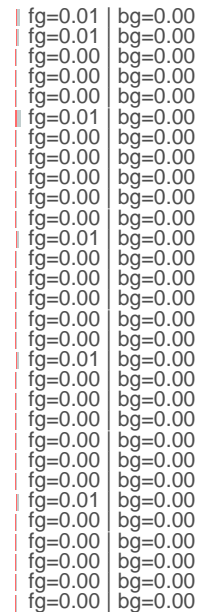


fraction genes in fg and bg



fraction

n=250/243 input genes with annotations