich
NC_022064.1_Myco_phg_Reprobate
KY224000.1_Myco_phg_Rich
NC_028742.1_Myco_phg_Baee
NC_051587.1_Myco_phg_Serendipitous
NC_023701.1_Myco_vir_Acadian
MK524490.1_Myco_phg_Donny
MT889381.1_Myco_phg_Suigeneris