Number of Samples with HG in this Specific GCF Context − GCF_1
count
15
10
5
0
HOG0000436
HOG0002187
HOG0000437
HOG0000438
HOG0002387
HOG0003066
OG0003344
HOG0002100
OG0003383
HOG0000113
HOG0000439
HOG0000440
HOG0000441
HOG0000442
HOG0000443
HOG0000444
HOG0000445
HOG0000446
HOG0000447
HOG0000448
HOG0000449
HOG0000450
HOG0000114
HOG0000451
reorder(HG, order)
core False True
population Clade_0
Tajima's D ("//" = NOT single−copy in GCF context)
GCF
GCF_1
0
5000
10000
15000
20000
core False True
Tajimas_D
−1 0 1 2
proportion 0.25 0.50 0.75 1.00
Beta−RD ("//" = NOT single−copy in GCF context)
GCF
GCF_1
0
5000
10000
15000
20000
Beta_RD
0.8 0.9 1.0
core False True
proportion 0.25 0.50 0.75 1.00
Proportion of Samples with HG in Multi−Copy Across Genome ("//" = NOT single−copy in GCF context)
GCF
GCF_1
0
5000
10000
15000
20000
core False True
Prop_Multi_Copy
0.000 0.025 0.050 0.075 0.100
proportion 0.25 0.50 0.75 1.00

Number of Samples with HG in this Specific GCF Context − GCF_3
count
reorder(HG, order)
core False True
population Clade_0
Tajima's D ("//" = NOT single−copy in GCF context)
GCF
GCF_3
proportion 0.2 0.4 0.6 0.8 1.0
core False True
Tajimas_D
−1.0 −0.5 0.0 0.5 1.0
Beta−RD ("//" = NOT single−copy in GCF context)
Beta_RD
0.990 0.995 1.000 1.005 1.010
Proportion of Samples with HG in Multi−Copy Across Genome ("//" = NOT single−copy in GCF context)
Prop_Multi_Copy
0

Number of Samples with HG in this Specific GCF Context − GCF_4
count
1.00
0.75
0.50
0.25
0.00
HOG0000649
HOG0000650
HOG0000651
HOG0000652
HOG0000653
OG0003143
OG0003144
OG0003145
HOG0002628
OG0003146
HOG0002044
HOG0000121
reorder(HG, order)
core
False
True
population
Clade_0
Tajima's D ("//" = NOT single−copy in GCF context)
GCF
GCF_4
0
2000
4000
6000
proportion
1
core
False
True
Beta−RD ("//" = NOT single−copy in GCF context)
GCF
GCF_4
0
2000
4000
6000
proportion
1
core
False
True
Proportion of Samples with HG in Multi−Copy Across Genome ("//" = NOT single−copy in GCF context)
GCF
GCF_4
0
2000
4000
6000
proportion
1
core
False
True
Prop_Multi_Copy
0.0
0.1
0.2
0.3
0.4
0.5

Number of Samples with HG in this Specific GCF Context − GCF_2
count
10.0
7.5
5.0
2.5
0.0
HOG0000385
HOG0000384
HOG0000383
OG0003301
HOG0002137
HOG0002136
HOG0002135
HOG0002050
HOG0002134
HOG0002133
HOG0002322
HOG0002293
HOG0002248
reorder(HG, order)
core False True
population Clade_0
Tajima's D ("//" = NOT single−copy in GCF context)
GCF
GCF_2
0
2500
5000
7500
core False True
proportion 0.25 0.50 0.75 1.00
Tajimas_D
−1 0 1
Beta−RD ("//" = NOT single−copy in GCF context)
GCF
GCF_2
0
2500
5000
7500
core False True
Beta_RD
0.98 0.99 1.00 1.01
proportion 0.25 0.50 0.75 1.00
Proportion of Samples with HG in Multi−Copy Across Genome ("//" = NOT single−copy in GCF context)
GCF
GCF_2
0
2500
5000
7500
Prop_Multi_Copy
0
core False True
proportion 0.25 0.50 0.75 1.00