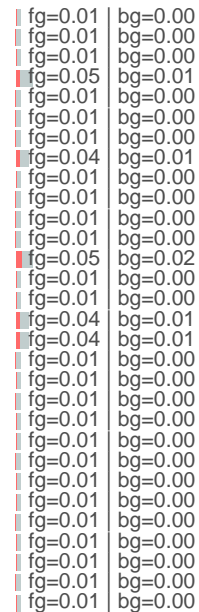


fraction genes in fg and bg



fraction

n=70/71 input genes with annotations