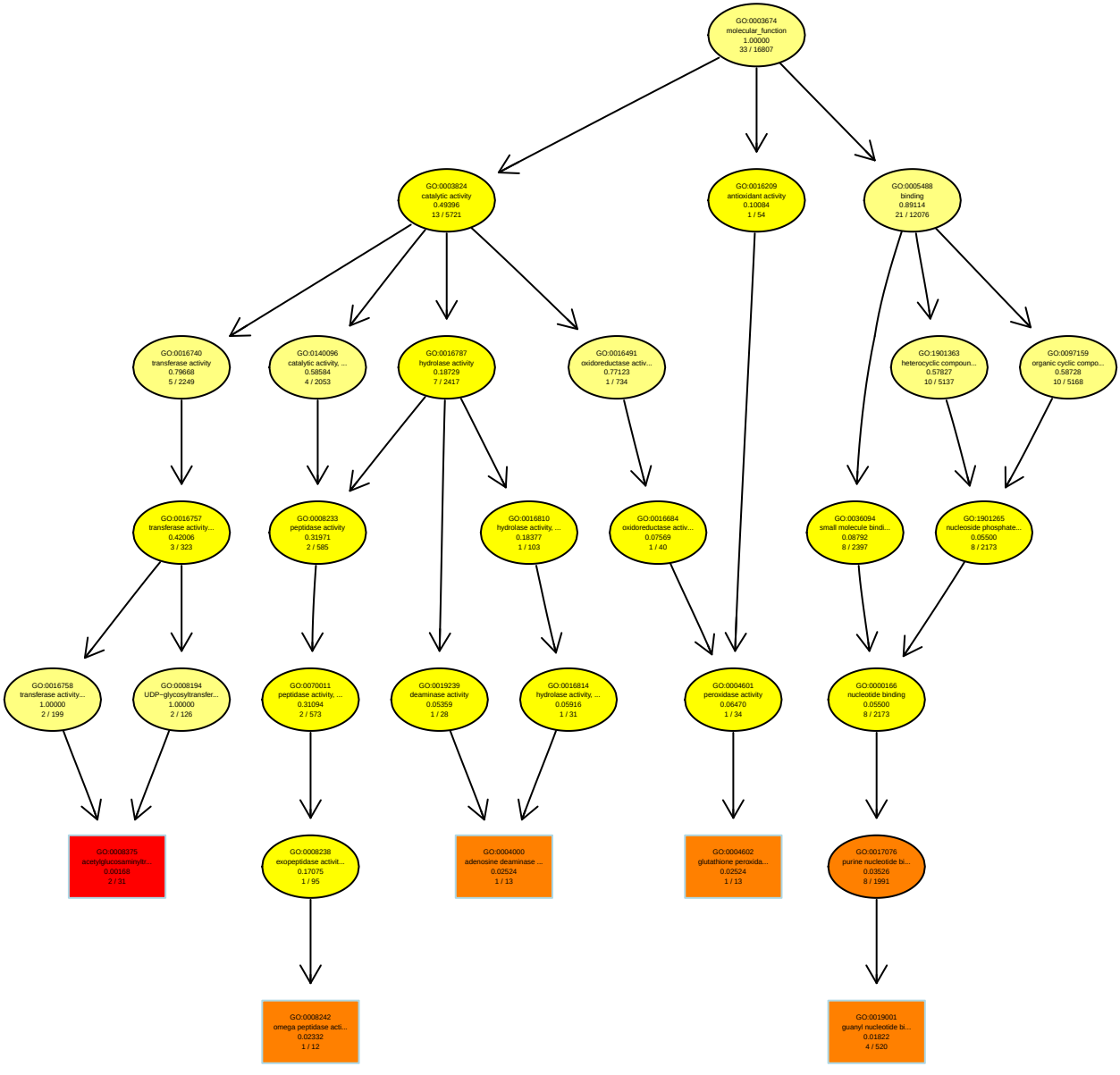
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GO:0003824
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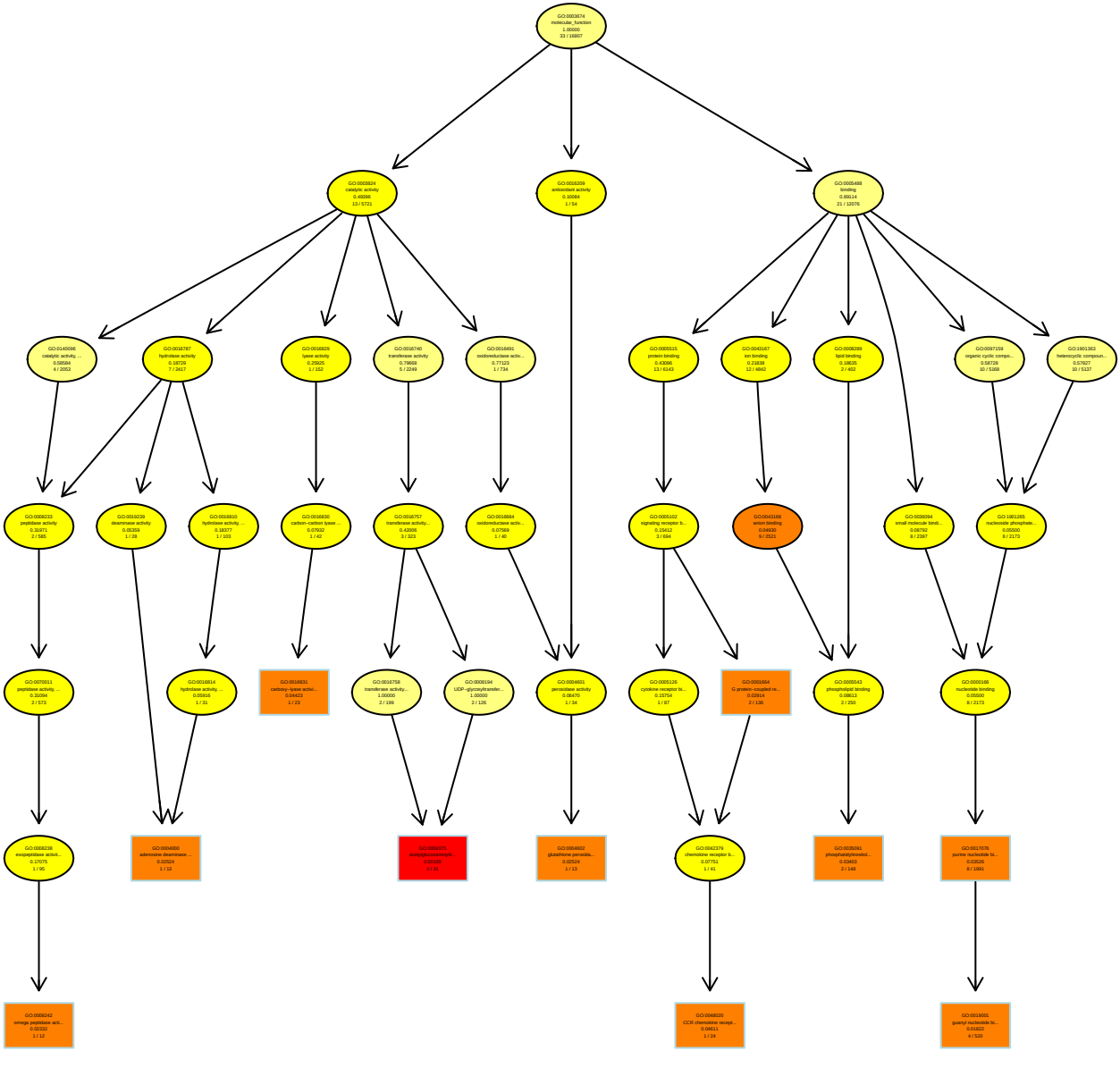
0.49396
13 / 5721
GO:0016209
antioxidant activity
0.10084
1 / 54
GO:0005488
binding
0.89114
21 / 12076
GO:0016740
transferase activity
0.79668
5 / 2249
GO:0140096
catalytic activity, ...
0.58584
4 / 2053
GO:0016787
hydrolase activity
0.18729
7 / 2417
GO:0016491
oxidoreductase activ...
0.77123
1 / 734
GO:1901363
heterocyclic compoun...
0.57827
10 / 5137
GO:0097159
organic cyclic compo...
0.58728
10 / 5168
GO:0016757
transferase activity...
0.42006
3 / 323
GO:0008233
peptidase activity
0.31971
2 / 585
GO:0016810
hydrolase activity, ...
0.18377
1 / 103
GO:0016684
oxidoreductase activ...
0.07569
1 / 40
GO:0036094
small molecule bindi...
0.08792
8 / 2397
GO:1901265
nucleoside phosphate...
0.05500
8 / 2173
GO:0016758
transferase activity...
1.00000
2 / 199
GO:0008194
UDP−glycosyltransfer...
