

GOseq significant terms (FDR = 0.05) — Depth 8 — MF

GO Term

purine ribonucleoside triphospha...

ribonucleotide binding

0.00

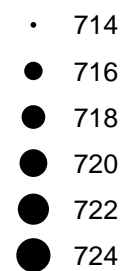
0.05

0.10

0.15

Fold Enrichment

numDEInCat



negLogFDR

