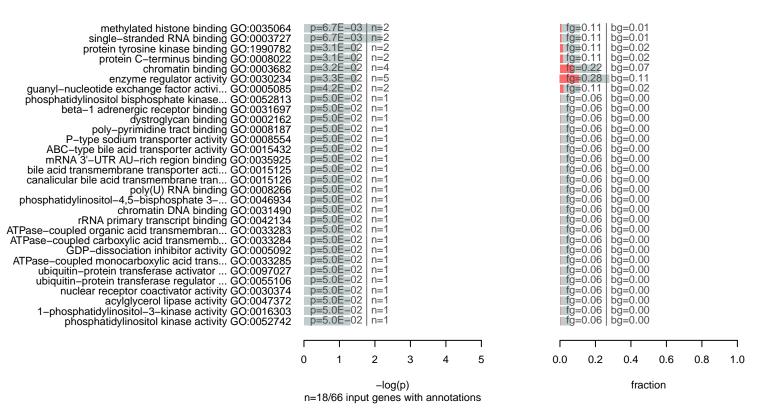
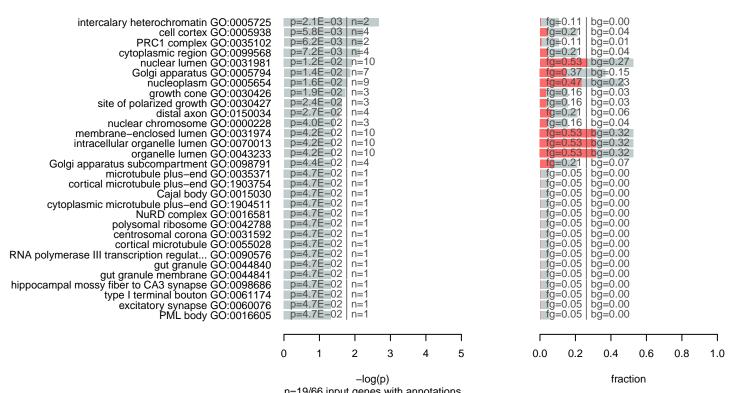


GO:MF fraction genes in fg and expected value Elav_Nvec_vc1.1_XM_001636587.3



GO:CC fraction genes in fg and expected value Elav_Nvec_vc1.1_XM_001636587.3



n=19/66 input genes with annotations