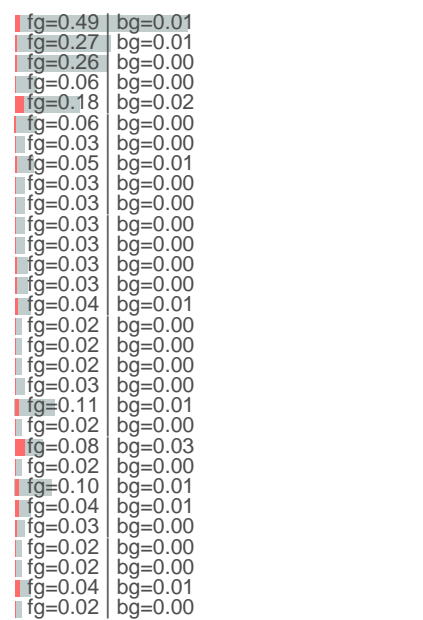
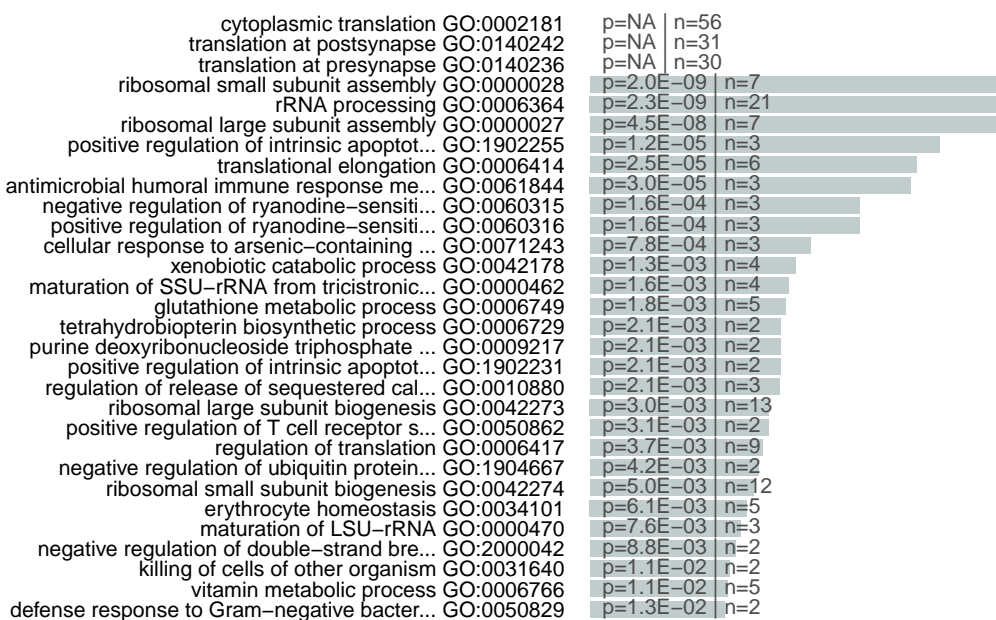


**GO:BP**  
**gc007a**

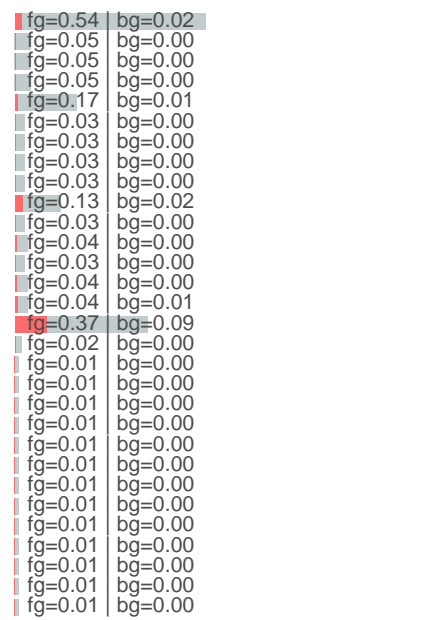
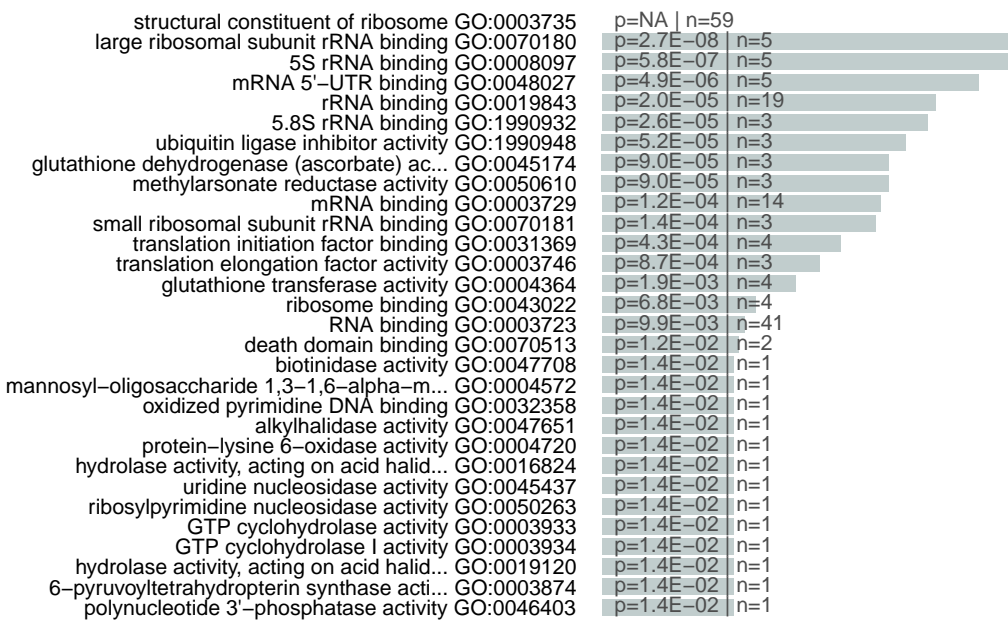
fraction genes in fg and expected value



