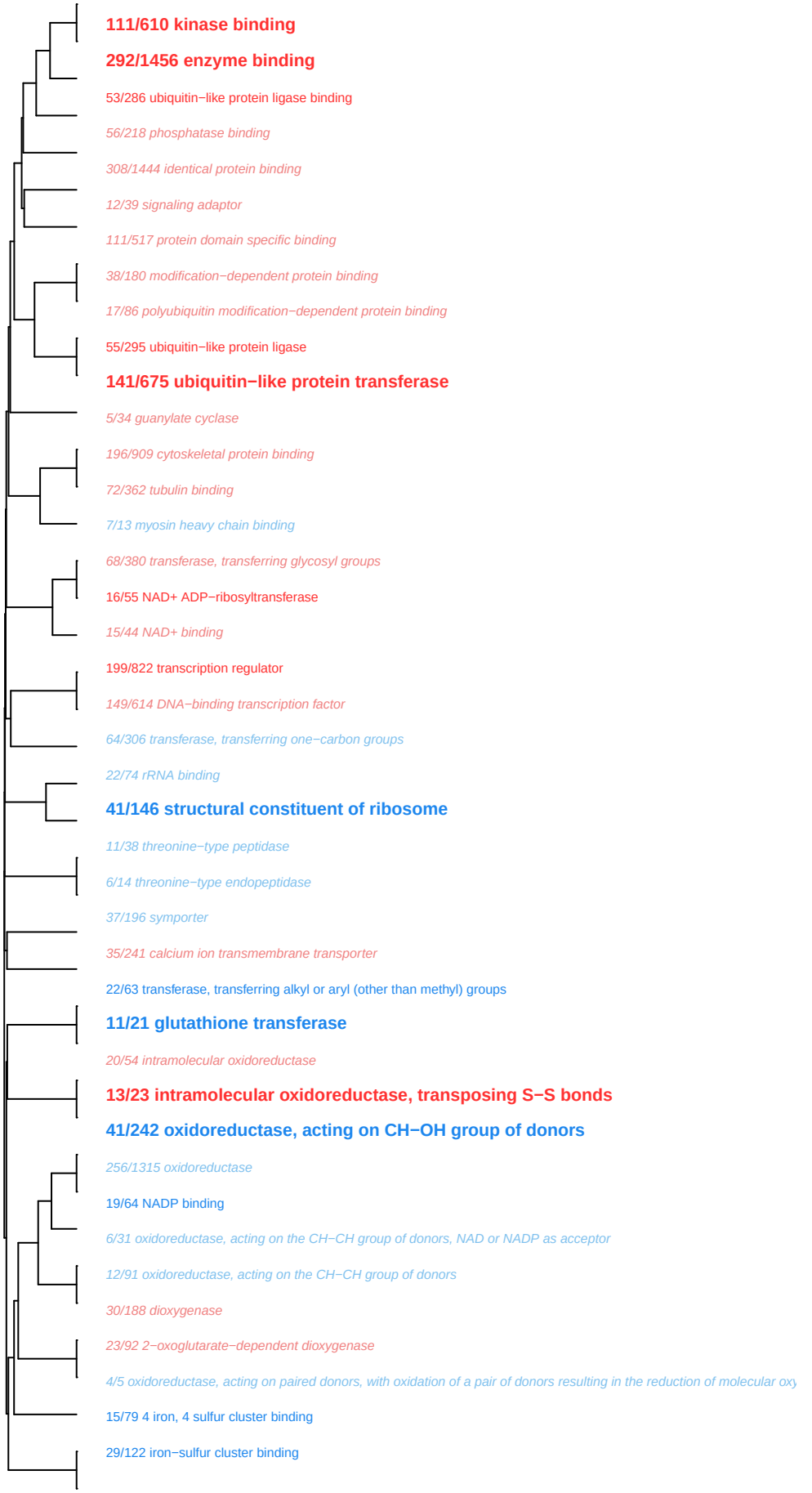
111/610 kinase binding
292/1456 enzyme binding
53/286 ubiquitin−like protein ligase binding
56/218 phosphatase binding
308/1444 identical protein binding
12/39 signaling adaptor
111/517 protein domain specific binding
38/180 modification−dependent protein binding
17/86 polyubiquitin modification−dependent protein binding
55/295 ubiquitin−like protein ligase
141/675 ubiquitin−like protein transferase
5/34 guanylate cyclase
196/909 cytoskeletal protein binding
72/362 tubulin binding
7/13 myosin heavy chain binding
68/380 transferase, transferring glycosyl groups
16/55 NAD+ ADP−ribosyltransferase
15/44 NAD+ binding
199/822 transcription regulator
149/614 DNA−binding transcription factor
64/306 transferase, transferring one−carbon groups
22/74 rRNA binding
41/146 structural constituent of ribosome
11/38 threonine−type peptidase
6/14 threonine−type endopeptidase
37/196 symporter
35/241 calcium ion transmembrane transporter
22/63 transferase, transferring alkyl or aryl (other than methyl) groups
11/21 glutathione transferase
20/54 intramolecular oxidoreductase
13/23 intramolecular oxidoreductase, transposing S−S bonds
41/242 oxidoreductase, acting on CH−OH group of donors
256/1315 oxidoreductase
19/64 NADP binding
6/31 oxidoreductase, acting on the CH−CH group of donors, NAD or NADP as acceptor
12/91 oxidoreductase, acting on the CH−CH group of donors
30/188 dioxygenase
23/92 2−oxoglutarate−dependent dioxygenase
4/5 oxidoreductase, acting on paired donors, with oxidation of a pair of donors resulting in the reduction of molecular oxy
15/79 4 iron, 4 sulfur cluster binding
29/122 iron−sulfur cluster binding
p < 0.001
p < 0.01
p < 0.05