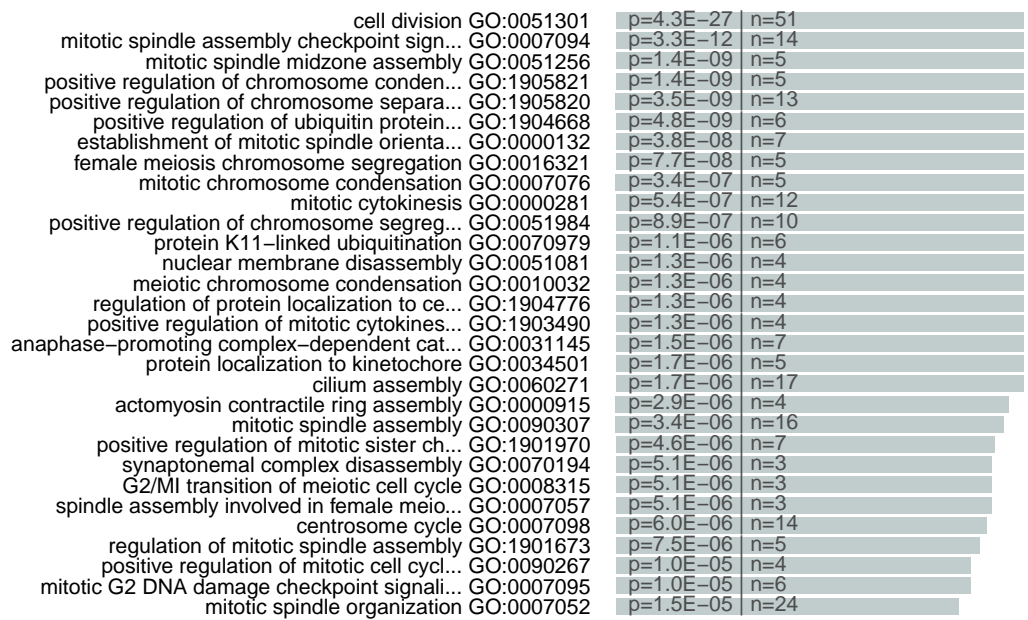
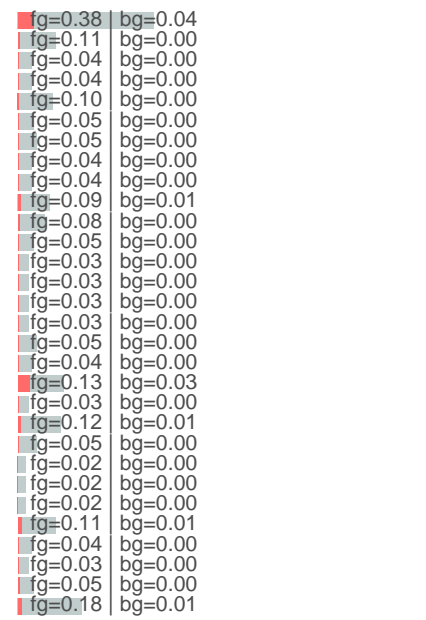


GO:BP
gc021

fraction genes in fg and expected value



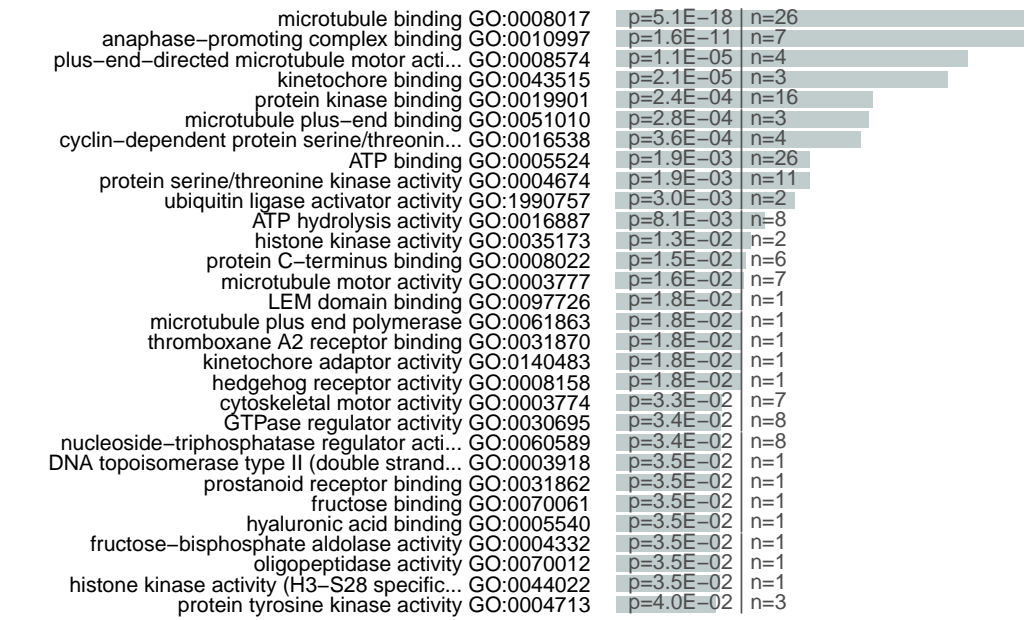
0 1 2 3 4 5
-log(p)
n=133/193 input genes with annotations



