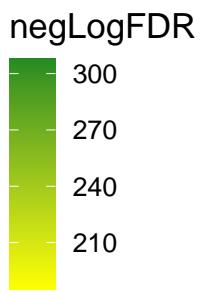


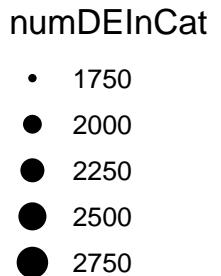
# GOseq significant terms (FDR = 0.05) — Depth 1 — MF

GO Term

binding



catalytic activity



0.00

0.05

0.10

0.15

Fold Enrichment