

p=1.1E-03 | n=2 p=3.2E-03 | n=2 fg=0.04 fg=0.04 bg=0.00 bg=0.00 G protein-coupled adenosine receptor act... GO:0001609 heterotrimeric G-protein binding GO:0032795
phospholipase C activity GO:0004629
transaminase activity GO:0008483
G-protein beta/gamma-subunit complex bin... GO:003170
vitamin B6 binding GO:0070279 p=6.3E-03 n=2 p=1.0E-02 n=2 fg=0.04 bg=0.00 fg=0.04 bg=0.00 p=1.0E-02 n=2 p=1.5E-02 n=2 fğ=0.04 bg=0.00 bg=0.00 fg=0.04 vitamin B6 binding GO:0070279
nucleoside–triphosphatase regulator acti... GO:0060589
transferase activity, transferring nitro... GO:0016769
guanyl–nucleotide exchange factor activi... GO:0005085
ornithine–oxo–acid transaminase activity GO:0004587
RNA polymerase I core binding GO:0001042
Rho–dependent protein serine/threonine k... GO:0072518
RNA strand–exchange activity GO:0034057
ATPase–coupled inorganic anion transmemb... GO:0043225
aminophospholipid flippase activity GO:0015247
selenium binding GO:0008430
phosphatidylglycerol binding GO:1901612
histone H3–tri/di/monomethyl–lysine–4 de... GO:0034647
4–hydroxyglutamate transaminase activity GO:0047578
nickel cation binding GO:01901615
adipokinetic hormone receptor activity GO:0097003
methylenetetrahydrofolate dehydrogenase ... GO:0004486 p=1.5E-02 p=2.0E-02 bg=0.00 bg=0.05 n=2 fg=0.04 n=7 $f_{q=0.13}$ fg=0.04 fg=0.08 p=2.1E-02 p=2.2E-02 n=2bg=0.00 n=4 bg=0.02 fg=0.02 fg=0.02 bg=0.00 p=3.3E-02 p=3.3E-02bg=0.00 bg=0.00 n=1p=3.3E-02 p=3.3E-02 fg=0.02 fg=0.02 n=1bg = 0.00p=3.3E-02 n=1fg=0.02 bg=0.00 fg=0.02 bg=0.00 p=3.3E-02n=1p=3.3E bg=0.00 n=1p=3.3E-02fg=0.02 n=1bg=0.00 p=3.3E-02 bg=0.00 n=1fg=0.02 p=3.3E-02bg=0.00 bg=0.00 n=1p=3.3E-02 fg=0.02 n=1fg=0.02 p=3.3E-02 p=3.3E-02 n=1bg=0.00 fg=0.02 bg=0.00 n=1adipokinetic hormone receptor activity GO:0097003
methylenetetrahydrofolate dehydrogenase ... GO:0004486
methylenetetrahydrofolate dehydrogenase ... GO:0004487
methylenetetrahydrofolate dehydrogenase ... GO:0004487
calmodulin–activated 3',5'–cyclic–GMP ph... GO:0044101
DNA binding, bending GO:0008301
methenyltetrahydrofolate cyclohydrolase ... GO:0004477
calmodulin–activated dual specificity 3'... GO:0004117 p=3.3E-02 fg=0.02 bg=0.00 n=1p=3.3E-02 fg=0.02 bg=0.00 n=1fg=0.02 p=3.3E-02 n=1bg=0.00 fg=0.02 bg=0.00 bg=0.00 p=3.3E-02 n=1p=3.3E-02 p=3.3E-02 fg=0.02 fg=0.02 n=1 n=1bq = 0.00p=3.3E-02 fg=0.02 bg=0.00 0 2 0.0 0.2 1 3 5 0.4 0.6 8.0 1.0

Fox_Nvec_vc1.1_XM_032379347.2

GO:CC Fox_Nvec_vc1.1_XM_032379347.2 fra

-log(p)
n=53/204 input genes with annotations

fraction genes in fg and expected valu

fraction

