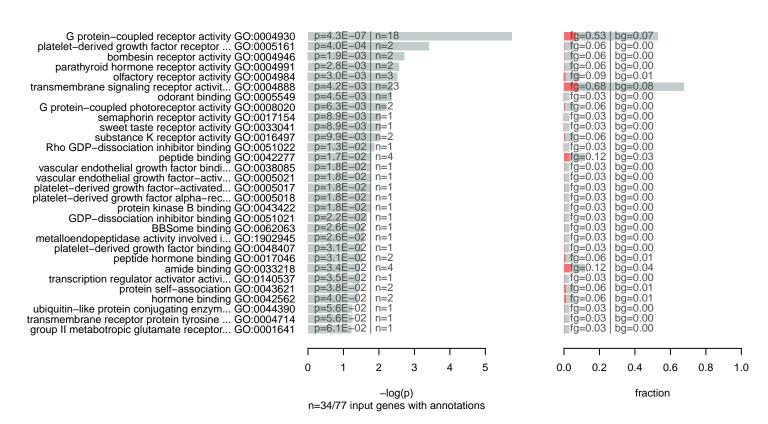


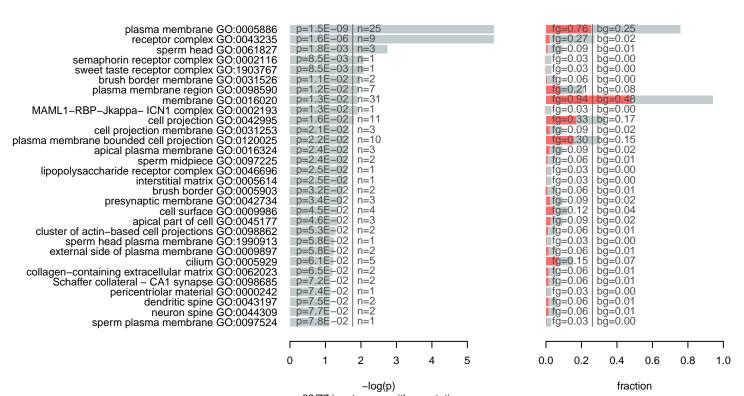
GO:MF gc014

fraction genes in fg and expected value



GO:CC gc014

fraction genes in fg and expected value



n=33/77 input genes with annotations