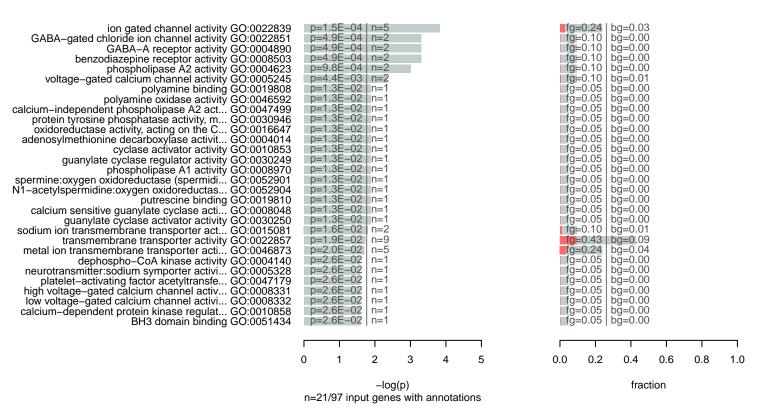
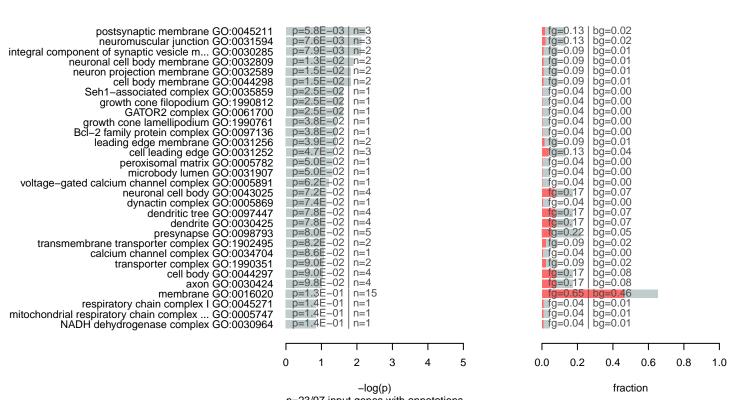


GO:MF fraction genes in fg and expected value Ncol_Nvec_vc1.1_XM_001630131.3



GO:CC fraction genes in fg and expected value Ncol_Nvec_vc1.1_XM_001630131.3



n=23/97 input genes with annotations