

calcium, potassium:sodium antiporter act... GO:0008273
lysophospholipase activity GO:0004622
potassium ion binding GO:0030955
sodium ion binding GO:0031402
voltage-gated potassium channel activity GO:0005249
voltage-gated sodium channel activity GO:0005248
histone H3-methyl-lysine-36 demethylase ... GO:0051864
intracellular ligand-gated ion channel a... GO:0005217
dynein light chain binding GO:0045503
ubiquitin conjugating enzyme binding GO:0031624 p=1.3E-03 | n=2 p=1.5E-03 | n=3 bg=0.00 bg=0.00 fg=0.02 fg=0.03 p=3.8E-03 n=2 p=3.8E-03 n=2 fg=0.02 bg=0.00 fg=0.02 bg=0.00 p=6.2E-03 n=3 p=7.5E-03 n=2 fg=0.03 bg=0.00 fg=0.02 bg=0.00 p=7.5E-03 n=2 p=1.0E-02 n=3 fg=0.02 bg=0.00 fg=0.03 bg=0.01 fğ=0.02 p=1.2E-02 n=2 p=1.3E-02 n=3 bg=0.00 ubiquitin conjugating enzyme binding GO:0043505
ubiquitin conjugating enzyme binding GO:0031624
ion gated channel activity GO:0022839
phosphatidylinositol–3,5–bisphosphate bi... GO:0080025
ligand–gated ion channel activity GO:0015276 fg=0.03 bg=0.01 fg=0.07 bg=0.02 p=1.5E-02 p=1.8F-02fg=0.02 n=2bg = 0.00fg=0.05 bg=0.01 ligand—gated channel activity GO:001270
ligand—gated channel activity GO:0022834
ubiquitin–like protein conjugating enzym... GO:0044390
calcium channel activity GO:0005262
calcium—release channel activity GO:0015278
outward rectifier rotassium channel acti p=1.9F-02fa=0.05 n=4ba = 0.01bğ=0.01 p=2.2E-02 fg=0.03 n=2.4F-0.2fg=0.05 n=4bg=0.01 p=2.4E n=2bg=0.00 outward rectifier potassium channel activity GO:0015278
outward rectifier potassium channel acti... GO:0015271
ligand–gated cation channel activity GO:0099094
siRNA binding GO:0035197
N–acetylglucosamine kinase activity GO:0045127
N–acylmannosamine kinase activity GO:0046624 ∥fg=0.02 p=3.2E-02n=2bg=0.00 p=3.4E-02 p=3.6E-02 bğ=0.01 n=3fg=0.01 fg=0.01 n=1bg=0.00 p=3.6E-02 bg=0.00 n=1p=3.6E-02 n=1 fg=0.01 bg=0.00 p=3.6E-02 fg=0.01 bg=0.00 n=1uridine–diphosphatase activity GO:0045134
fatty acid alpha–hydroxylase activity GO:0080132
glycerophosphodiester phosphodiesterase ... GO:0008889
rRNA (guanosine–2'–O–)–methyltransferase... GO:0070039
transmembrane receptor protein phosphata... GO:0019198 fg=0.01 p=3.6E-02bg=0.00 n=1p=3.6E-02 fg=0.01 bg=0.00 n=1n=1 fg=0.01 p=3.6E-02 bg=0.00 bg=0.00 bg=0.00 fg=0.01 p=3.6E-02 n=1p=3.6E-02 p=3.6E-02 fg=0.01 fg=0.01 n=1 muscle alpha–actinin binding GO:0051371 procollagen–proline 4–dioxygenase activi... GO:0004656 n=1bq = 0.00fg=0.01 p=3.6E-02 bg=0.00 2 0.0 1 3 5 0.2 0.4 0.6 8.0 1.0 fraction –log(p)

Ncol_Nvec_vc1.1_XM_032366646.2

GO:CC Ncol_Nvec_vc1.1_XM_032366646.2 fracti

n=86/305 input genes with annotations

fraction genes in fg and expected valu