0 1 2 3 4 5  
-log(p)  
n=115/159 input genes with annotations

**GO:MF**  
**gc007a**

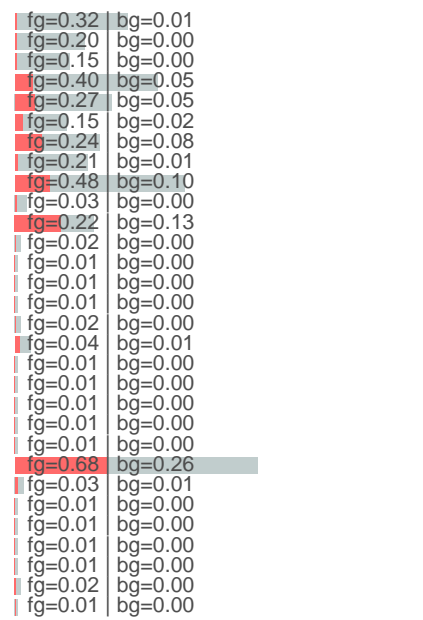
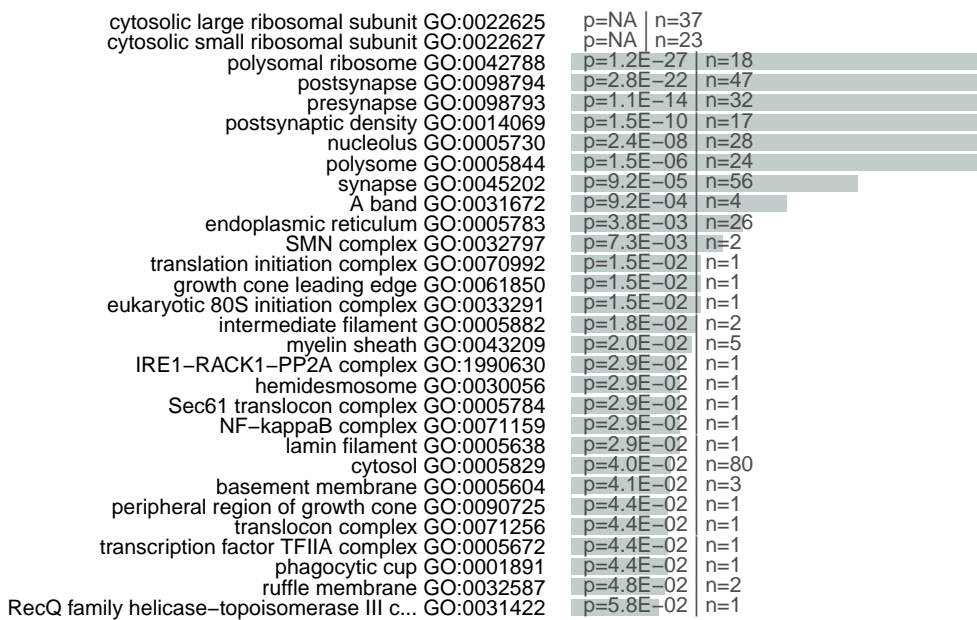
fraction genes in fg and expected value



0 1 2 3 4 5  
-log(p)  
n=110/159 input genes with annotations

**GO:CC**  
**gc007a**

fraction genes in fg and expected value



0 1 2 3 4 5  
-log(p)  
n=117/159 input genes with annotations