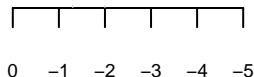
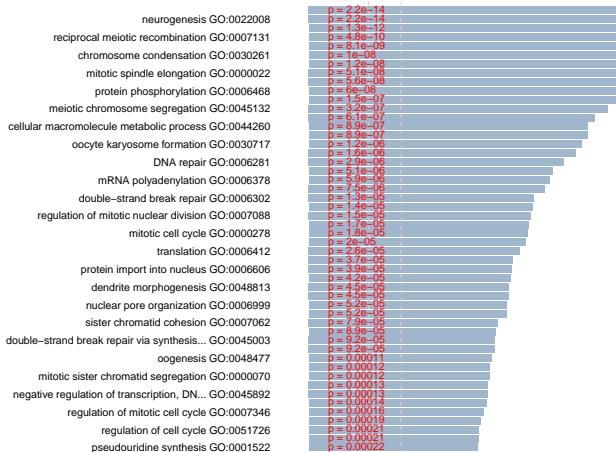


ALL.RUN-SNO_difneg top GO: BP fisher elim

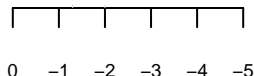
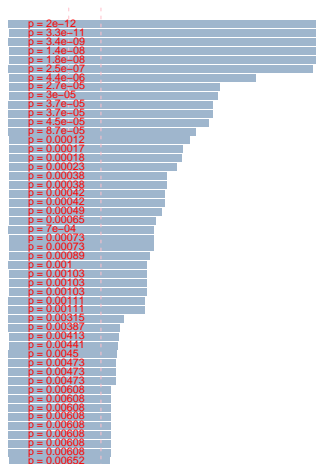


log(p)

n=1532/2666

ALL.RUN-SNO_difneg top GO: MF fisher elin

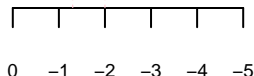
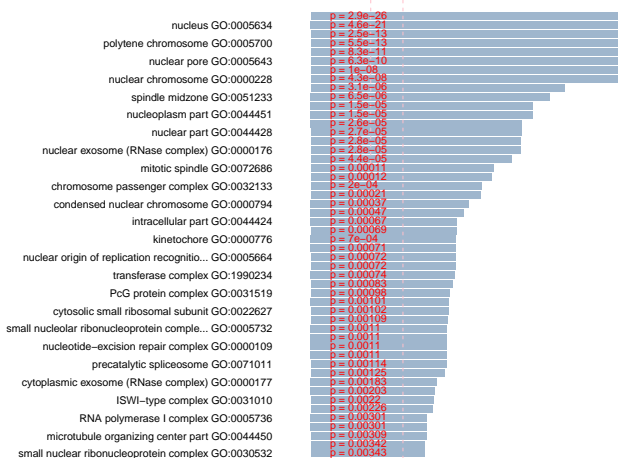
ATP-dependent RNA helicase activity GO:0004004
DNA-directed DNA polymerase activity GO:0003887
DNA binding GO:0003677
catalytic activity, acting on DNA GO:0140097
Y-form DNA binding GO:0000403
POZ domain binding GO:0031208
Ran GTPase binding GO:0008536
3'-5'-exoribonuclease activity GO:0000175
structural constituent of ribosome GO:0003735
cyclin-dependent protein serine/threonin... GO:0016538
helicase activity GO:0004386
sequence-specific DNA binding GO:0043565
endoribonuclease activity, producing 5'-... GO:0016891
ATP binding GO:0005524
3'-5'-exodeoxyribonuclease activity GO:0008296
endodeoxyribonuclease activity GO:0004520
DNA-directed 5'-3' RNA polymerase activ... GO:0003899
sequence-specific double-stranded DNA bi... GO:1990837
microtubule binding GO:0008017
polynucleotide adenyltransferase activ... GO:0004652
endonuclease activity, active with eithe... GO:0016894
guanine/thymine mispair binding GO:0032137
bent DNA binding GO:0003681
5S rRNA binding GO:0008097
S-adenosylmethionine-dependent methyltra... GO:0008757



log(p)

n=1344/2666

ALL.RUN-SNO_difneg top GO: CC fisher elin



log(p)

n=1117/2666