LM_GO_Molecular_Funct

Enriched terms (top to bottom):
- ubiquitin protein ligase binding (GO:003...)
- ubiquitin−like protein ligase binding (G...)
- RNA binding (GO:0003723)
- ATPase regulator activity (GO:0060590)
- ATPase activator activity (GO:0001671)
- C3HC4−type RING finger domain binding (G...)
- disordered domain specific binding (GO:0...)
- Hsp90 protein binding (GO:0051879)
- purine ribonucleoside triphosphate bindi...
- cadherin binding (GO:0045296)
- adenyl ribonucleotide binding (GO:003255...)
- kinase binding (GO:0019900)
- chemokine activity (GO:0008009)
- ATP binding (GO:0005524)
- chemokine receptor binding (GO:0042379)
- double−stranded RNA binding (GO:0003725)
- MHC class II protein complex binding (GO...)
- DNA polymerase binding (GO:0070182)
- deoxycytidine deaminase activity (GO:004...)
- CCR chemokine receptor binding (GO:00480...)

X-axis: Gene count (0, 10, 20, 30, 40)

P value legend: 1e−04, 2e−04, 3e−04