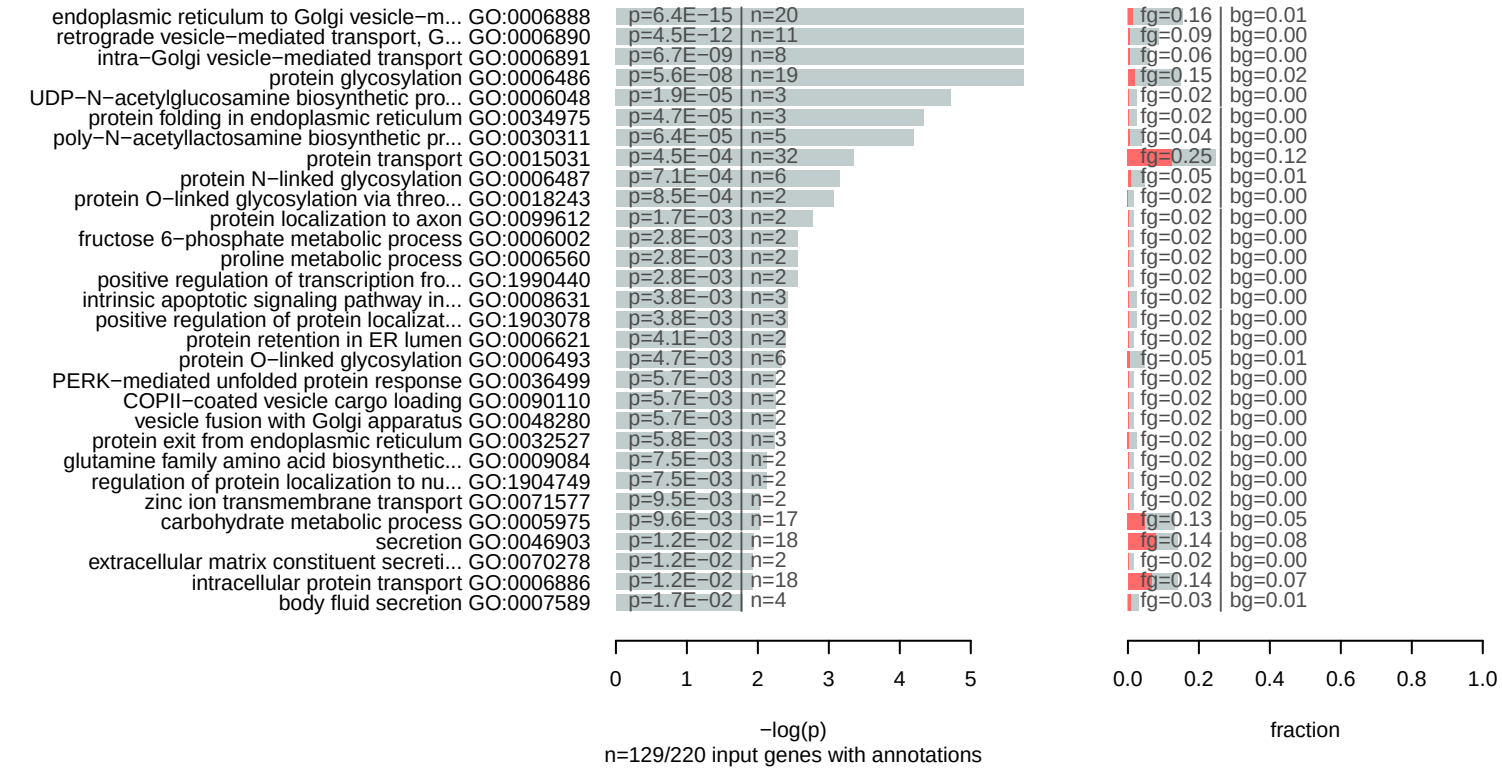
## GO:BP
## gc008