1.00000
2 / 126
GO:0070011
peptidase activity, ...
0.31094
2 / 573
GO:0019239
deaminase activity
0.05359
1 / 28
GO:0016814
hydrolase activity, ...
0.05916
1 / 31
GO:0004601
peroxidase activity
0.06470
1 / 34
GO:0000166
nucleotide binding
0.05500
8 / 2173
GO:0008375
acetylglucosaminyltr...
0.00168
2 / 31
GO:0008238
exopeptidase activit...
0.17075
1 / 95
GO:0004000
adenosine deaminase ...
0.02524
1 / 13
GO:0004602
glutathione peroxida...
0.02524
1 / 13
GO:0017076
purine nucleotide bi...
0.03526
8 / 1991
GO:0008242
omega peptidase acti...
0.02332
1 / 12
GO:0019001
guanyl nucleotide bi...
0.01822
4 / 520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antioxidant activity
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1 / 54
GO:0005488
binding
0.85114
21 / 12076
GO:0140096
catalytic activity, ...
0.58584
4 / 2053
GO:0016787
hydrolase activity
0.18719
7 / 2417
GO:0016829
lyase activity
0.23925
1 / 152
GO:0016740
transferase activity
0.78668
5 / 2249
GO:0016491
oxidoreductase activ...
0.77123
1 / 734
GO:0005515
protein binding
0.43086
13 / 6143
GO:0043167
ion binding
0.31838
12 / 4842
GO:0008289
lipid binding
0.18635
2 / 402
GO:0097159
organic cyclic compo...
0.58728
10 / 5168
GO:1901363
heterocyclic compoun...
0.57827
10 / 5137
GO:0008233
peptidase activity
0.31971
2 / 585
GO:0016229
dearminase activity
0.05269
1 / 28
GO:0016810
hydrolase activity, ...
0.16377
1 / 103
GO:0016830
carbon-carbon lyase ...
0.07632
1 / 42
GO:0016757
transferase activity, ...
0.42006
3 / 323
GO:0016684
oxidoreductase activ...
0.07569
1 / 40
GO:0005102
signaling receptor b...
0.15412
3 / 694
GO:0043168
anion binding
0.04930
8 / 2521
GO:0036094
small molecule bind...
0.08792
8 / 2397
GO:1901265
nucleoside phosphate...
0.05000
8 / 2173
GO:0070011
peptidase activity, ...
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2 / 573
GO:0016814
hydrolase activity, ...
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1 / 31
GO:0016831
carbon-lyase activi...
0.04423
1 / 23
GO:0016758
transferase activity, ...
1.00000
2 / 199
GO:0008194
UDP-glycosyltransfer...
1.00000
2 / 126
GO:0004601
peroxidase activity
0.06478
1 / 34
GO:0005126
cytokine receptor bi...
0.33754
1 / 87
GO:0031664
G protein-coupled re...
0.03014
2 / 136
GO:0005543
phospholipid binding
0.08613
2 / 250
GO:0000166
nucleotide binding
0.05500
8 / 2173
GO:0008238
exopeptidase activi...
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1 / 95
GO:0004000
adenosine deaminase ...
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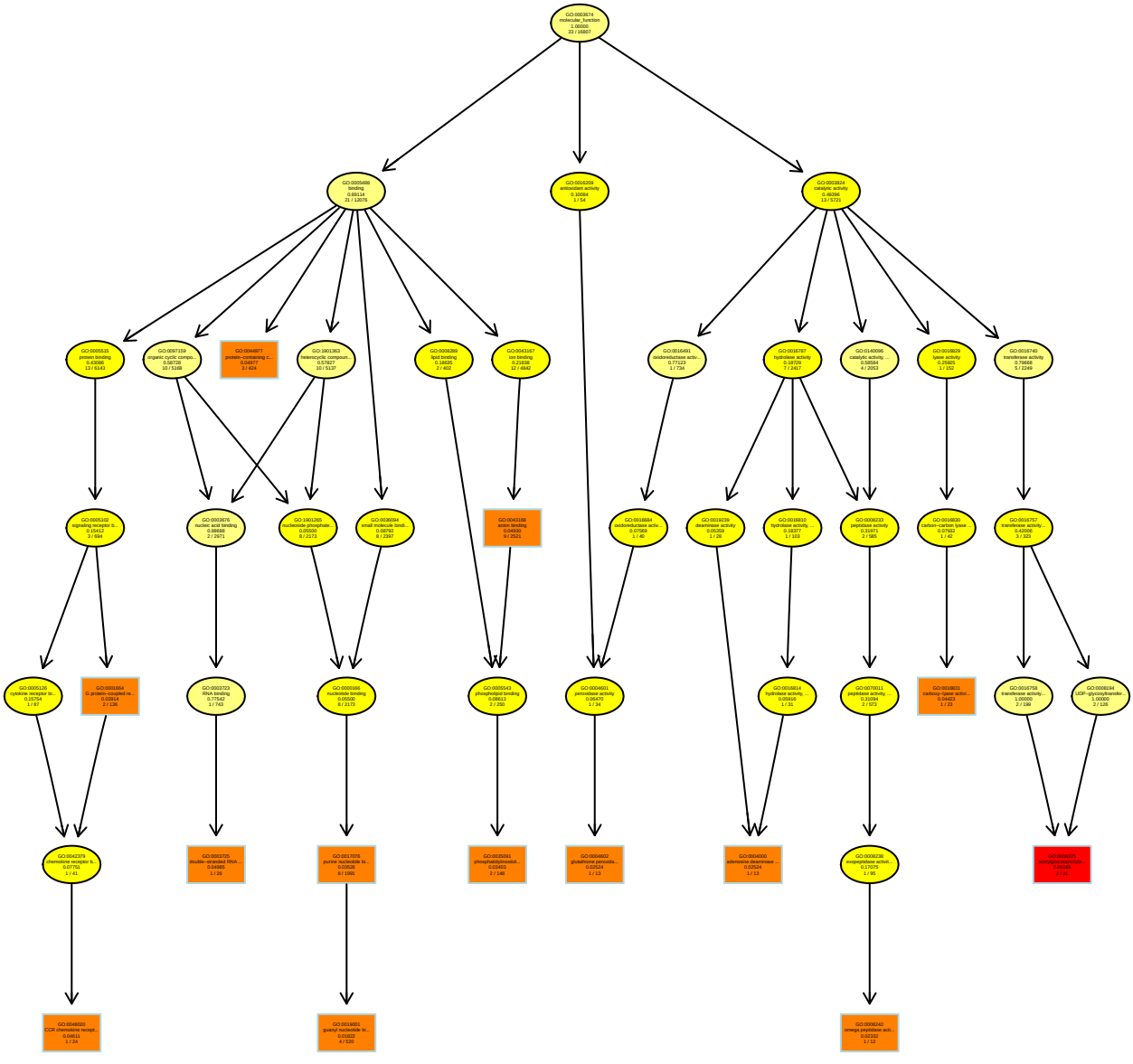
1 / 13
GO:0009975
acetylglucosaminyltr...
0.00148
2 / 31
GO:0004602
glutathione peroxida...
0.02524
1 / 13
GO:0042379
chemokine receptor b...
0.07751
1 / 41
GO:0035091
phosphoinositol bind...
0.03403
2 / 148
GO:0017076
purine nucleotide bi...
0.03526
8 / 1991
GO:0008242
omega peptidase acti...
0.02332
1 / 12
GO:0048020
CCR chemokine recept...
0.06611
1 / 24
GO:0019001
guanyl nucleotide bi...
0.01822
4 / 520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GO:0005199
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1 / 54
GO:0003824
catalytic activity
0.46356
13 / 5721
GO:0005515
protein binding
0.43095
13 / 6143
GO:0097159
organic cyclic compo...
0.58728
10 / 5168
GO:0048877
protein–containing c...
0.04977
3 / 424
GO:0001883
heterocyclic compoun...
0.57827
10 / 5137
GO:0008289
lipid binding
0.08035
2 / 452
GO:0043167
ion binding
0.21938
12 / 4842
GO:0016491
oxidoreductase activ...
0.77123
1 / 734
GO:0016787
hydrolase activity
0.18729
7 / 2417
GO:0140096
catalytic activity, ...
0.58694
4 / 2053
GO:0016829
lyase activity
0.35825
1 / 152
GO:0016740
transferase activity
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5 / 2249
GO:0005102
signaling receptor b...
0.15412
3 / 694
GO:0003676
nucleic acid binding
0.86688
2 / 2971
GO:1901265
nucleoside phosphate...
0.05500
0 / 2173
GO:0030234
enzyme regulator a...
0.08792
0 / 2397
GO:0043168
anion binding
0.04930
0 / 2021
GO:0016684
oxidoreductase activ...
0.07569
1 / 40
GO:0019239
deaminase activity
0.05359
1 / 28
GO:0016810
hydrolase activity, ...
