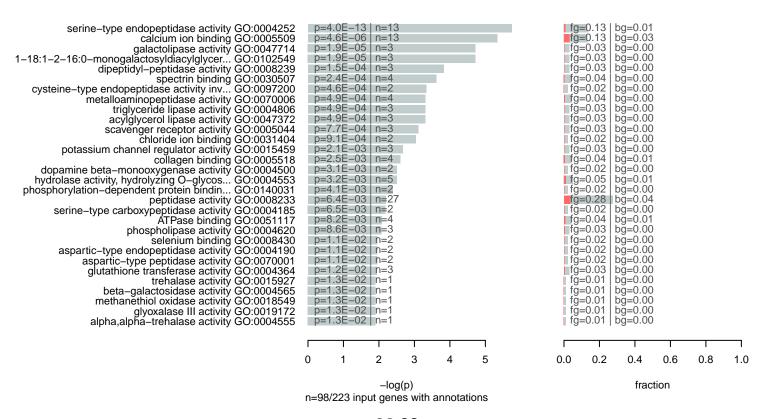


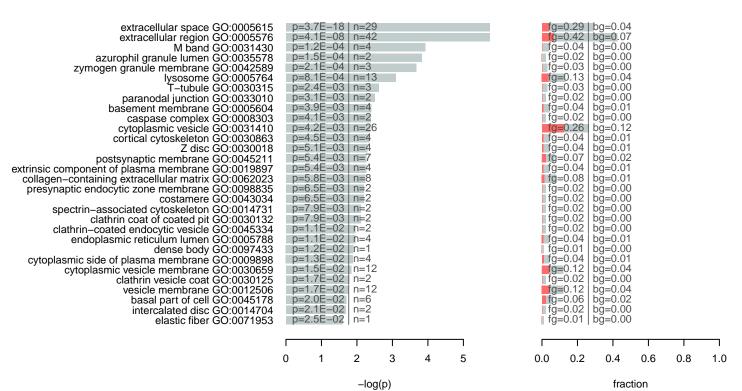
GO:MF gc022

fraction genes in fg and expected valu



GO:CC gc022

fraction genes in fg and expected value



n=99/223 input genes with annotations