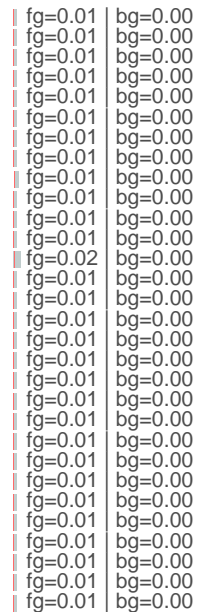


fraction genes in fg and bg



fraction

n=132/92 input genes with annotations