0.18377
1 / 153
GO:0008233
peptidase activity
0.31971
2 / 985
GO:0016830
carbon–carbon lyase ...
0.07932
1 / 42
GO:0016757
transferase activity, ...
0.42006
3 / 323
GO:0005126
cytokine receptor bi...
0.11254
1 / 47
GO:0001664
G protein–coupled re...
0.02934
0 / 136
GO:0003723
RNA binding
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1 / 743
GO:0005185
nucleoside binding
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0 / 2175
GO:0005543
phospholipid binding
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GO:0004601
peroxidase activity
0.04475
1 / 34
GO:0016814
hydrolase activity, ...
0.09914
1 / 31
GO:0070011
peptidase activity, ...
0.31059
2 / 953
GO:0016831
carboxy–lyase activi...
0.04423
1 / 20
GO:0016758
transferase activity, ...
1.00000
2 / 199
GO:0008194
UDP–glycosyltransfer...
1.00000
2 / 139
GO:0042379
chemokine receptor b...
0.07791
1 / 41
GO:0003725
double–stranded RNA ...
0.04665
1 / 26
GO:0017076
purine nucleotide bi...
0.05536
0 / 1991
GO:0035091
phosphatidylinositol ...
0.03403
2 / 148
GO:0004602
glutathione peroxida...
0.02524
1 / 13
GO:0004000
adenosine deaminase ...
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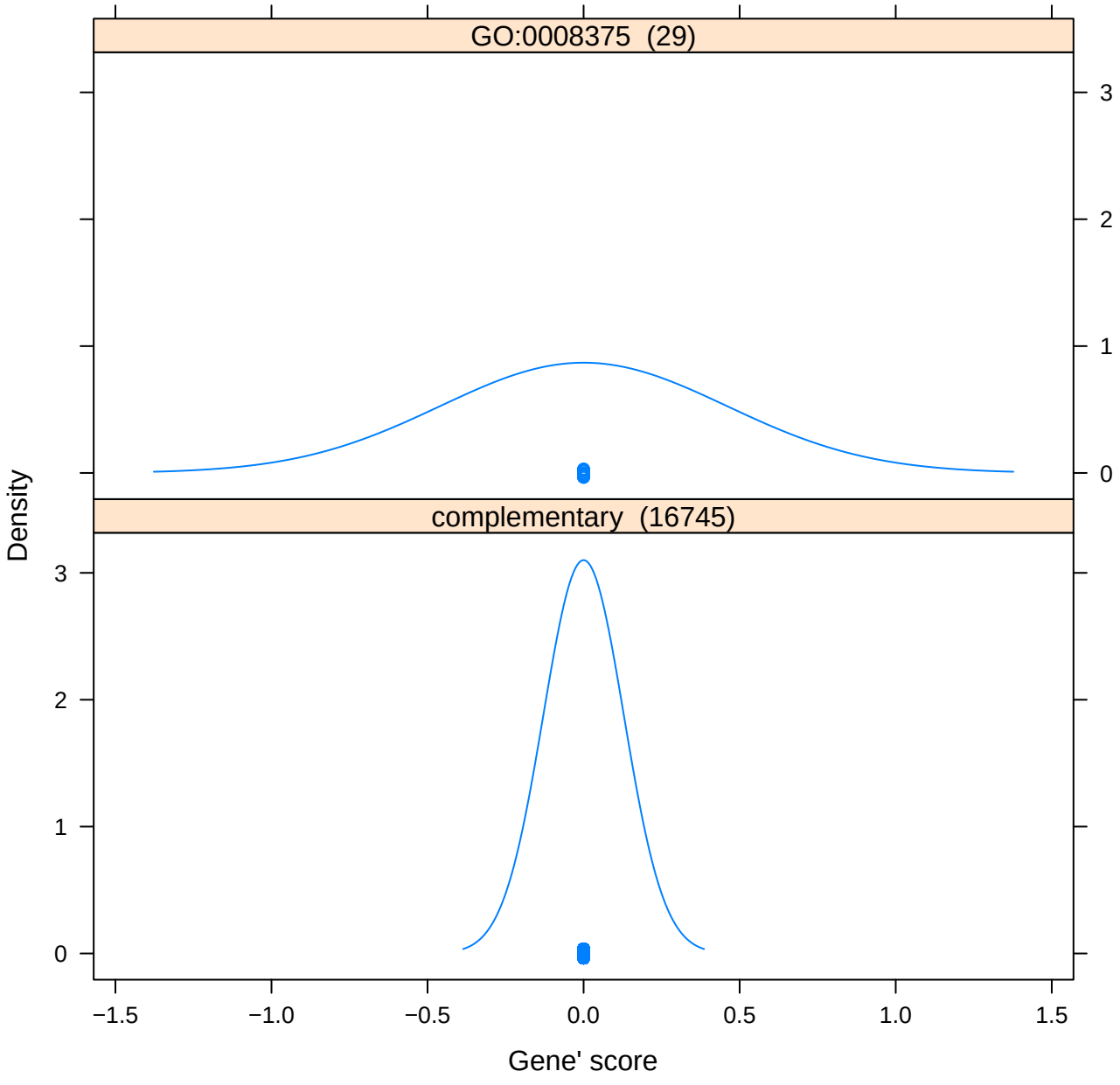
1 / 13
GO:0008236
exopeptidase activit...
0.17075
1 / 95
GO:0008270
zinc ion binding
0.06168
2 / 31
GO:0048020
CCR chemokine recept...
