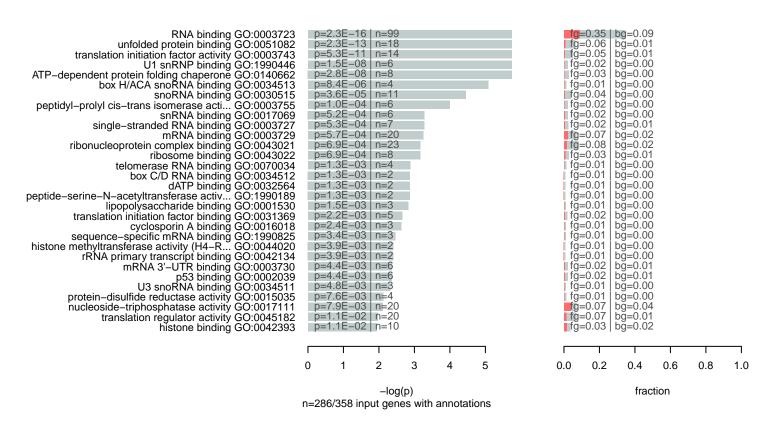


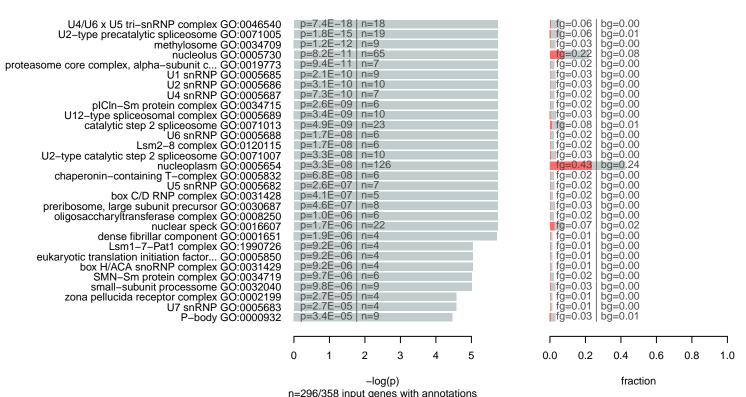
## GO:MF gc007b

## fraction genes in fg and expected value



## GO:CC gc007b

## fraction genes in fg and expected value



n=296/358 input genes with annotations