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
2e+05
150000
1e+05
50000
0
KX452698
KJ477686
KJ477685
AF158101
KM607003
KM607002
HM137666
KJ477684
KJ668714
LR215722
KR269718
MF001354
MH837626
LR215723
MH355584
MG781190
MG781191
KY290975
KY608967
LR215724
MF327007
MH992510
KX130861
KX130862
JN986846
MH809535
KF208315
MH550421
FJ839692
KM606998
KM606996
KM606995
KM606997
KM606994
KM606999
JX128259
DQ904452
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KT184312
MF158045
LN881732
MH553563
MF327008
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MG775043
HM035025
LN881737
LN881734
LN881726
LN881729
HE956711
LN881733
LN881728
LN881736
KT184308
KT184311
KX130864
LC348380
MH751506
KU867876
JN202312
MH051915
KX009778
KF925357
KR233165
KU925172
MF158046
FJ839693
KM607004
HM997020
KP869105
MF001360
AP018932
MH992121
KM501444
KX452694
LC348379
KX828711
AP011113
KM607000
MH051916
MH243439
MF001359
MG867727