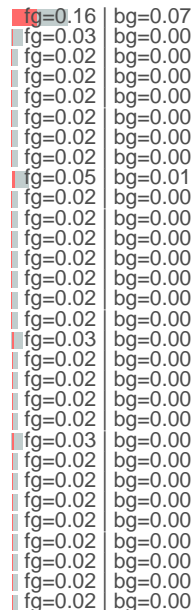


fraction genes in fg and bg



fraction

n=49/53 input genes with annotations