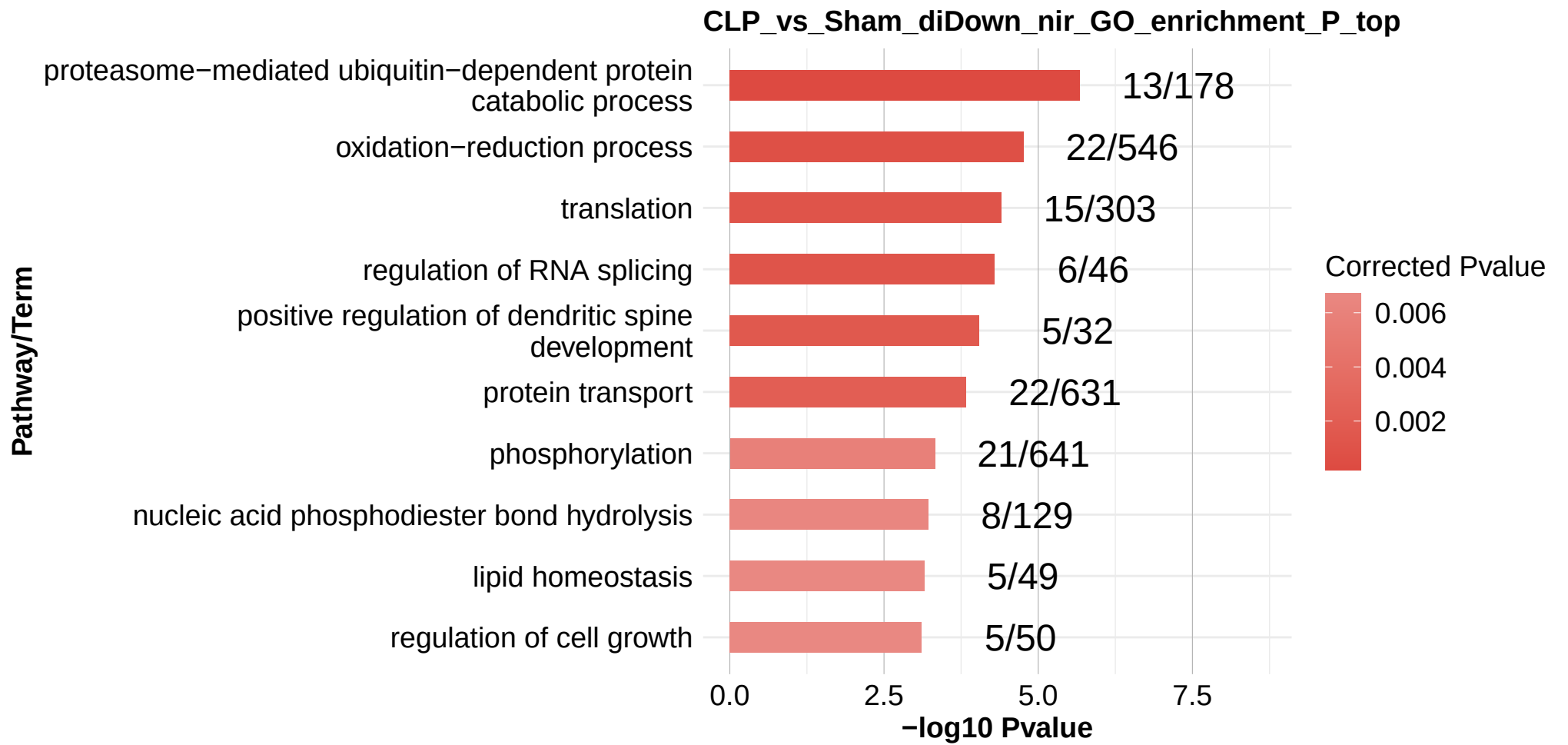
CLP_vs_Sham_diDown_nir_GO_enrichment_P_top
proteasome−mediated ubiquitin−dependent protein catabolic process
oxidation−reduction process
translation
regulation of RNA splicing
positive regulation of dendritic spine development
protein transport
phosphorylation
nucleic acid phosphodiester bond hydrolysis
lipid homeostasis
regulation of cell growth
13/178
22/546
15/303
6/46
5/32
22/631
21/641
8/129
5/49
5/50
Pathway/Term
0.0
2.5
5.0
7.5
−log10 Pvalue
Corrected Pvalue
0.006
0.004
0.002