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
60000
40000
20000
0
MF140402.1_Myco_phg_Chancellor
MH779514.1_Myco_phg_Paito
KX641265.1_Myco_phg_Taheera
KX641266.1_Myco_phg_Terror
MK524494.1_Myco_phg_Rabbs
NC_031102.1_Myco_phg_Sneeze
MW291028.1_Myco_phg_Peeb
MG099946.1_Myco_phg_LouisV14
MH779507.1_Myco_phg_Hotshotbaby7
MF668268.1_Myco_phg_Aroostook
MF919497.1_Myco_phg_Chance64
MN444875.1_Myco_phg_Jonghyun
MH045569.1_Myco_phg_Schiebel
NC_012788.1_Myco_phg_Angel
MH001456.1_Myco_phg_CLED96
MH590605.1_Myco_phg_Cherrybomb426
MF668272.1_Myco_phg_Gideon
KT365399.1_Myco_phg_Annihilator
MK305884.1_Myco_phg_BQuat
MH479920.1_Myco_phg_Mowgli
KM923970.1_Myco_phg_Gomashi
MH779513.1_Myco_phg_Olga
KX443326.1_Myco_phg_BruceB
KX664455.1_Myco_phg_Zombie
MH001455.1_Myco_phg_Remy19
MH077584.1_Myco_phg_Phish
KT355472.1_Myco_phg_Cedasite
MK310146.1_Myco_phg_Crespo
KT355474.1_Myco_phg_Frosty24
MK524493.1_Myco_phg_Darionha
GQ303261.1_Myco_phg_Hope
JN699002.1_Myco_phg_Avrafan
KX588251.1_Myco_phg_Jane
MH371119.1_Myco_phg_OctaviousRex
MH371116.1_Myco_phg_DMoney
MK433277.1_Myco_phg_Renaissance
MH919475.1_Myco_phg_Camri
KT347314.2_Myco_phg_Phreak
EU568876.1_Myco_phg_BPs
MK433279.1_Myco_phg_Kareem
MH450127.1_Myco_phg_Plagueis
MH779509.1_Myco_phg_Kasen3
DQ398042.2_Myco_phg_Halo
MH779515.1_Myco_phg_Sweets
KC787111.1_Myco_phg_Sedge,
KC787103.1_Myco_phg_Chy2,
KC787109.1_Myco_phg_Legendre,
KC787112.1_Myco_phg_Clark,
KC787107.1_Myco_phg_Bo4
KC787108.1_Myco_phg_DNAIII
KC787104.1_Myco_phg_Chy3,
KC787110.1_Myco_phg_Leo
KJ725374.1_Myco_phg_Guo1
MH779505.1_Myco_phg_Grizzly
MW119570.1_Myco_phg_Lang
MT818422.1_Myco_phg_Periodt
NC_023705.1_Myco_phg_Liefie
KJ410133.1_Myco_phg_Jolie2
MN234185.1_Myco_phg_Lemuria
MN234219.1_Myco_phg_Mercurio
NC_028861.1_Myco_phg_FlagStaff
MF141540.1_Myco_phg_Avocado
NC_028922.1_Myco_phg_Cambiare
MN234183.1_Myco_phg_Antsirabe
MK494105.1_Myco_phg_Pinnie
NC_028791.1_Myco_phg_MOOREtheMARYer