

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

GO Term

binding

0.00

0.05

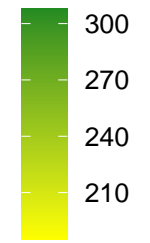
Fold Enrichment

0.10

0.15

catalytic activity

negLogFDR



numDEInCat

