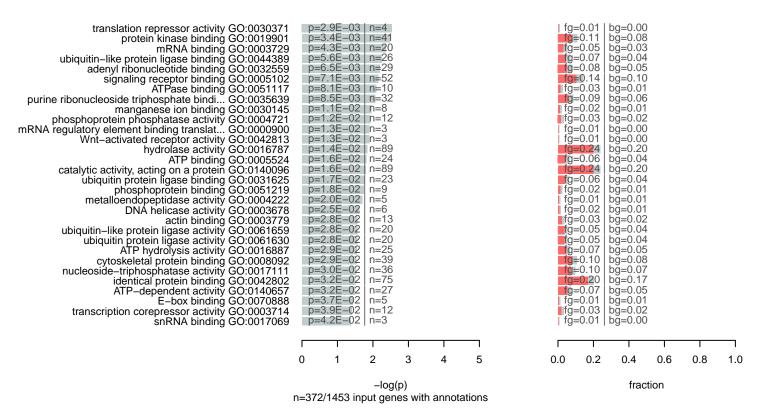
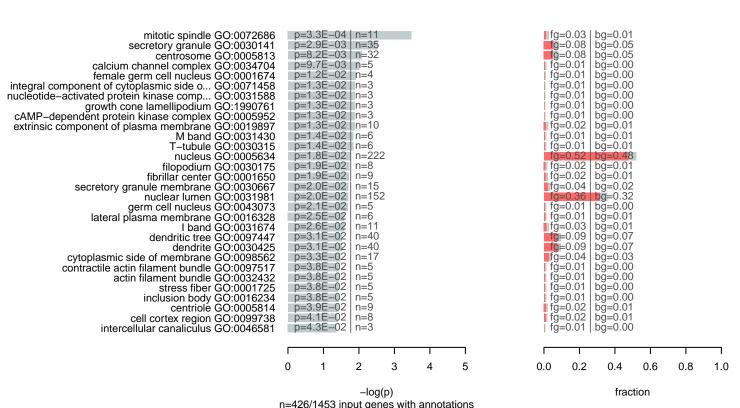


GO:MF fraction genes in fg and expected value Elav_Nvec_vc1.1_XM_032364887.2



GO:CC fraction genes in fg and expected value Elav_Nvec_vc1.1_XM_032364887.2



n=426/1453 input genes with annotations