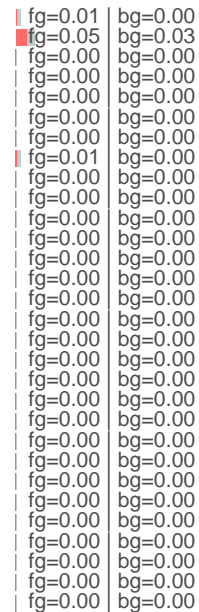


fraction genes in fg and bg



fraction

n=247/227 input genes with annotations