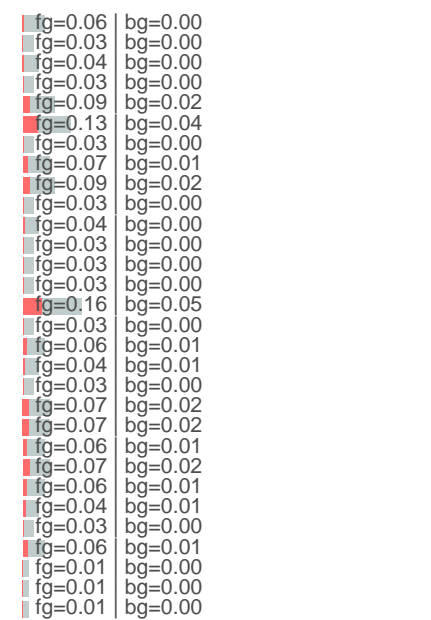
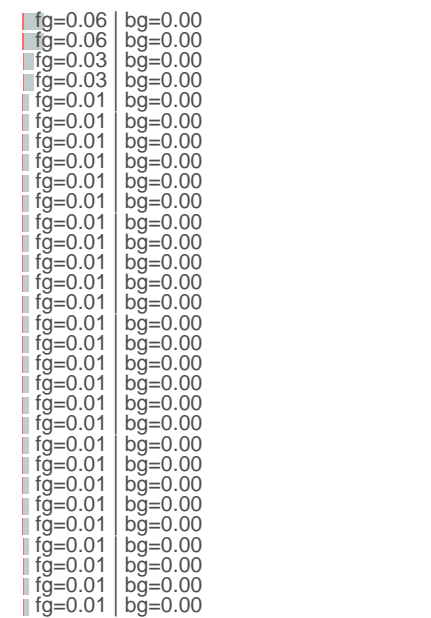


## fraction genes in fg and expected value



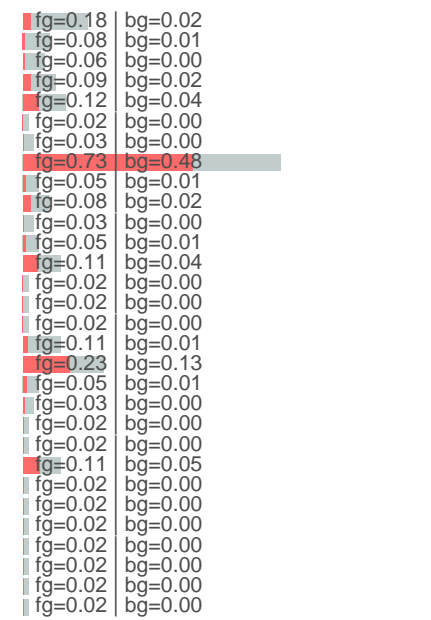
fraction

## fraction genes in fg and expected value



fraction

## fraction genes in fg and expected value



0 1 2 3 4 5

$-\log(p)$

n=66/134 input genes with annotations

fraction