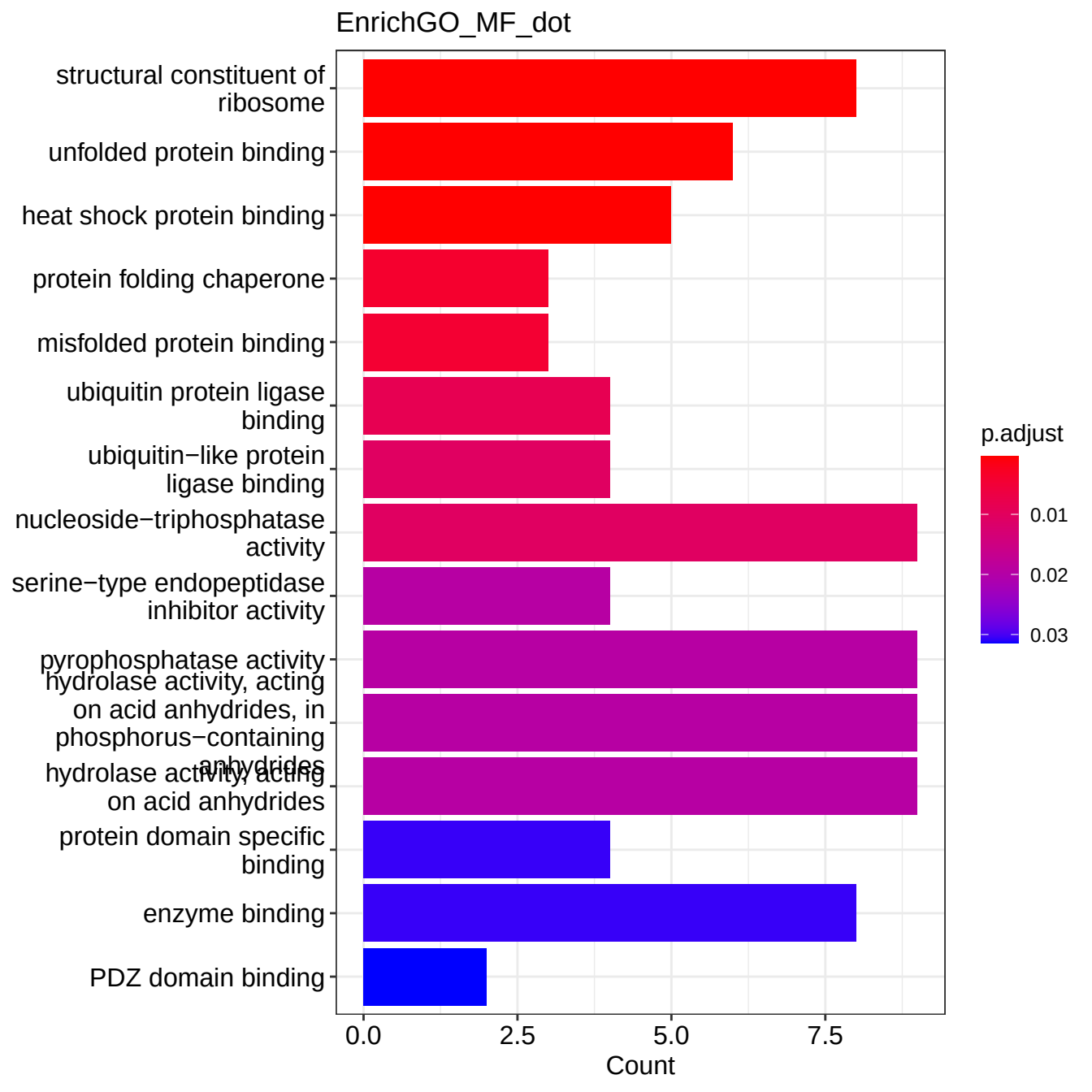
EnrichGO_MF_dot
structural constituent of ribosome
unfolded protein binding
heat shock protein binding
protein folding chaperone
misfolded protein binding
ubiquitin protein ligase binding
ubiquitin−like protein ligase binding
nucleoside−triphosphatase activity
serine−type endopeptidase inhibitor activity
pyrophosphatase activity
hydrolase activity, acting on acid anhydrides, in phosphorus−containing anhydrides
hydrolase activity, acting on acid anhydrides
protein domain specific binding
enzyme binding
PDZ domain binding
0.0
2.5
5.0
7.5
Count
p.adjust
0.01
0.02
0.03