0.04611
1 / 24
GO:0019001
guanyl nucleotide bi...
0.01822
0 / 520
GO:0008242
omega peptidase acti...
0.02332
1 / 12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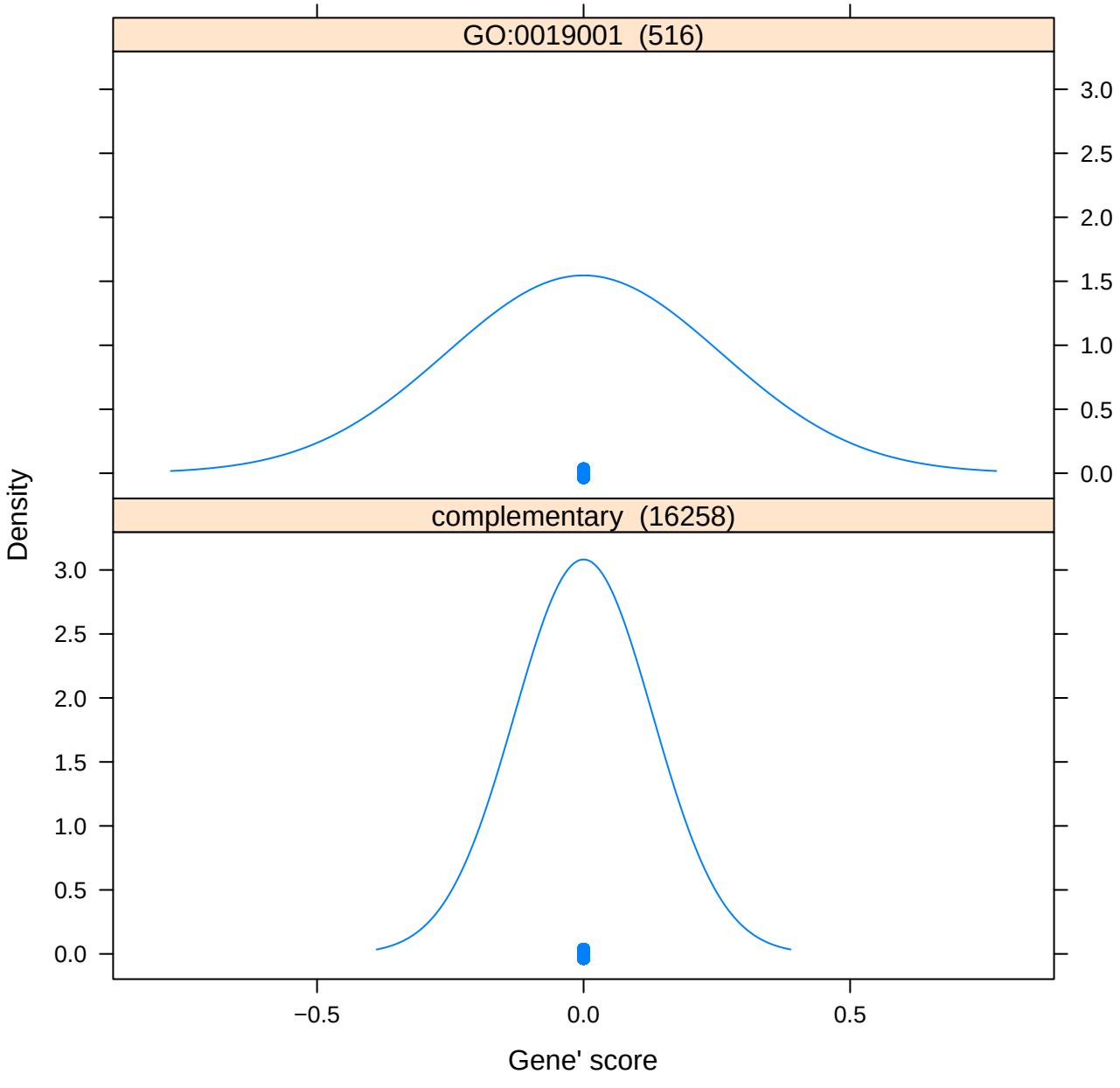
0
1
2
3


GO:0019001 (516)
complementary (16258)
Density
Gene' score
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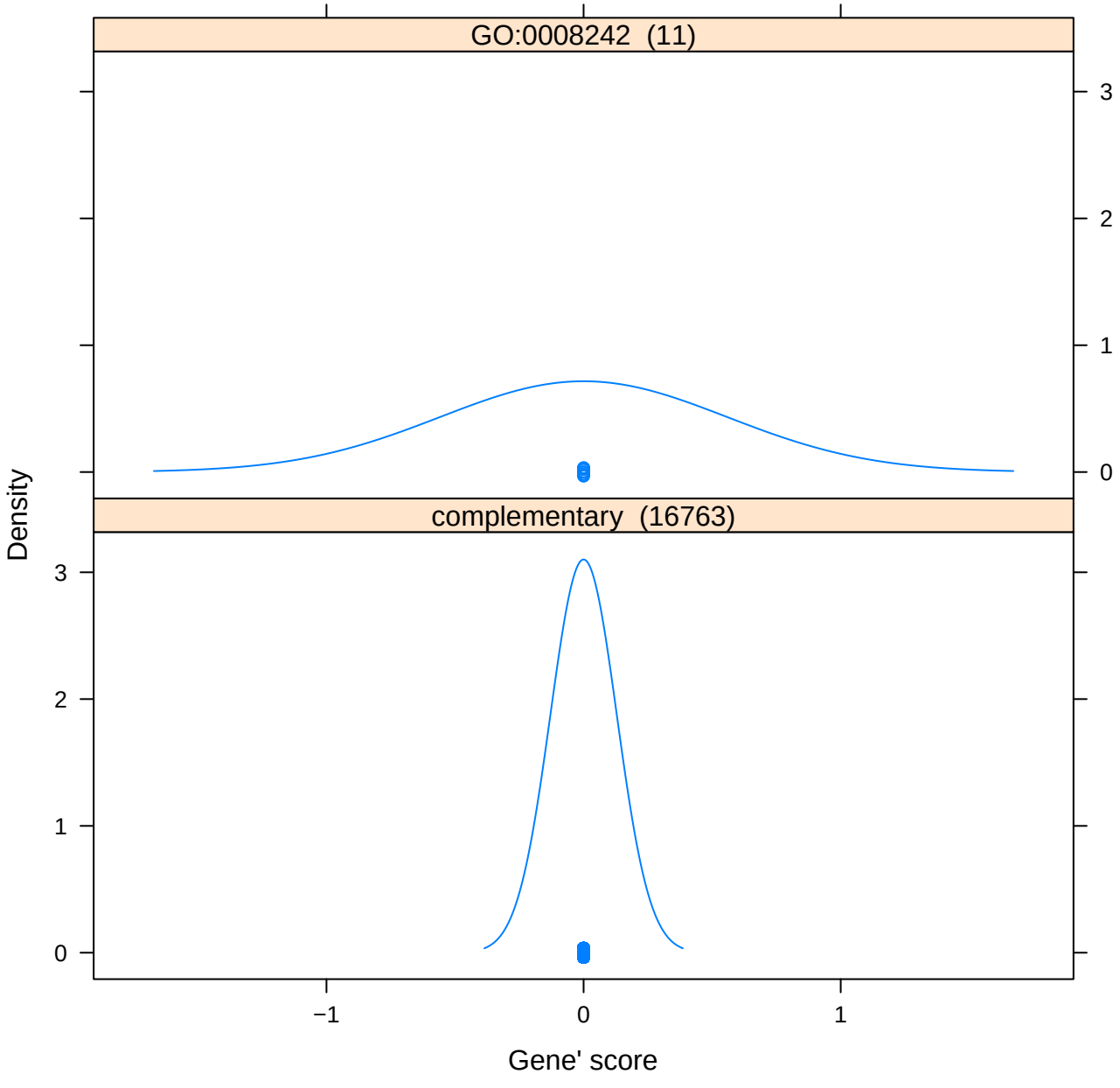
GO:0008242 (11)
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Density
Gene' score
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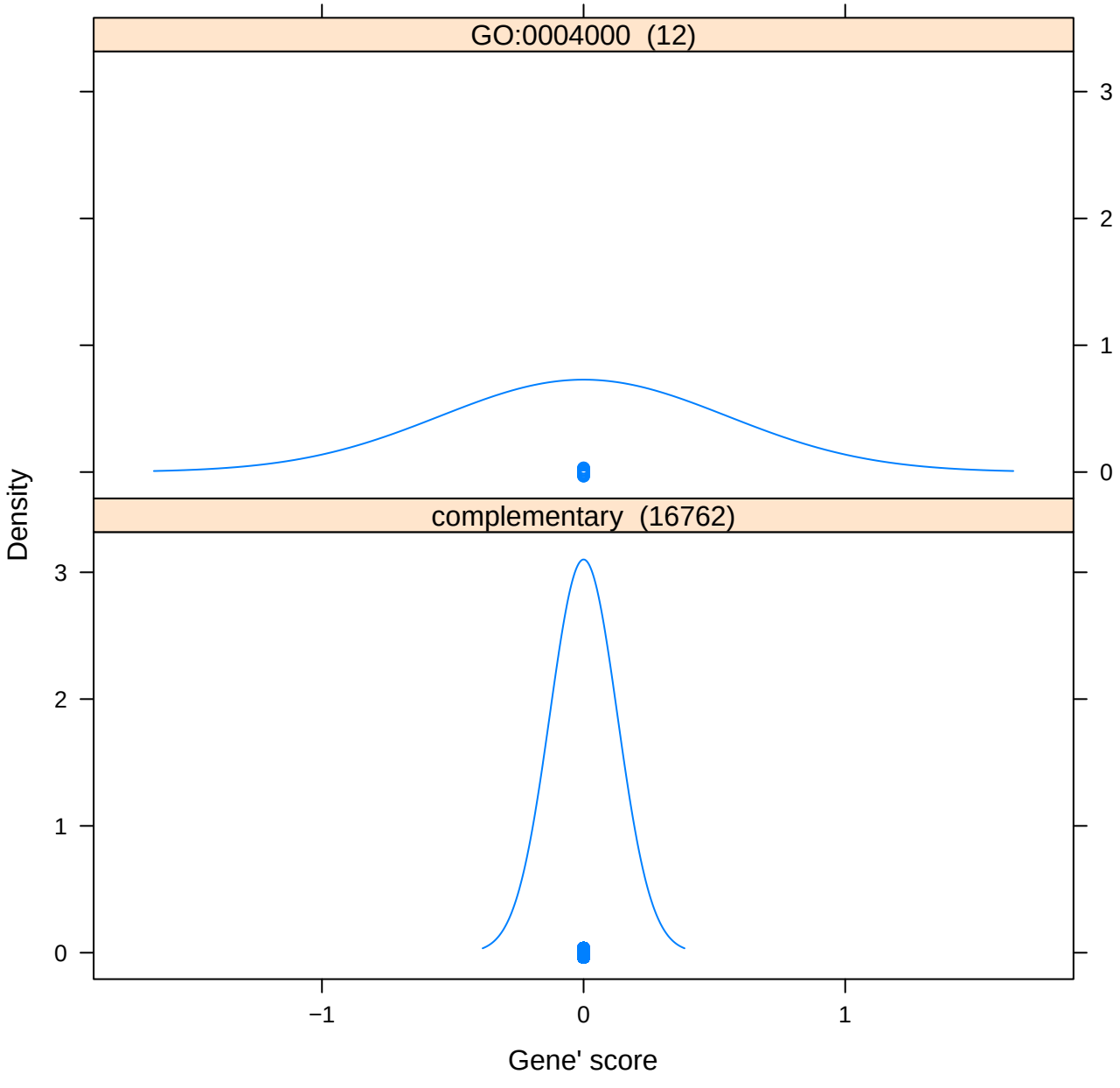
0
1
2
3


GO:0004000 (12)
complementary (16762)
Density
Gene' score
