7/42 translation initiation factor 95/424 RNA binding p < 0.014/5 peroxiredoxin p < 0.0566/289 structural molecule p < 0.126/103 structural constituent of ribosome 13/50 structural constituent of cytoskeleton 6/8 myosin II binding 24/78 unfolded protein binding 5/15 calmodulin-dependent protein kinase 47/228 protein serine/threonine kinase 73/267 anion transmembrane transporter 3/8 sulfur amino acid transmembrane transporter 39/135 organic anion transmembrane transporter 139/582 ion transmembrane transporter 15/46 L-amino acid transmembrane transporter 9/25 neutral amino acid transmembrane transporter 11/23 sulfur compound transmembrane transporter 49/192 monovalent inorganic cation transmembrane transporter 9/33 organic acid:sodium symporter 28/110 sodium ion transmembrane transporter 17/61 symporter 39/131 secondary active transmembrane transporter 4/6 sulfate transmembrane transporter 69/222 active transmembrane transporter 5/12 anion: anion antiporter 16/51 inorganic anion transmembrane transporter 3/5 inorganic anion exchanger 8/28 chloride transmembrane transporter 16/68 voltage-gated ion channel 57/268 passive transmembrane transporter 31/172 ion gated channel 15/107 ligand-gated ion channel 98/431 cation transmembrane transporter 31/165 divalent inorganic cation transmembrane transporter 64/305 metal ion transmembrane transporter 2/16 ATP-dependent microtubule motor 22/86 motor 7/7 aldehyde dehydrogenase [NAD(P)+] 5/5 aldehyde dehydrogenase (NAD+) 9/25 calcium-dependent phospholipid binding 3/7 SUMO transferase 5/11 cyclic-di-GMP binding