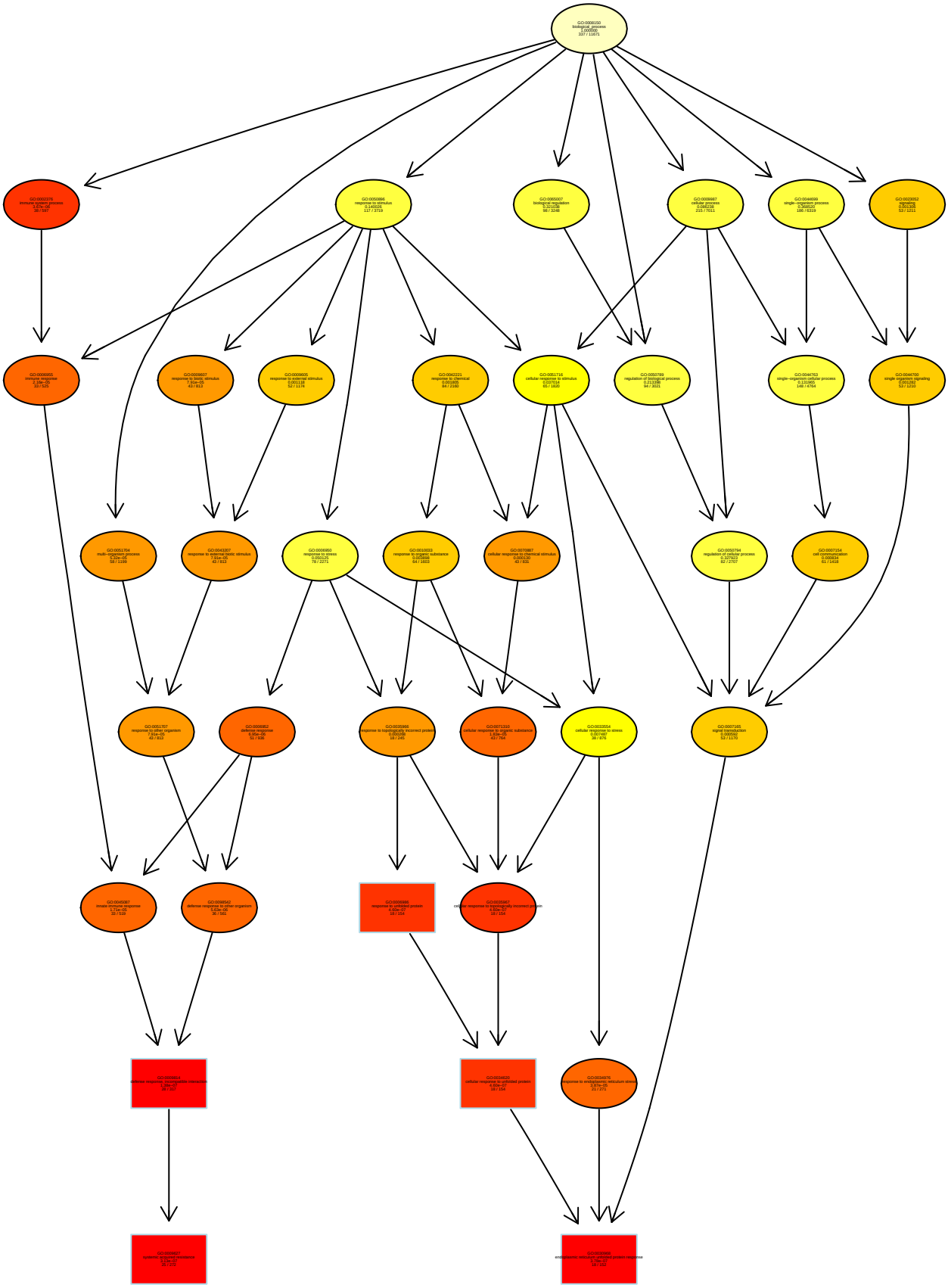
GO:0008150
biological_process
1.000000
337 / 11671
GO:0002376
immune system process
1.07e-06
38 / 597
GO:0050896
response to stimulus
0.146026
117 / 3719
GO:0065007
biological regulation
0.321076
98 / 3248
GO:0008987
cellular process
0.084238
215 / 7011
GO:0044699
single-organism process
0.368626
166 / 6319
GO:0023052
signaling
0.501308
53 / 1211
GO:0006955
immune response
2.14e-06
23 / 525
GO:0006607
response to DNA damage stimulus
8.1e-05
43 / 813
GO:0009605
response to external stimulus
0.011118
52 / 1174
GO:0042221
response to chemical
0.001408
84 / 2160
GO:0051716
cellular response to stimulus
0.007014
65 / 1820
GO:0050789
regulation of biological process
0.213382
94 / 3021
GO:0044763
single-organism cellular process
0.210365
148 / 4764
GO:0044700
single organism signaling
0.001413
53 / 1210
GO:0051704
multi-organism process
0.000112
52 / 1186
GO:0043207
response to external biotic stimulus
0.000112
43 / 812
GO:0006950
response to stress
0.000000
79 / 2271
GO:0010033
response to organic substance
0.000000
64 / 1803
GO:0070887
cellular response to chemical stimulus
0.000000
42 / 826
GO:0050794
regulation of cellular process
0.000000
62 / 2707
GO:0007154
cell communication
0.000000
61 / 1418
GO:0051707
response to other organism
7.91e-05
42 / 813
GO:0006952
defense response
6.95e-06
51 / 936
GO:0033993
response to lipid
0.000258
18 / 245
GO:0071310
cellular response to organic substance
1.93e-05
42 / 764
GO:0033554
cellular response to stress
0.007487
39 / 876
GO:0007165
signal transduction
0.000592
52 / 1170
GO:0045087
innate immune response
1.77e-05
33 / 519
GO:0098542
defense response to other organism
5.63e-06
36 / 561
GO:0006986
response to unfolded protein
4.65e-07
18 / 154
GO:0035967
cellular response to topologically incorrect protein
4.65e-07
18 / 154
GO:0009614
defense response, incompatible interaction
1.24e-07
31 / 317
GO:0034620
cellular response to unfolded protein
4.65e-07
18 / 154
GO:0034976
response to endoplasmic reticulum stress
3.97e-05
21 / 373
GO:0002227
systemic innate immune response
2.25e-07
27 / 272
GO:0030968
endoplasmic reticulum unfolded protein response
2.79e-07
18 / 152