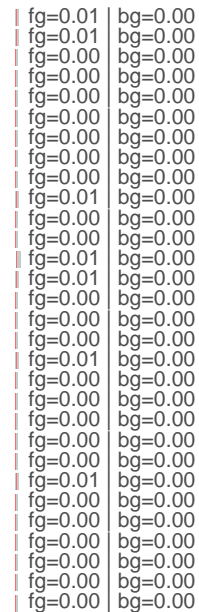


fraction genes in fg and bg



fraction

n=190/161 input genes with annotations