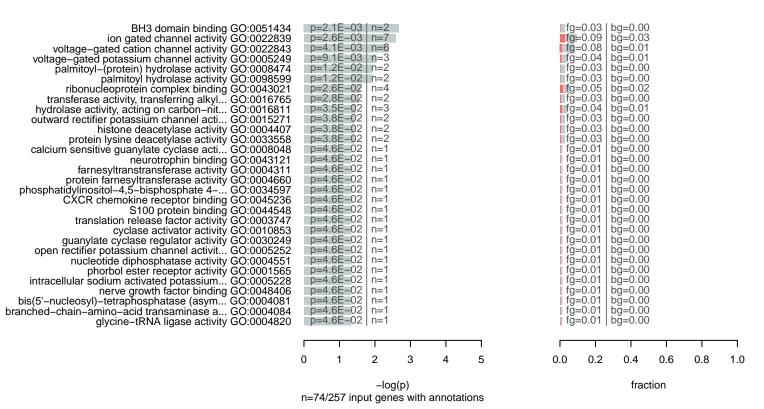
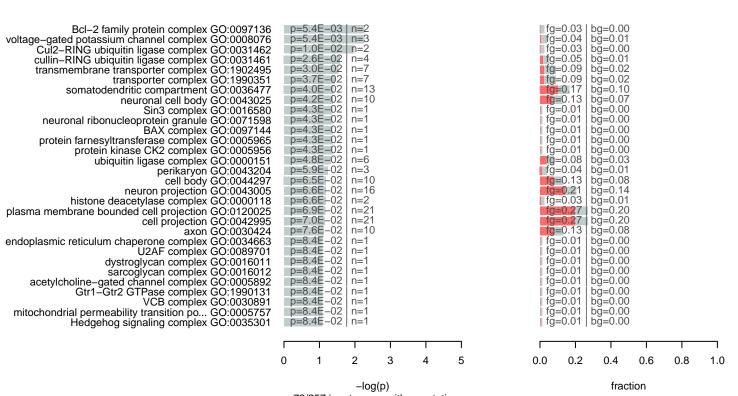


GO:MF fraction genes in fg and expected value Ncol_Nvec_vc1.1_XM_032381562.2



GO:CC fraction genes in fg and expected value Ncol_Nvec_vc1.1_XM_032381562.2



n=78/257 input genes with annotations