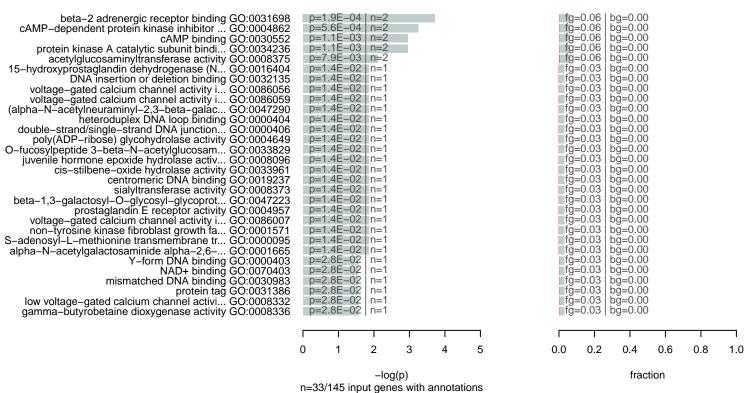
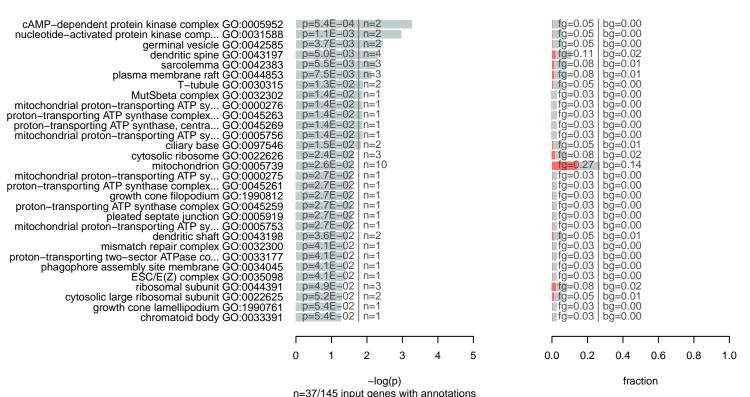


Elav_Nvec_vc1.1_XM_001632368.2



GO:CC fraction genes in fg and expected valu Elav_Nvec_vc1.1_XM_001632368.2



n=37/145 input genes with annotations