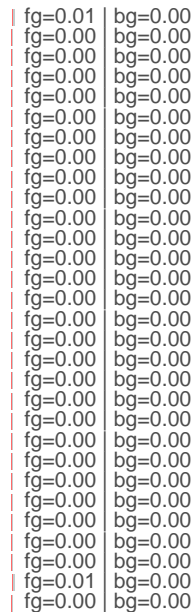


fraction genes in fg and bg



fraction

n=247/211 input genes with annotations