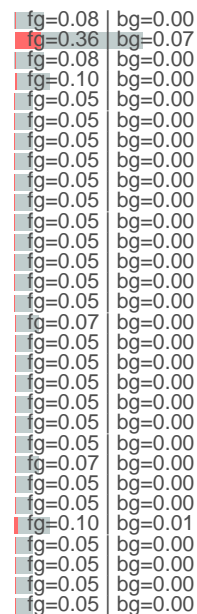
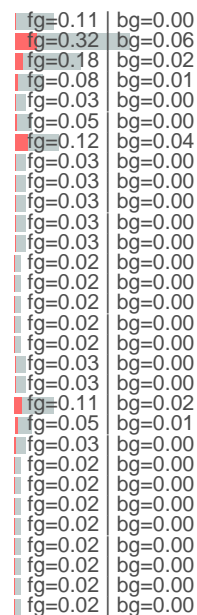


fraction genes in fg and expected value

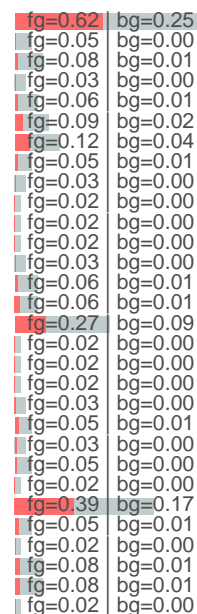


A horizontal axis labeled "fraction" with tick marks at 0.0, 0.2, 0.4, 0.6, 0.8, and 1.0.

fraction genes in fg and expected value



fraction genes in fg and expected value



0 1 2 3 4 5

$-\log(p)$

n=64/195 input genes with annotations