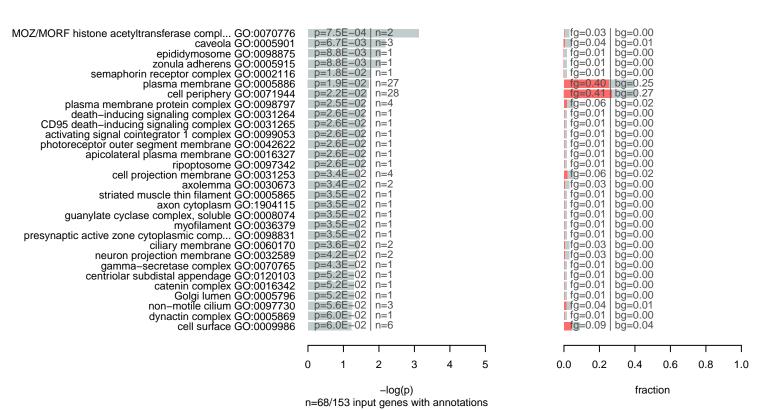


GO:CC gc023 fraction genes in fg and expected value

fraction



-log(p)
n=68/153 input genes with annotations