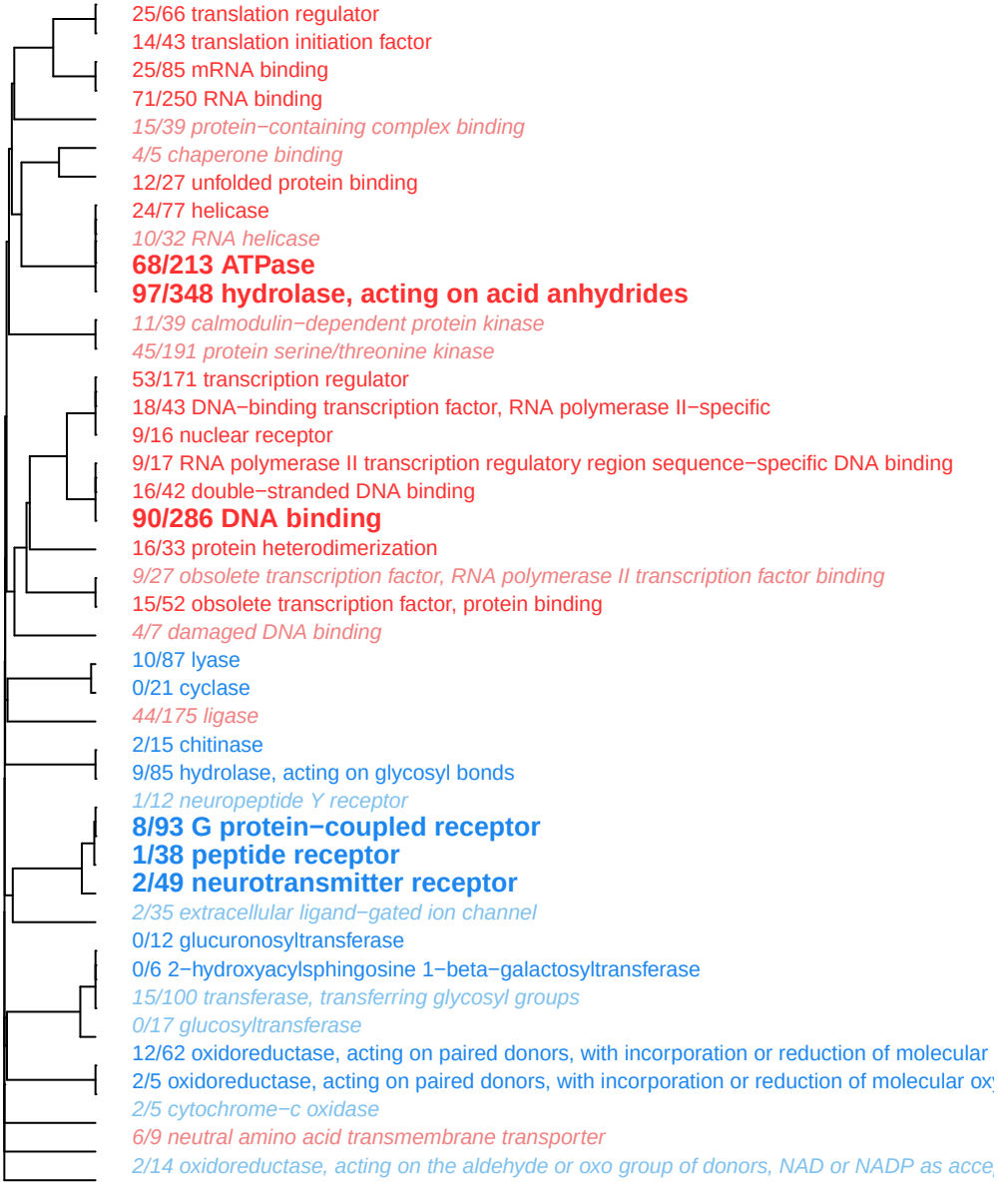
25/66 translation regulator
14/43 translation initiation factor
25/85 mRNA binding
71/250 RNA binding
15/39 protein−containing complex binding
4/5 chaperone binding
12/27 unfolded protein binding
24/77 helicase
10/32 RNA helicase
68/213 ATPase
97/348 hydrolase, acting on acid anhydrides
11/39 calmodulin−dependent protein kinase
45/191 protein serine/threonine kinase
53/171 transcription regulator
18/43 DNA−binding transcription factor, RNA polymerase II−specific
9/16 nuclear receptor
9/17 RNA polymerase II transcription regulatory region sequence−specific DNA binding
16/42 double−stranded DNA binding
90/286 DNA binding
16/33 protein heterodimerization
9/27 obsolete transcription factor, RNA polymerase II transcription factor binding
15/52 obsolete transcription factor, protein binding
4/7 damaged DNA binding
10/87 lyase
0/21 cyclase
44/175 ligase
2/15 chitinase
9/85 hydrolase, acting on glycosyl bonds
1/12 neuropeptide Y receptor
8/93 G protein−coupled receptor
1/38 peptide receptor
2/49 neurotransmitter receptor
2/35 extracellular ligand−gated ion channel
0/12 glucuronosyltransferase
0/6 2−hydroxyacylsphingosine 1−beta−galactosyltransferase
15/100 transferase, transferring glycosyl groups
0/17 glucosyltransferase
12/62 oxidoreductase, acting on paired donors, with incorporation or reduction of molecular
2/5 oxidoreductase, acting on paired donors, with incorporation or reduction of molecular ox
2/5 cytochrome−c oxidase
6/9 neutral amino acid transmembrane transporter
2/14 oxidoreductase, acting on the aldehyde or oxo group of donors, NAD or NADP as acce
p < 0.01
p < 0.05
p < 0.1