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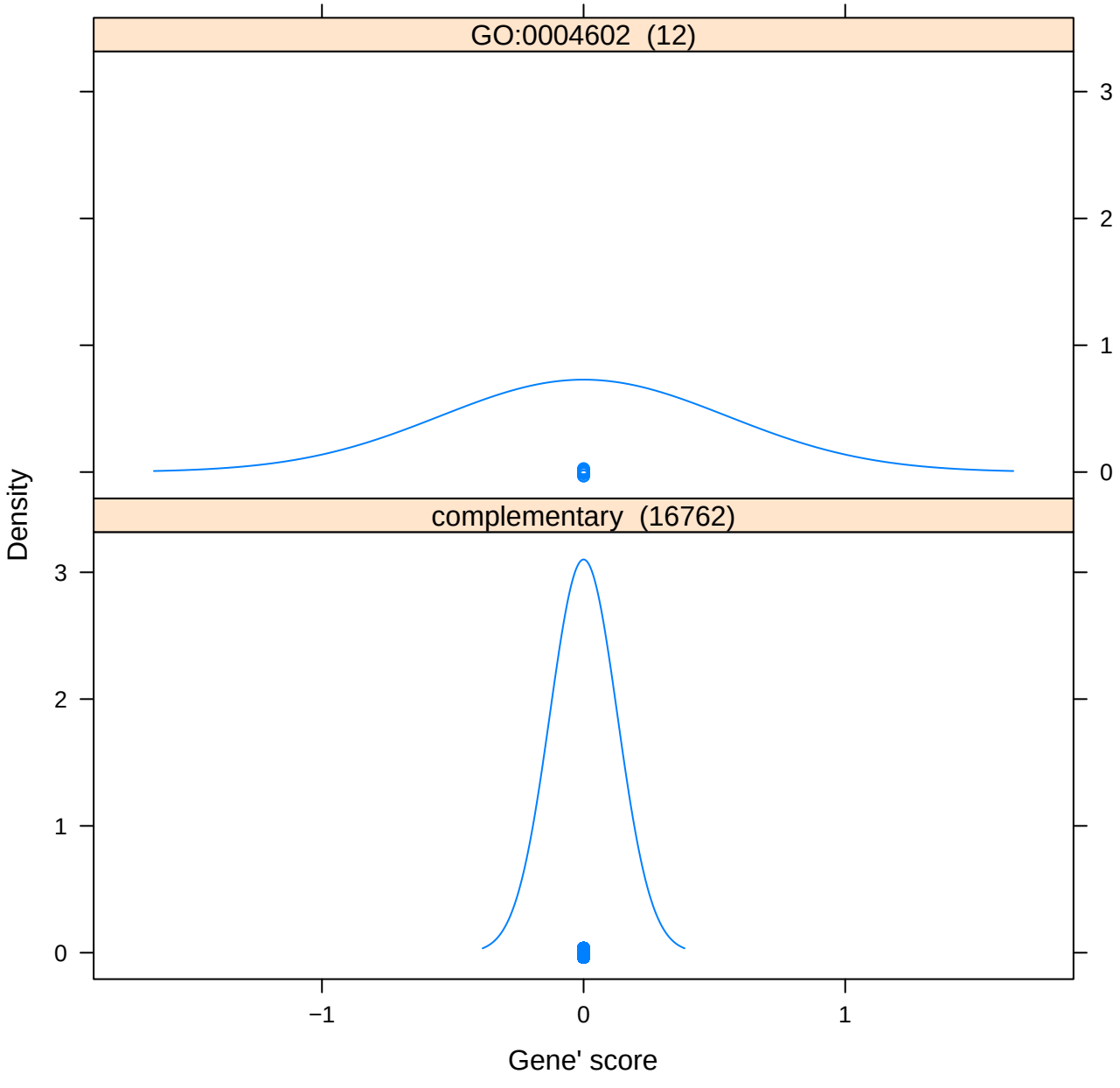
0
1
2
3


GO:0004602 (12)
complementary (16762)
Density
Gene' score
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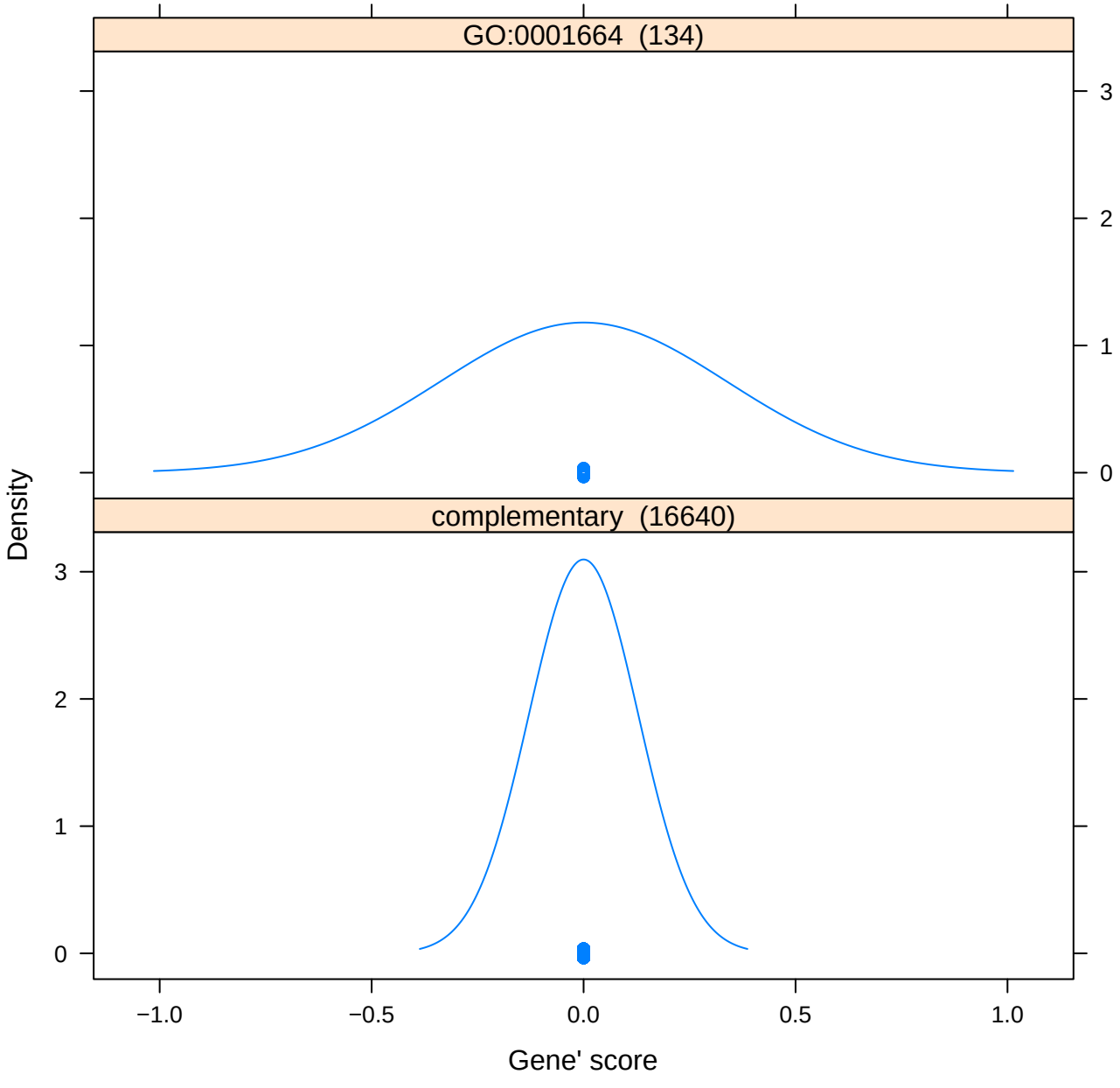
1
2
3


GO:0001664 (134)
complementary (16640)
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Gene' score
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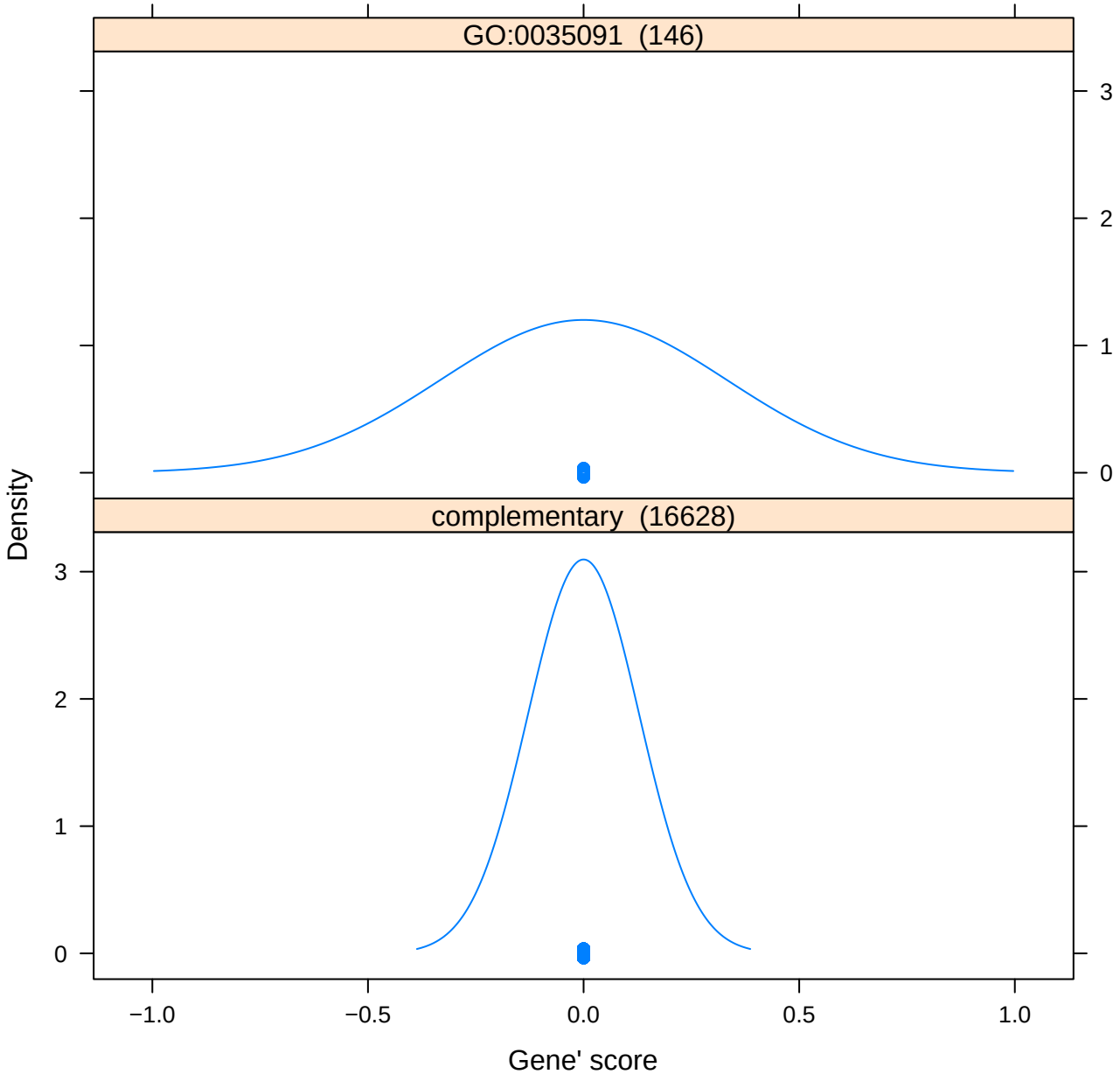
2
3


GO:0035091 (146)
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Gene' score
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0.5
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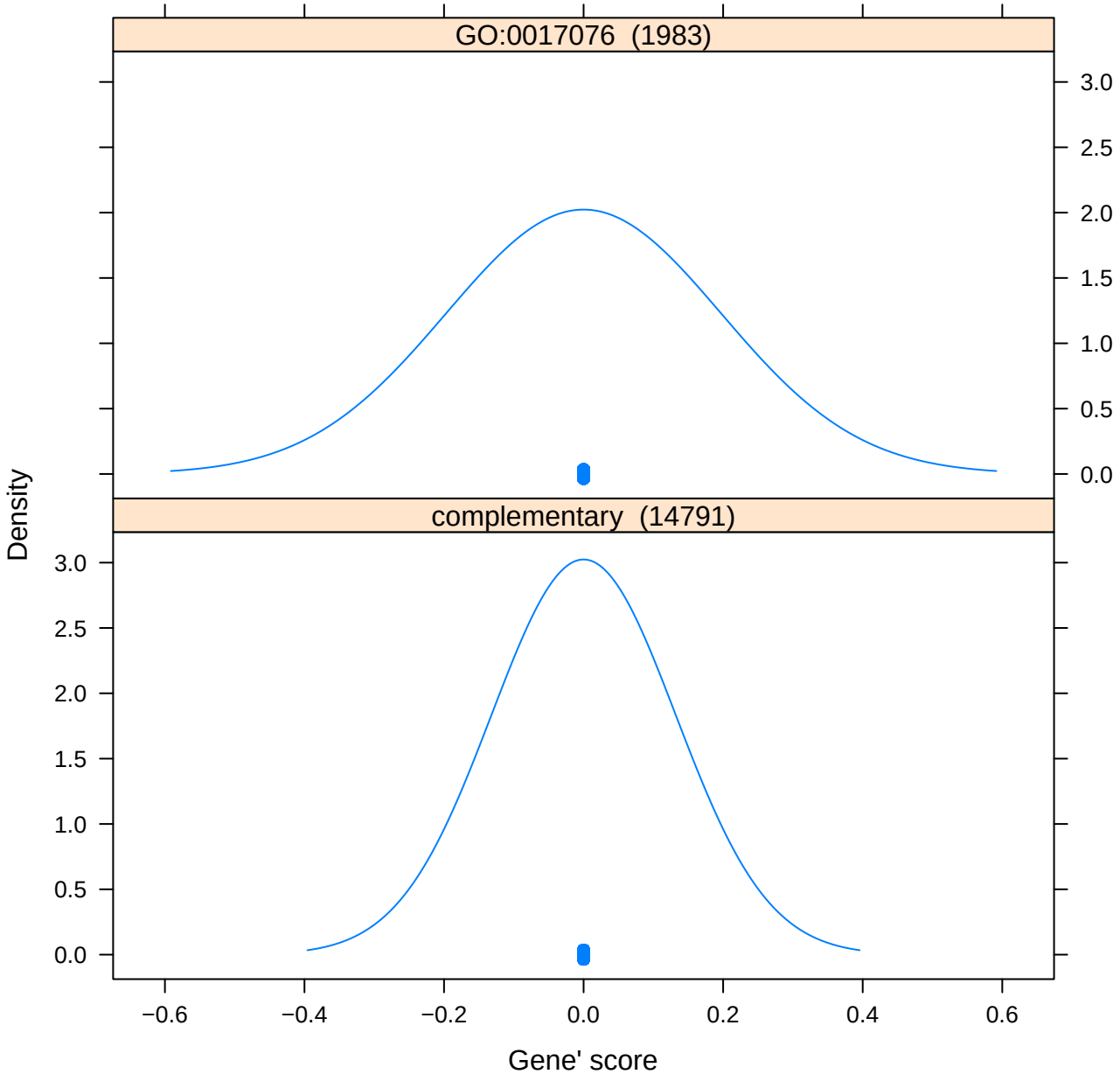