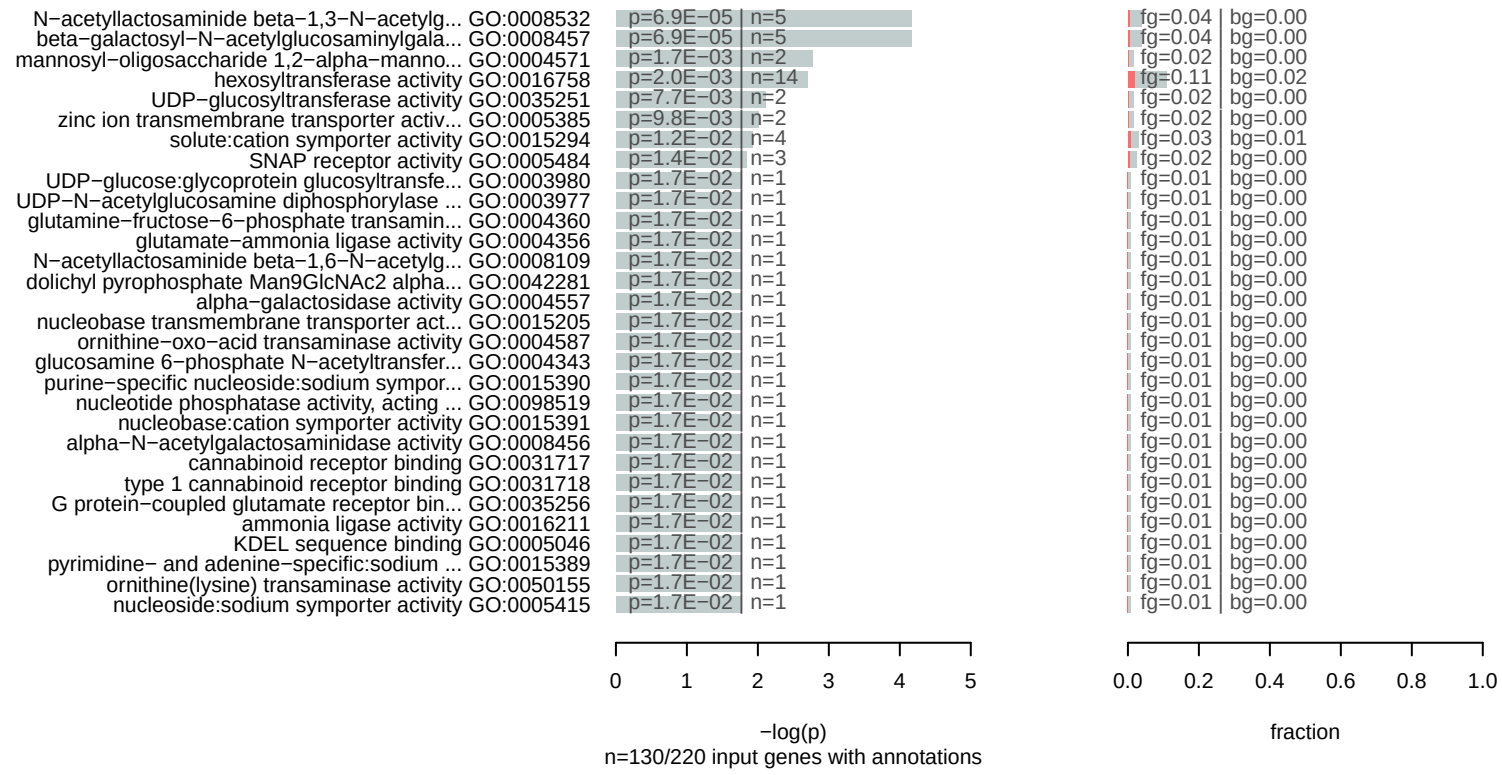
**fraction genes in fg and expected valu**




