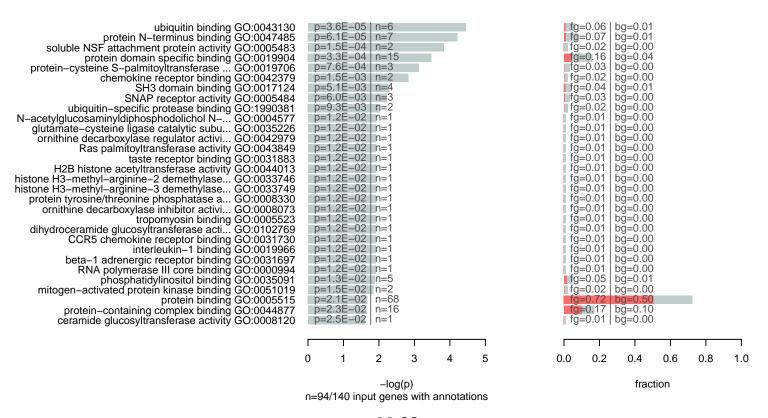


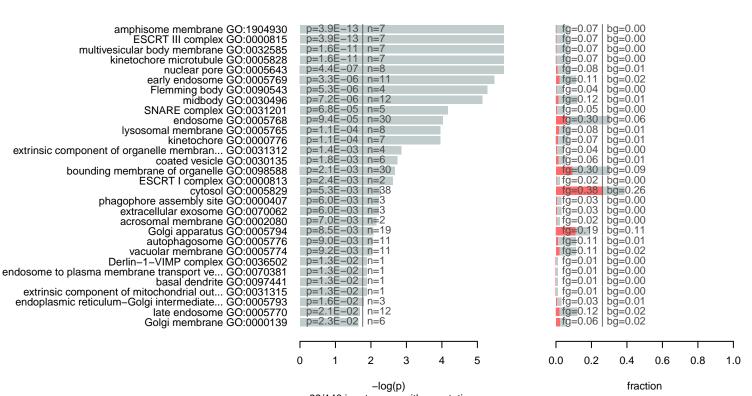
GO:MF gc032

fraction genes in fg and expected value



GO:CC gc032

fraction genes in fg and expected value



n=99/140 input genes with annotations