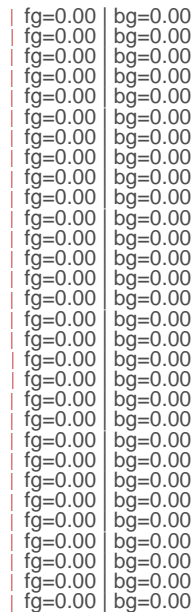


fraction genes in fg and bg



fraction

n=432/450 input genes with annotations