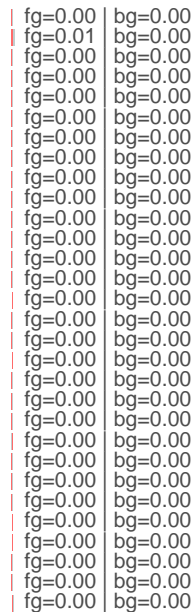


fraction genes in fg and bg



fraction

n=833/983 input genes with annotations