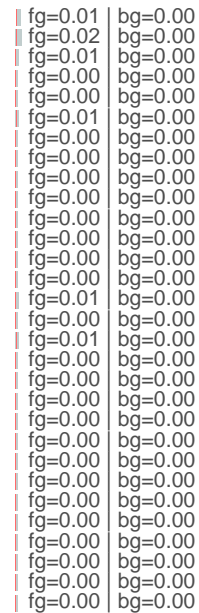


fraction genes in fg and bg



fraction

n=189/169 input genes with annotations