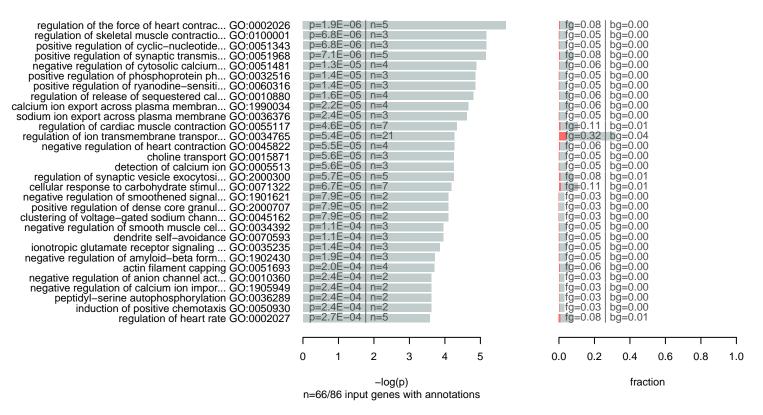
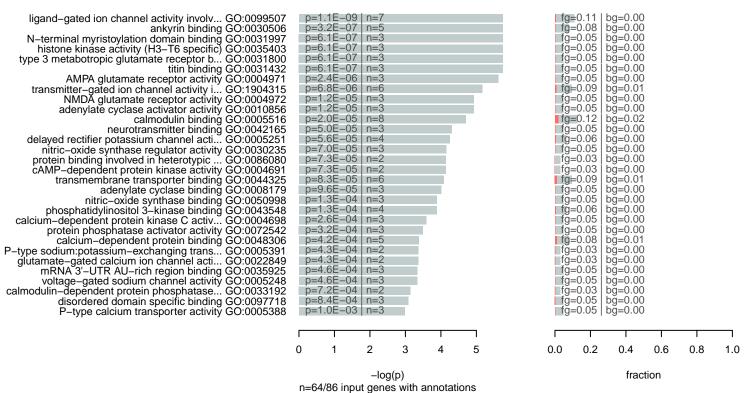
## GO:BP CniBil\_gain



#### GO:MF CniBil\_gain

### fraction genes in fg and expected valu



# GO:CC

CniBil\_gain

### fraction genes in fg and expected valu

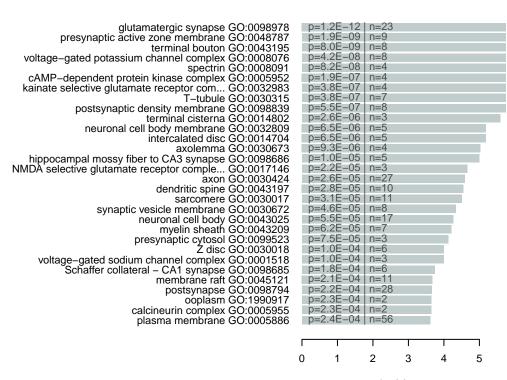
fg=0.35 | bg=0.06 fg=0.14 | bg=0.01

bg=0.01

bg=0.01

fg=0.12

fg=0.12



fg=0.06 bg=0.00 fg=0.06 bg=0.00 bg=0.00 fg=0.06 fg=0.11 bg=0.01 fg=0.12 fg=0.05 bg=0.01 bg=0.00 fg=0.08 bg=0.00 fg=0.08 fg=0.06 bg=0.00 bg=0.00 fg=0.08 | bg=0.00 | bg=0.00 fg=0.05 bg=0.06 fg=0.15 bg=0.02 fg=0.17 bg=0.02 fg=0.12 bg=0.01 fg=0.26 fg=0.11 bg=0.06 bg=0.02 fg=0.05 fg=0.09 bg=0.00 bg=0.01 fg=0.05 bg=0.00 bg=0.01 fg=0.09 fg=0.17 bg=0.03 bg=0.07 fq = 0.42fg=0.03 fg=0.03 bg=0.00 bg=0.00 bg=0.38 0.0 0.2 0.4 0.6 0.8 1.0 fraction

-log(p) n=66/86 input genes with annotations