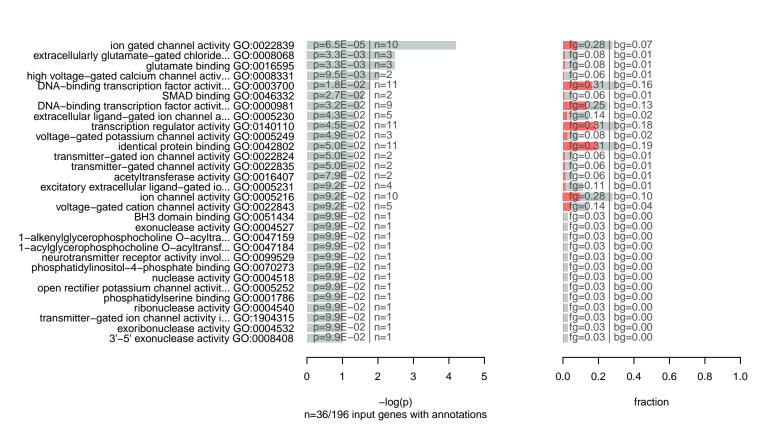
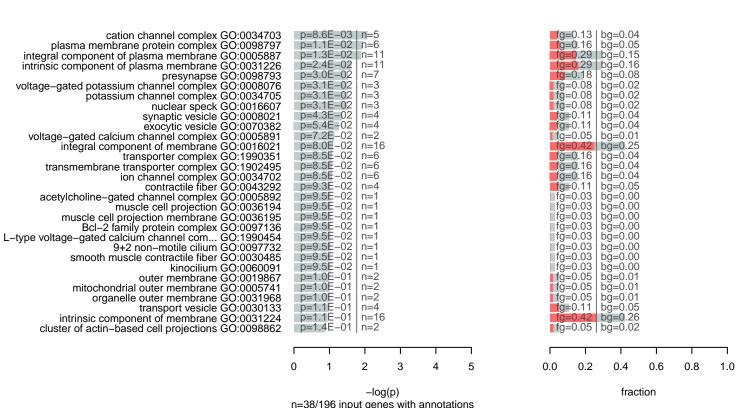


GO:MF fraction genes in fg and expected value Ncol\_Nvec\_vc1.1\_XM\_048728967.1



GO:CC fraction genes in fg and expected value Ncol\_Nvec\_vc1.1\_XM\_048728967.1



n=38/196 input genes with annotations