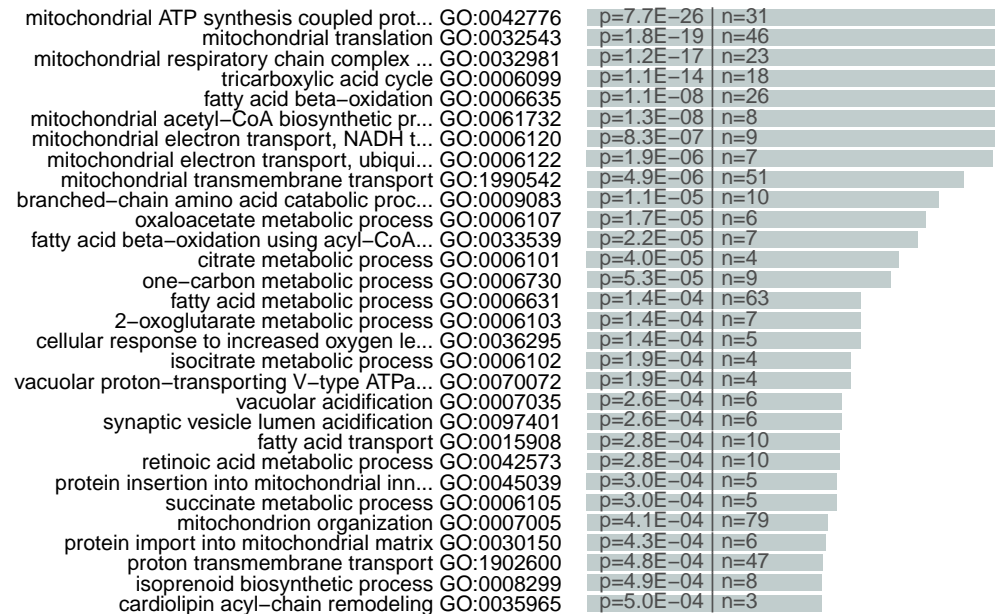
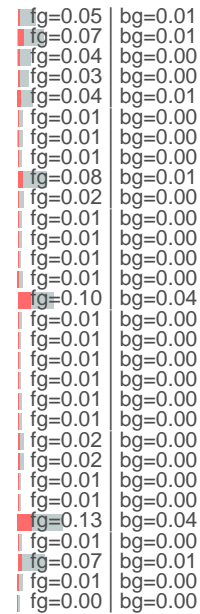


GO:BP  
gc002a

fraction genes in fg and expected value



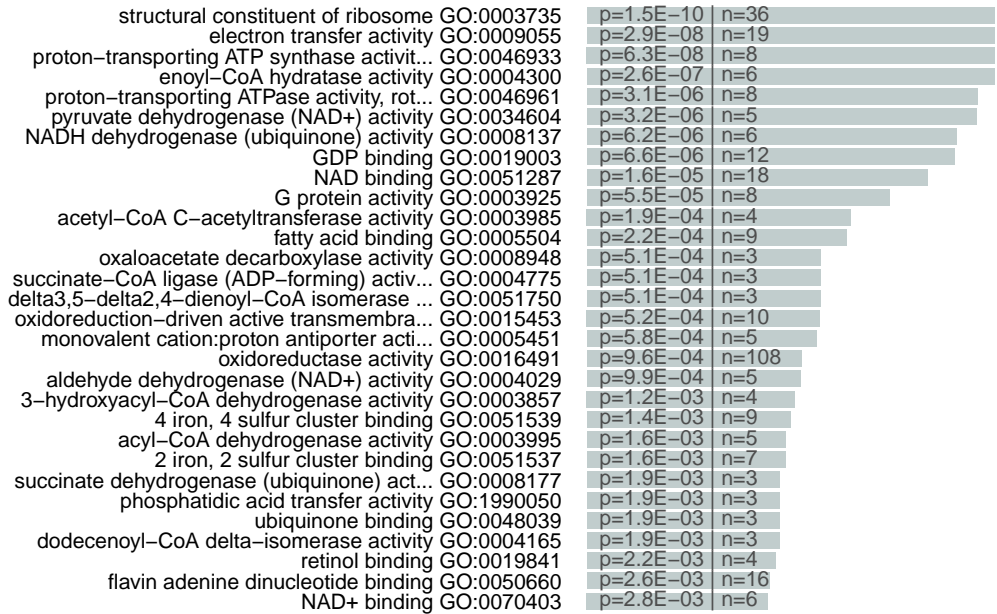
0 1 2 3 4 5  
-log(p)  
n=627/994 input genes with annotations