0
1
2
3


GO:0017076 (1983)
complementary (14791)
Density
Gene' score
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−0.4
−0.2
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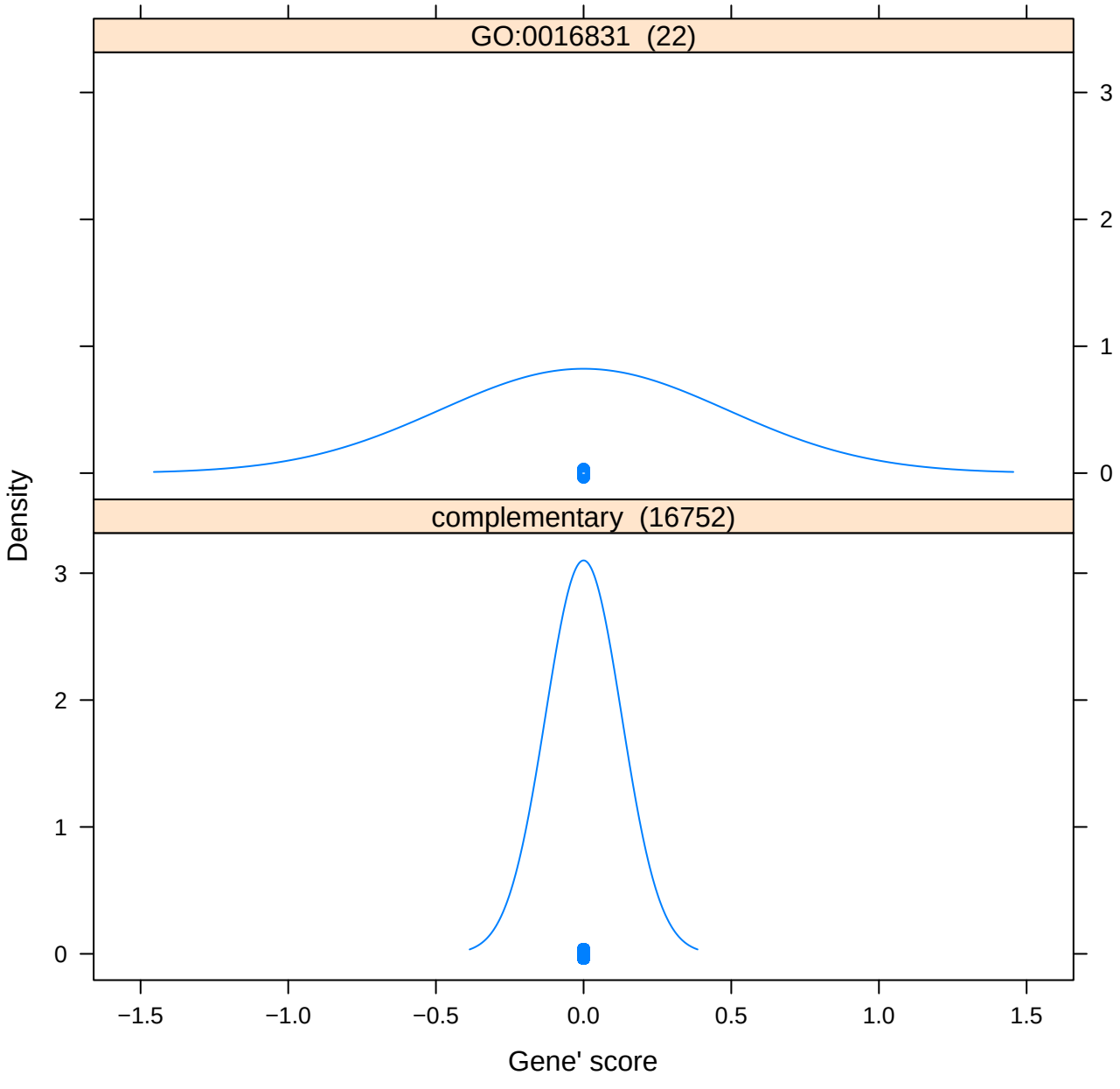
0.4
0.6


GO:0016831 (22)
complementary (16752)
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Gene' score
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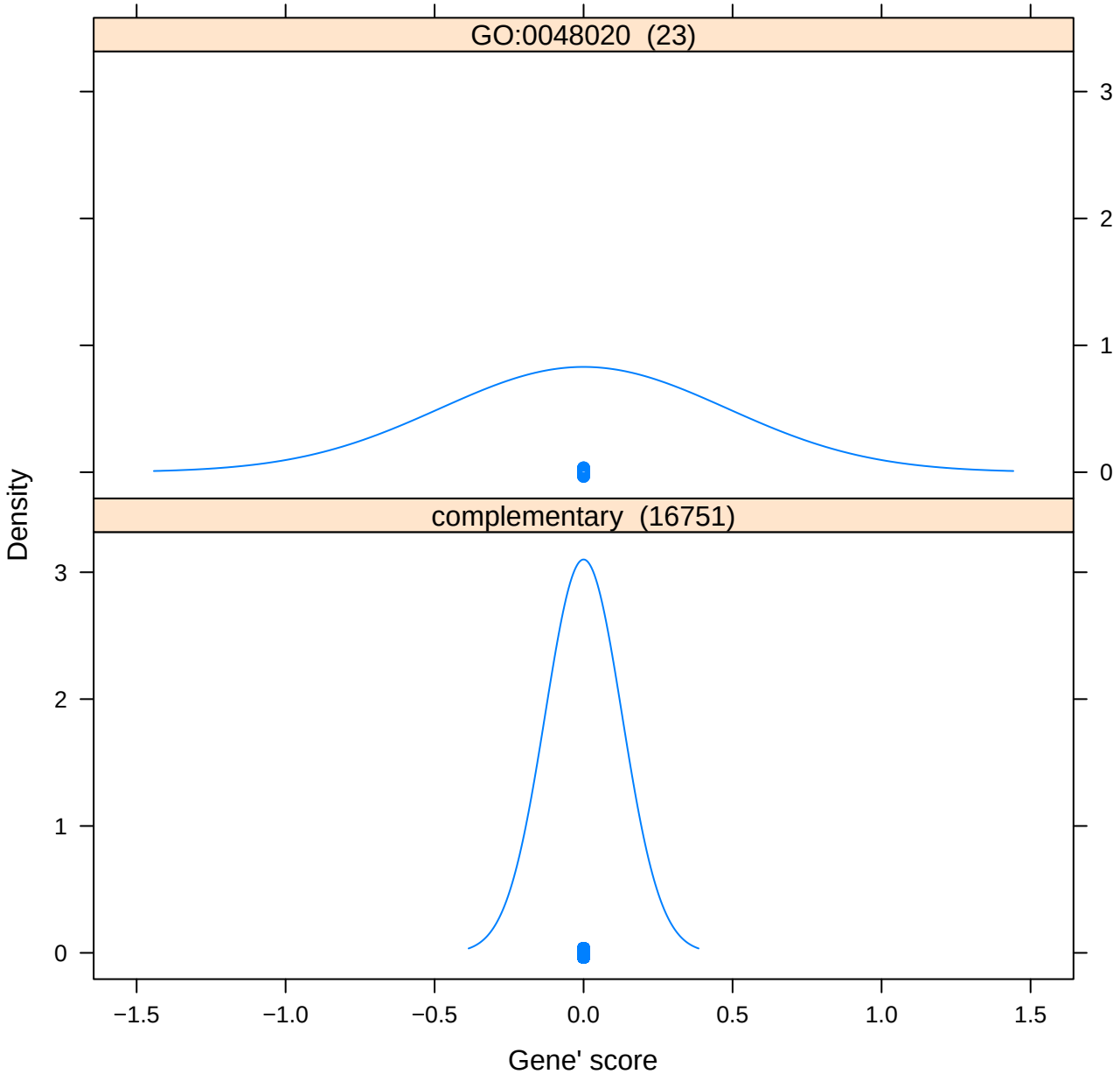
0
1
2
3


GO:0048020 (23)
complementary (16751)
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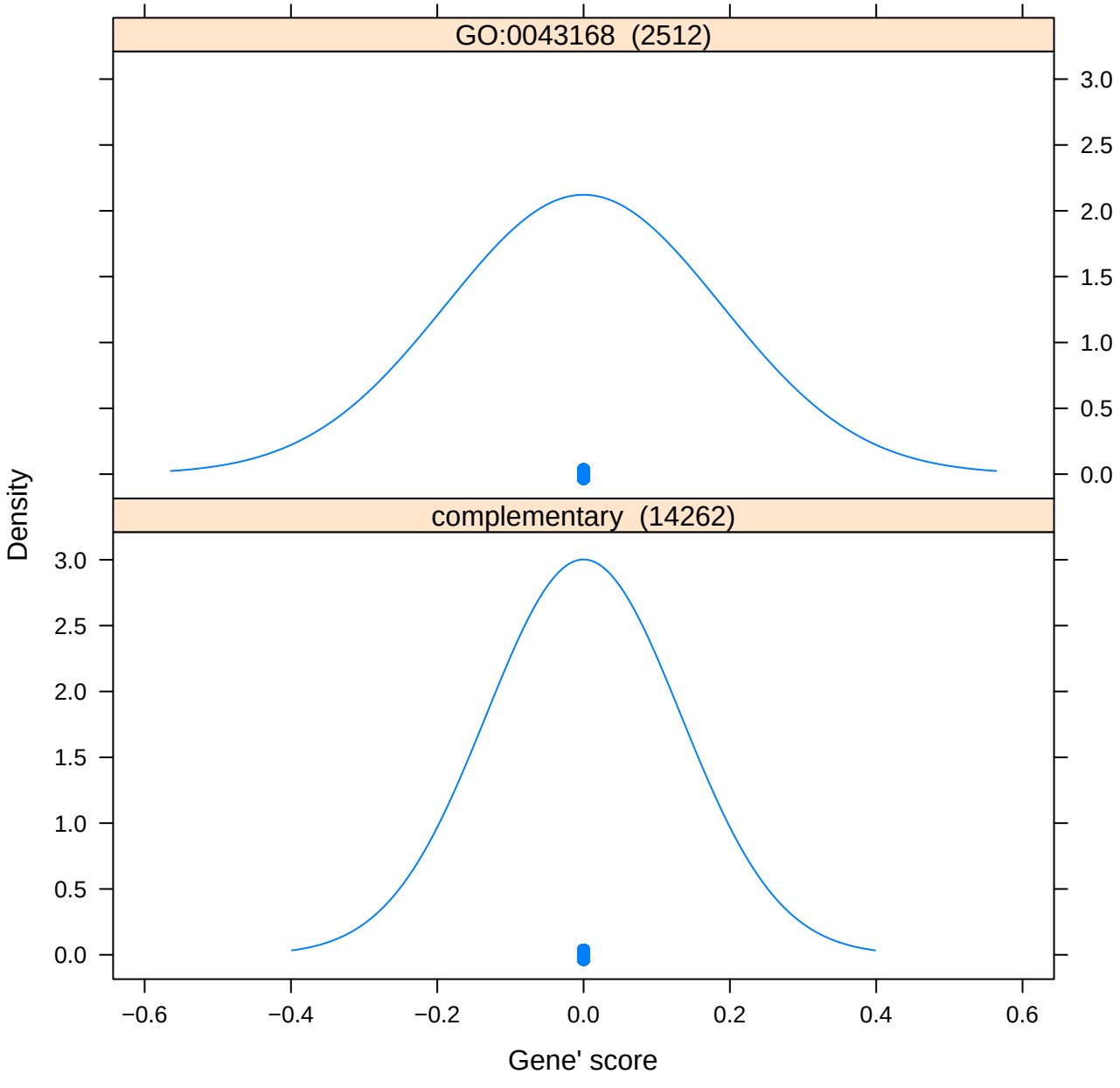
1.5
0
1
2
3


GO:0043168 (2512)
complementary (14262)
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Gene' score
