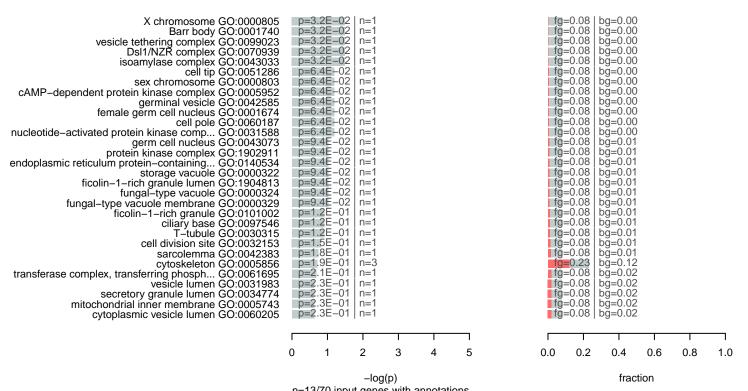


GO:CC fraction genes in fg and expected valu Fox_Nvec_vc1.1_XM_032376930.1



n=13/70 input genes with annotations