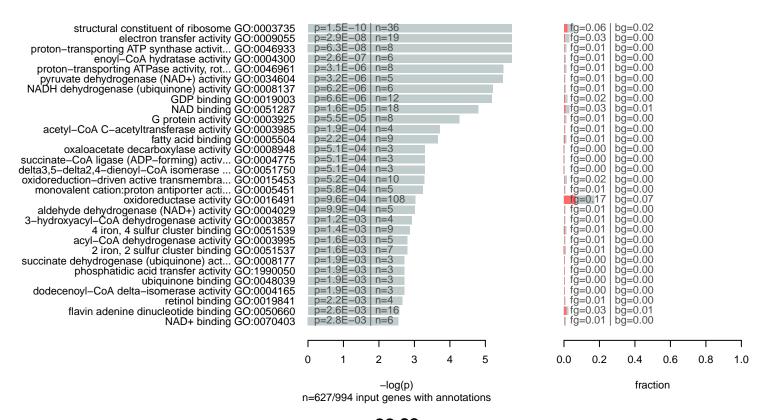


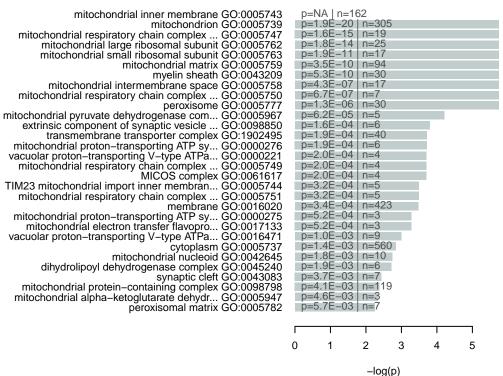
## GO:MF gc002a

## fraction genes in fg and expected value



## GO:CC gc002a

## fraction genes in fg and expected value



fg=0.25 | bg=0.05 fg=0.48 | bg=0.15 bg=0.00 bg=0.01 fg=0.03 fg=0.04 fg=0.03 bg=0.00 bg=0.03 fg=0.15 fg=0.05 fg=0.03 bg=0.01 bg=0.01 fg=0.01 fg=0.05 bg=0.00 bg=0.02 fg=0.01 bg=0.00 bg=0.00 bg=0.01 fa=0.01 fg=0.06 fg=0.01 fg=0.01 bg=0.00 bg=0.00 fg=0.01 bg=0.00 fg=0.01 bg=0.00  $f\bar{q} = 0.01$ bg=0.00 fg=0.01 bg=0.00 bg=0.00 fg=0.00 fg=0.00 fg=0.01 bg=0.00 bg=0.00 fg=0.02 bg=0.01 fg=0.01 bg=0.00 bg=0.00 bg=0.03 fg=0.01 fg=0.19 fg=0.00 bq = 0.00| fg=0.01 | bg=0.00 0.0 0.2 0.4 0.6 0.8 1.0

fraction

-log(p) n=640/994 input genes with annotations