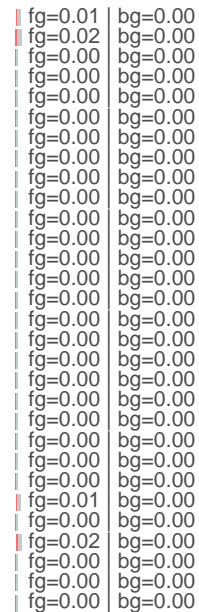


fraction genes in fg and bg



fraction

n=193/180 input genes with annotations