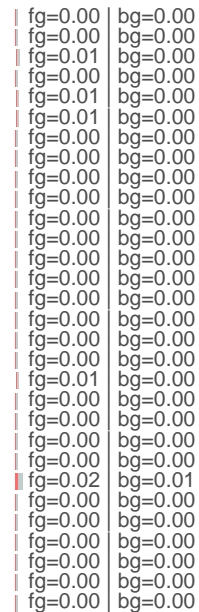


fraction genes in fg and bg



fraction

n=240/223 input genes with annotations