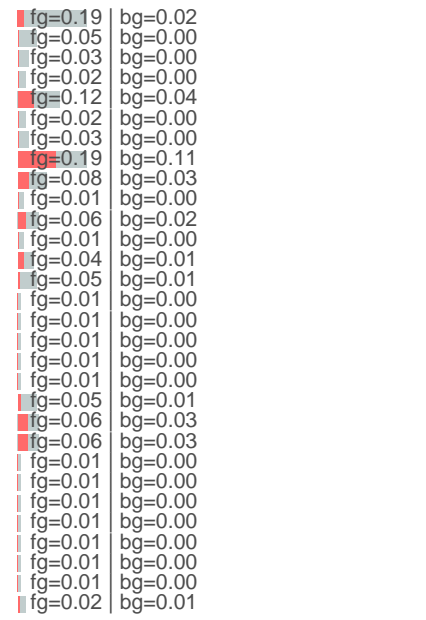
0.0 0.2 0.4 0.6 0.8 1.0
fraction

GO:MF
gc021

fraction genes in fg and expected value



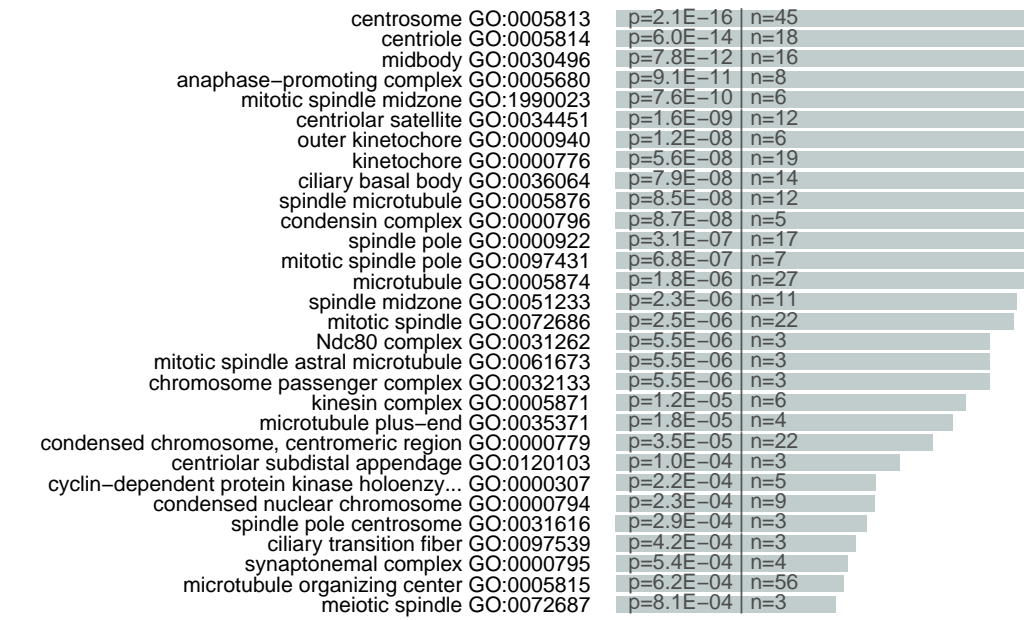
0 1 2 3 4 5
-log(p)
n=135/193 input genes with annotations



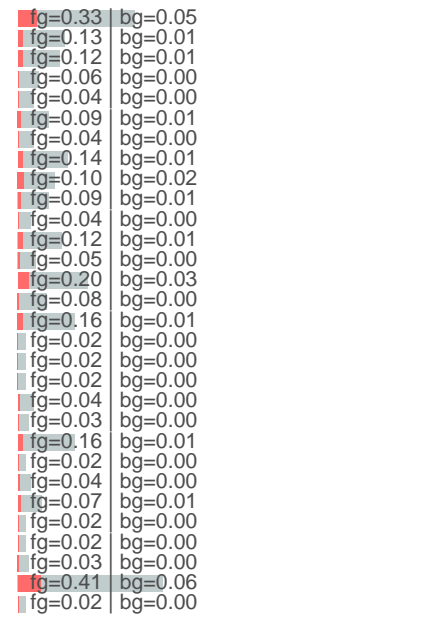
0.0 0.2 0.4 0.6 0.8 1.0
fraction

GO:CC
gc021

fraction genes in fg and expected value



0 1 2 3 4 5
-log(p)
n=137/193 input genes with annotations



0.0 0.2 0.4 0.6 0.8 1.0
fraction