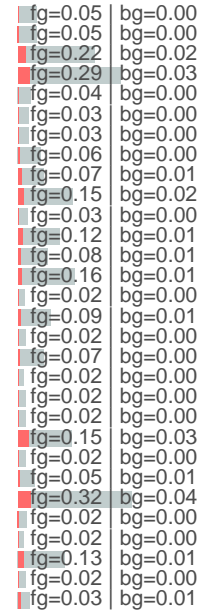
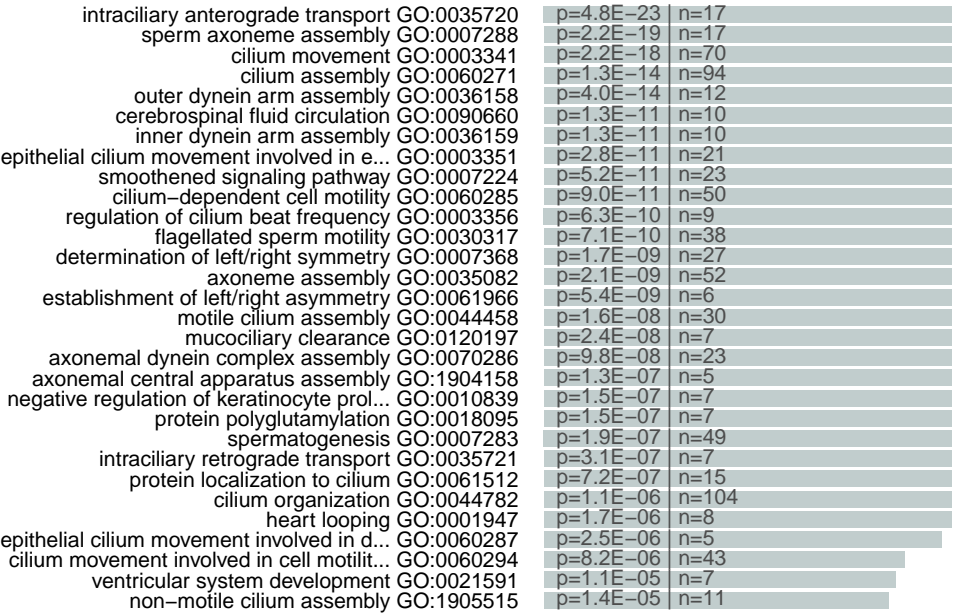


GO:BP
gc030

fraction genes in fg and expected value

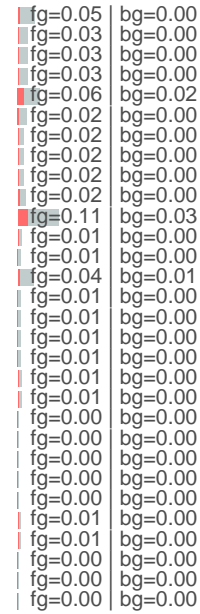
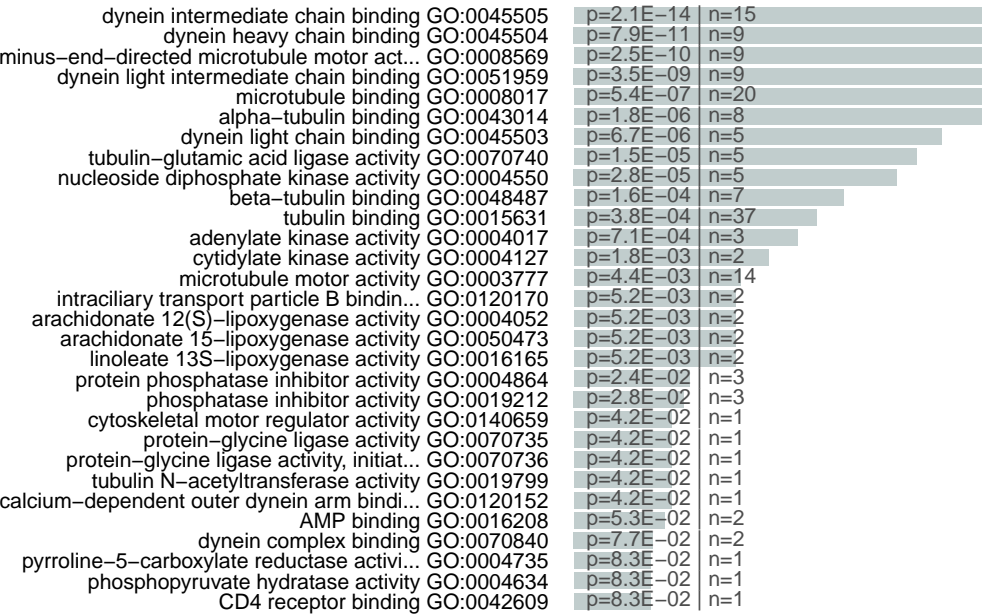


-log(p)
n=324/468 input genes with annotations

fraction

GO:MF
gc030

fraction genes in fg and expected value

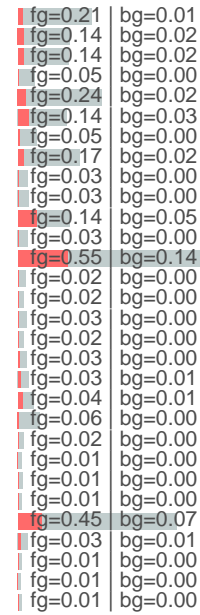
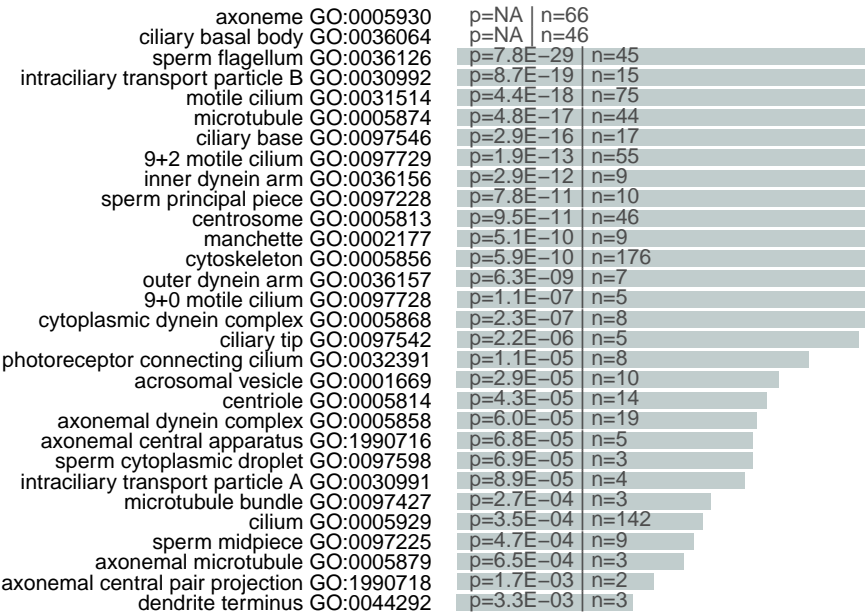


-log(p)
n=324/468 input genes with annotations

fraction

GO:CC
gc030

fraction genes in fg and expected value



-log(p)
n=318/468 input genes with annotations

fraction