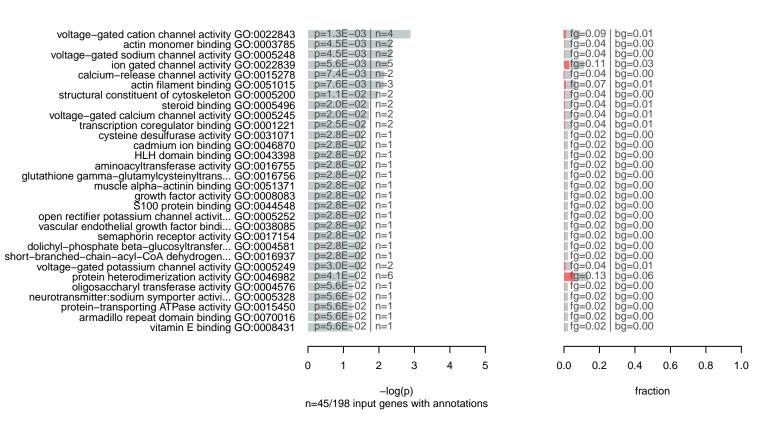


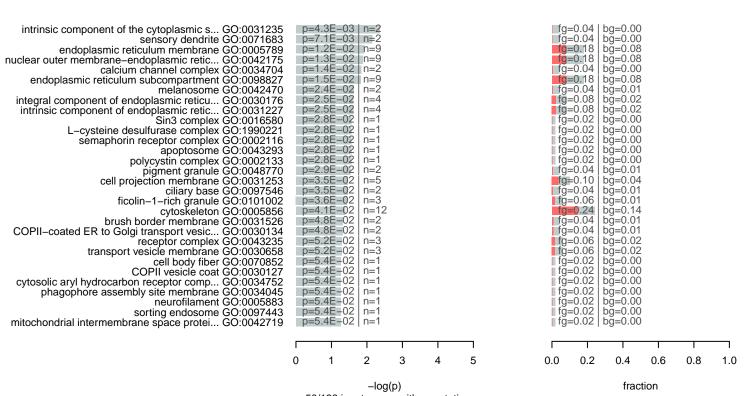
GO:MF Ncol\_Nvec\_vc1.1\_XM\_032376608.2

fraction genes in fg and expected value



GO:CC Ncol\_Nvec\_vc1.1\_XM\_032376608.2

fraction genes in fg and expected value



n=50/198 input genes with annotations