



7/14 ATPase–coupled intramembrane lipid transporter  
52/178 primary active transmembrane transporter  
6/10 apolipoprotein receptor  
3/18 sphingolipid transporter  
23/75 syntaxin binding  
31/106 SNARE binding  
9/21 SNAP receptor  
5/6 protein transmembrane transporter  
13/42 transition metal ion transmembrane transporter  
12/21 muscle alpha–actinin binding  
24/51 actinin binding  
343/909 cytoskeletal protein binding  
165/409 actin binding  
69/164 actin filament binding  
17/35 structural constituent of muscle  
14/29 ankyrin binding  
7/17 laminin–1 binding  
402/1211 calcium ion binding  
16/48 dystroglycan binding  
11/13 myosin heavy chain binding  
23/62 myosin binding  
21/47 extracellular matrix structural constituent conferring tensile strength  
143/400 extracellular matrix structural constituent  
263/791 structural molecule  
30/146 structural constituent of ribosome  
19/45 ErbB–2 class receptor binding  
91/280 motor  
5/14 nuclear export signal receptor  
14/49 signal sequence binding  
11/42 structural constituent of nuclear pore  
534/1437 RNA binding  
5/6 second spliceosomal transesterification  
20/39 snRNA binding  
67/174 mRNA binding  
24/74 rRNA binding  
6/6 5S rRNA binding  
15/33 snoRNA binding  
50/101 translation regulator, nucleic acid binding  
35/58 translation initiation factor  
64/138 translation regulator  
12/23 translation initiation factor binding  
25/50 ribosome binding  
49/119 ribonucleoprotein complex binding  
114/427 catalytic, acting on RNA  
15/45 DNA–directed 5'–3' RNA polymerase  
12/64 RNA methyltransferase  
29/123 catalytic, acting on a tRNA  
57/149 transcription coactivator  
88/260 transcription coregulator  
6/11 ATP–dependent peptidase  
13/25 ATPase regulator  
7/12 ATPase activator  
43/102 unfolded protein binding  
38/97 ubiquitin–like protein binding  
112/286 ubiquitin–like protein ligase binding  
12/14 threonine–type endopeptidase  
41/122 organic acid binding  
17/43 L–ascorbic acid binding  
113/306 carbohydrate binding  
6/8 galactose binding  
42/99 monosaccharide binding  
5/5 transferase, transferring aldehyde or ketonic groups  
4/5 protein–glucosylgalactosylhydroxylsine glucosidase  
70/193 hydrolase, acting on glycosyl bonds  
13/35 dipeptidase  
131/371 metallopeptidase  
10/30 dipeptidyl–peptidase  
53/171 exopeptidase  
13/23 peptidase activator  
32/101 antiporter  
7/18 nucleotide–sugar transmembrane transporter  
26/83 solute:sodium symporter  
77/196 symporter  
20/42 ion gated channel  
12/20 intracellular chloride channel  
11/14 phospholipid scramblase  
22/44 delayed rectifier potassium channel  
39/100 voltage–gated potassium channel  
56/158 potassium ion transmembrane transporter  
5/7 glutathione disulfide oxidoreductase  
30/98 electron transfer  
9/20 protein disulfide oxidoreductase  
6/29 FMN binding  
11/17 nucleoside kinase  
7/9 L–alanine:2–oxoglutarate aminotransferase  
6/11 3',5'–cyclic–AMP phosphodiesterase  
16/61 sulfotransferase  
7/8 proteasome–activating ATPase  
4/6 oxidoreductase, acting on single donors with incorporation of molecular oxygen, incorporation of one atom of oxygen (internal monooxygenases or internal mixed function oxidases)  
4/6 peroxiredoxin  
9/12 MATH domain binding

p < 0.001  
p < 0.01  
p < 0.05