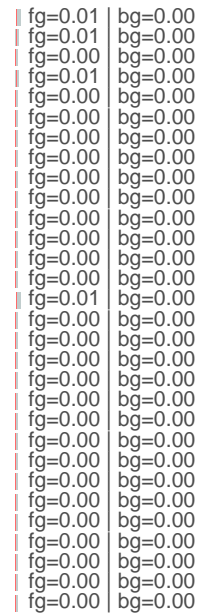


fraction genes in fg and bg



fraction

n=178/167 input genes with annotations