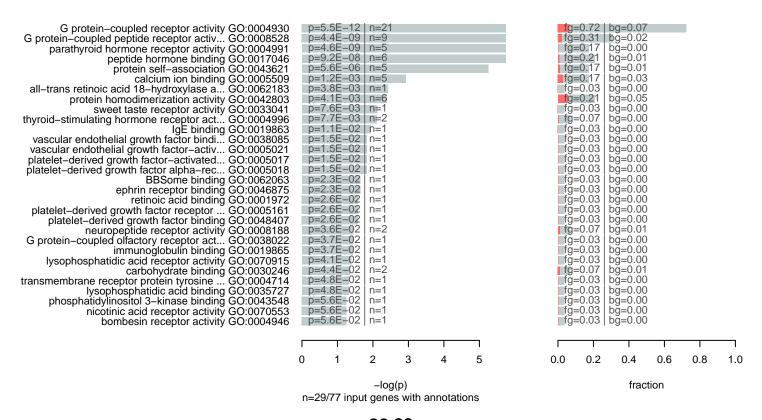


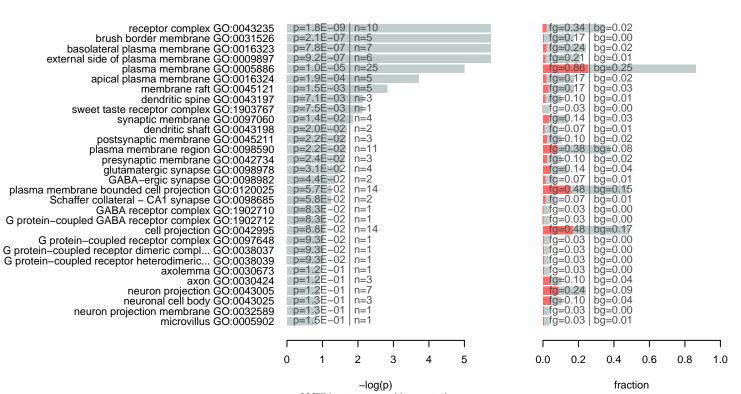


## fraction genes in fg and expected valu



## GO:CC gc020

## fraction genes in fg and expected value



n=29/77 input genes with annotations