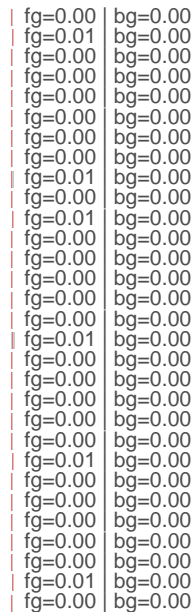


fraction genes in fg and bg



fraction

n=271/248 input genes with annotations