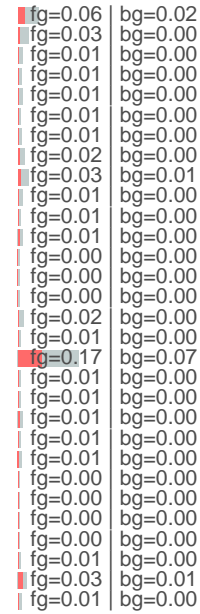
0.0 0.2 0.4 0.6 0.8 1.0  
fraction

GO:MF  
gc002a

fraction genes in fg and expected value



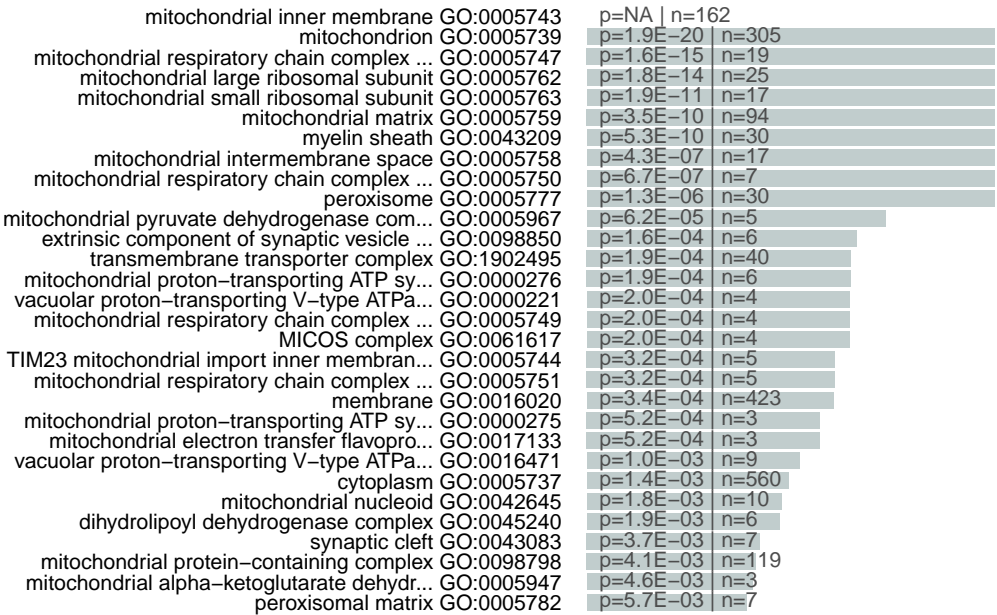
0 1 2 3 4 5  
-log(p)  
n=627/994 input genes with annotations



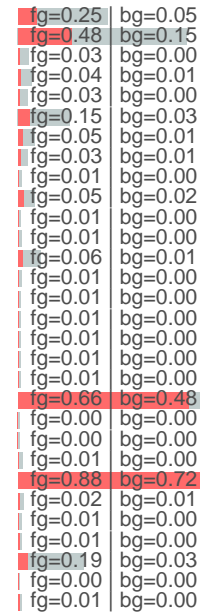
0.0 0.2 0.4 0.6 0.8 1.0  
fraction

GO:CC  
gc002a

fraction genes in fg and expected value



0 1 2 3 4 5  
-log(p)  
n=640/994 input genes with annotations



0.0 0.2 0.4 0.6 0.8 1.0  
fraction