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
Genome_length
15000
10000
5000
0
MH045556.1
MN234191.1
MH825700.1
MT316458.1
MK801730.1
MH371118.1
MH371111.1
MT498040.1
MN369745.1
MT521990.1
MN585990.1
MT639647.1
MN284892.1
MT498051.1
MH744416.1
MT639642.1
MH045561.1
MH651181.1
MK878899.1
MT498046.1
MH371124.1
MH576963.1
MT316462.1
MT316459.1
MN329679.1
MK620897.1
MK894434.1
MN428062.1
MT776807.1
MN010759.1
MN010755.1
MN586008.1
MH045557.1
MN428064.1
MK524502.1
MT498038.1
MH399783.1
MT498065.1
MH779512.1
MT310864.1
MT498047.1
MN586010.1
MT498067.1