n=129/220 input genes with annotations

## GO:MF
## gc008

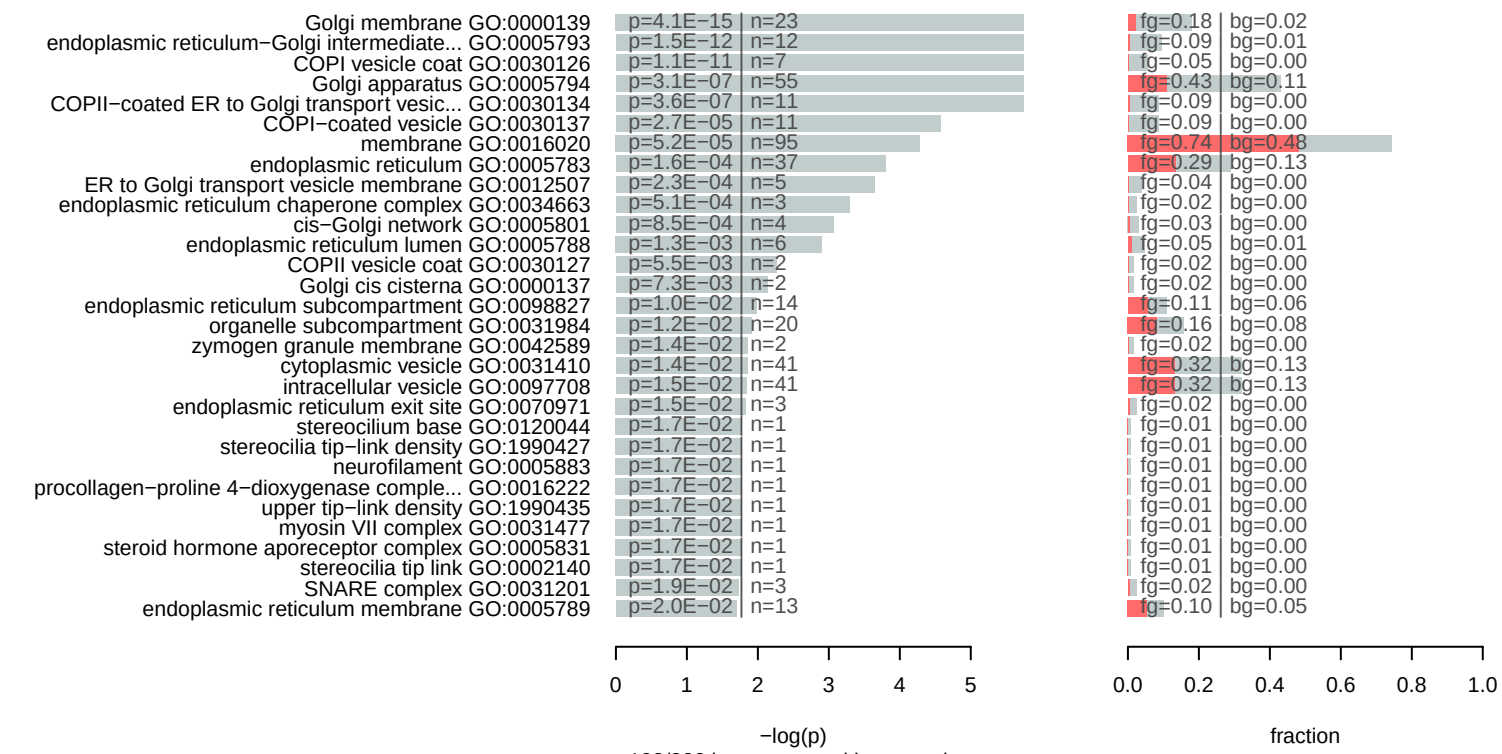
**fraction genes in fg and expected valu**




n=130/220 input genes with annotations

## GO:CC
## gc008

**fraction genes in fg and expected valu**




n=128/220 input genes with annotations