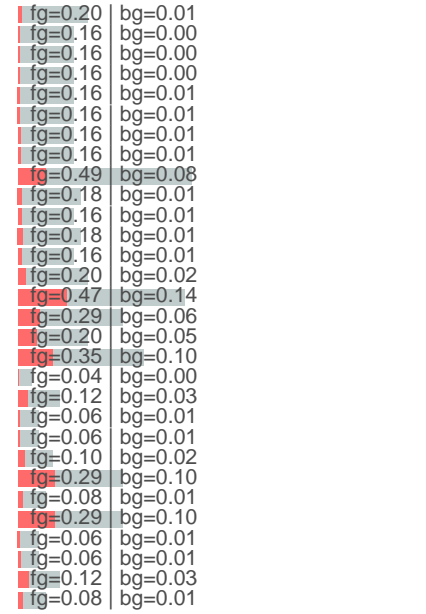
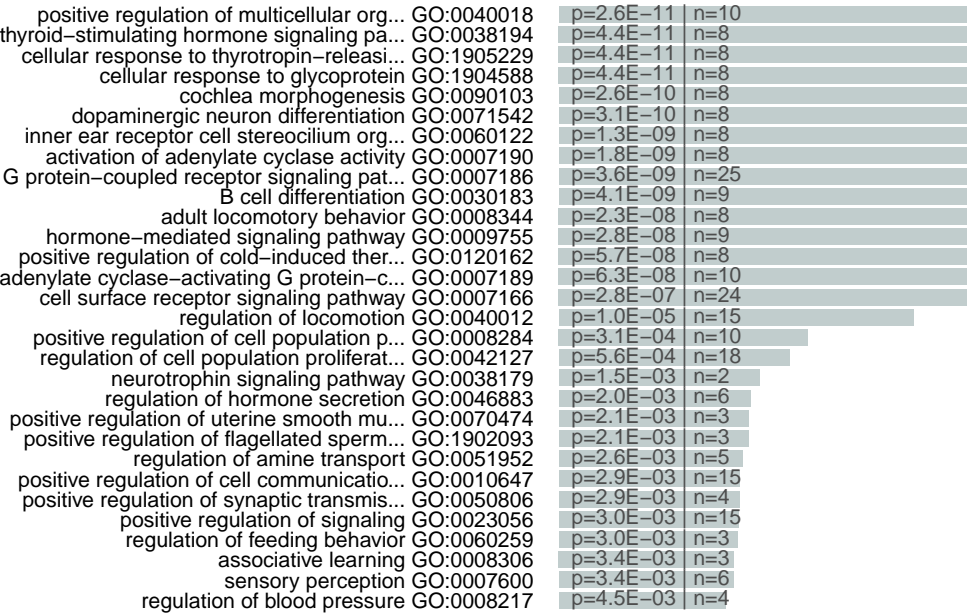


GO:BP
gc006

fraction genes in fg and expected value

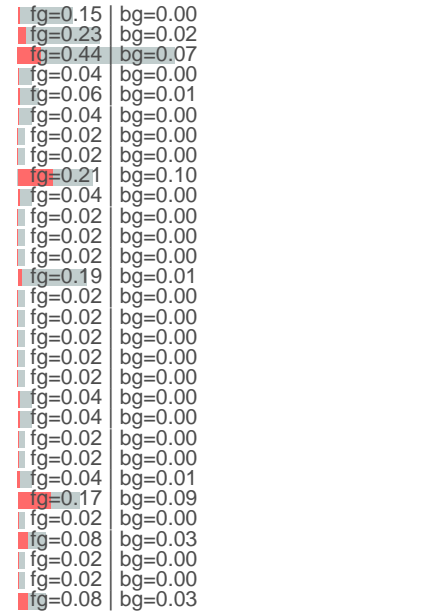
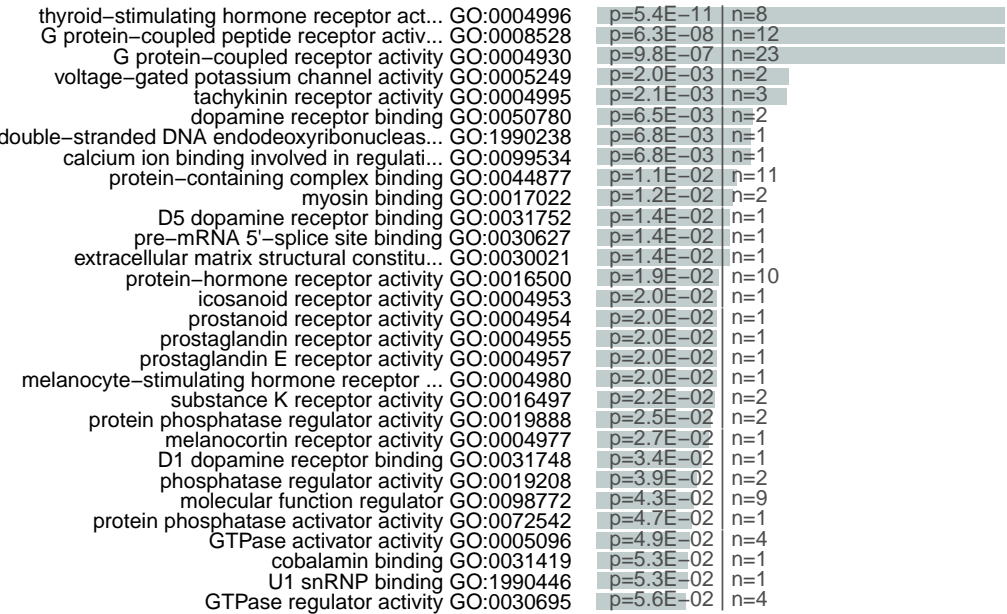


