GO enrichment analysis transcription factor binding mRNA processing structural molecule activity chromosome organization isomerase activity -Gene_count 50 cytoskeletal protein binding -100 150 cytoskeleton organization -RNA binding brown_BP log_pvalue DNA-binding transcription factor activity -20 cellular nitrogen compound metabolic process -15 DNA binding -10 5 biological_process -GTPase activity oxidoreductase activity biosynthetic process ion binding -

0.0 0.1 0.2 0.3 0.4 0.5

0.0 0.1 0.2 0.3 0.4 0.5