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1.5
1.0
0.5
0.0
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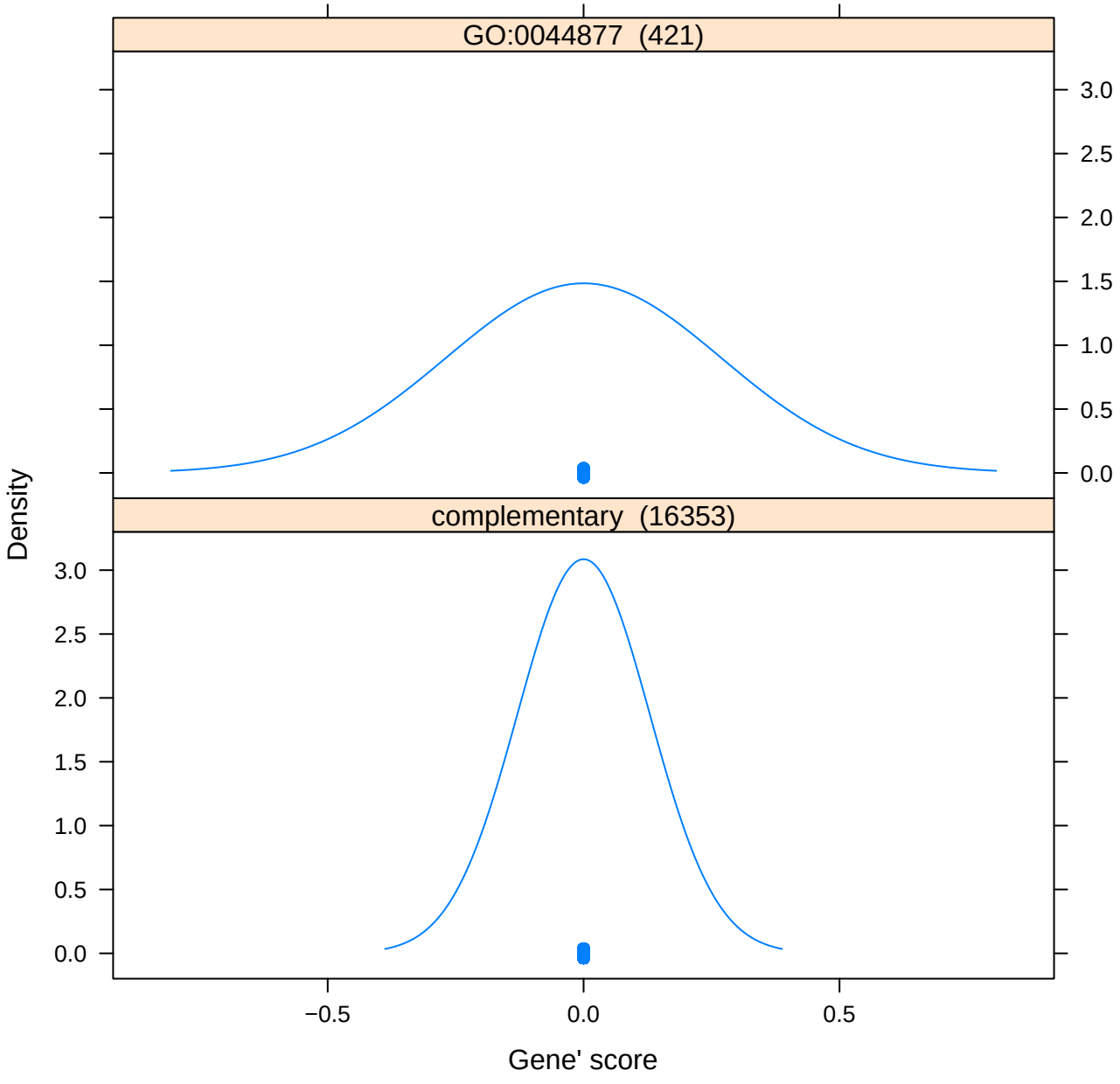
−0.4
−0.2
0.0
0.2
0.4
0.6


GO:0044877 (421)
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Gene' score
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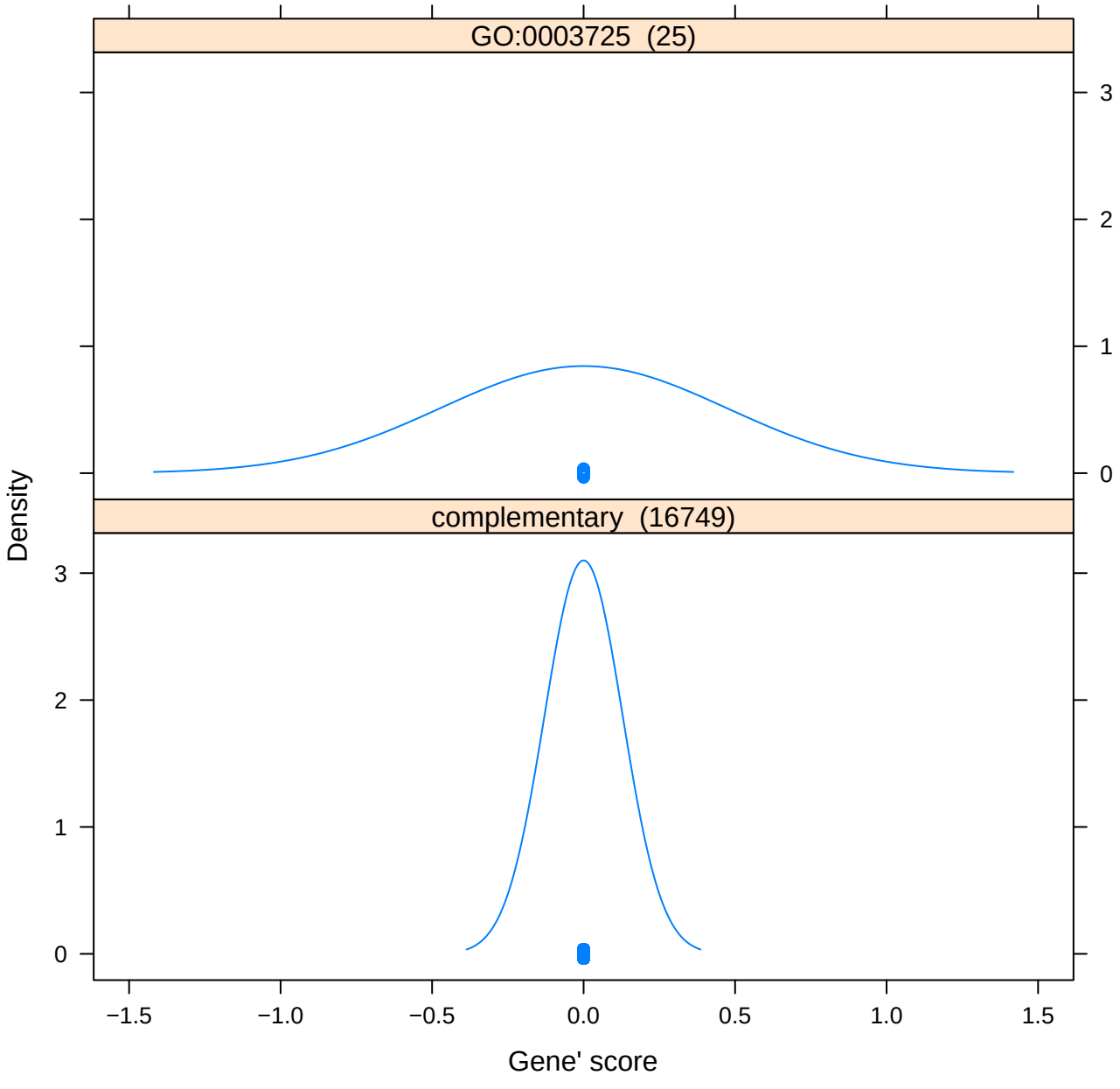
0.5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complementary (16749)
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Gene' score
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−0.5
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0.5
1.0
1.5
0
1
2
3