-log(p)
n=51/182 input genes with annotations

fraction

GO:MF
gc006

fraction genes in fg and expected value

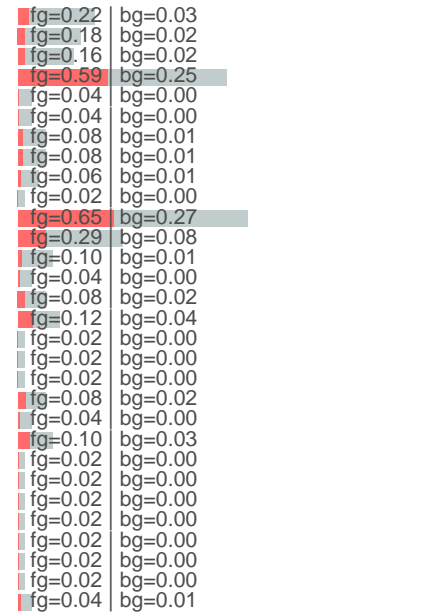
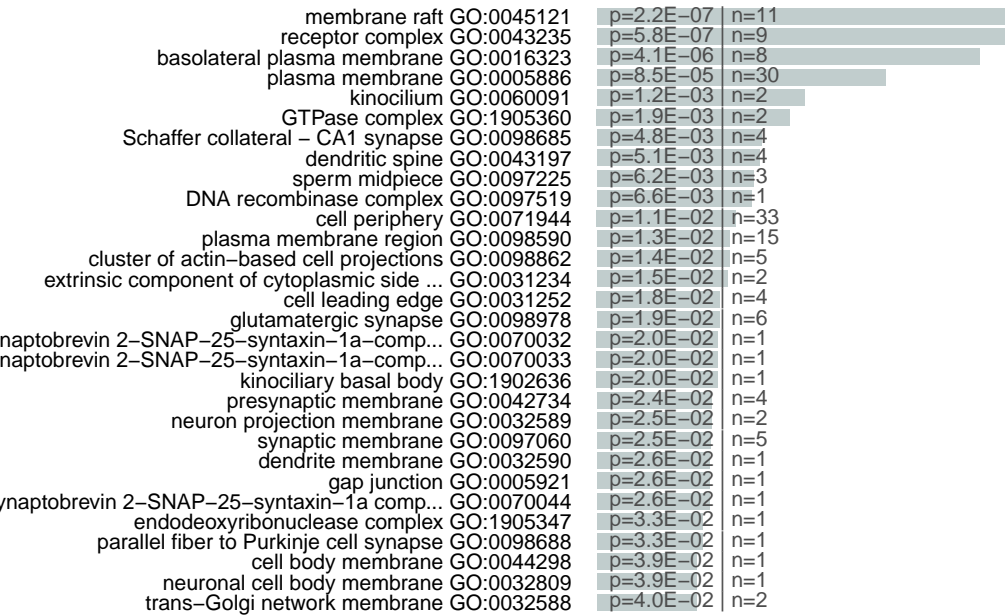


-log(p)
n=52/182 input genes with annotations

fraction

GO:CC
gc006

fraction genes in fg and expected value



-log(p)
n=51/182 input genes with annotations

fraction