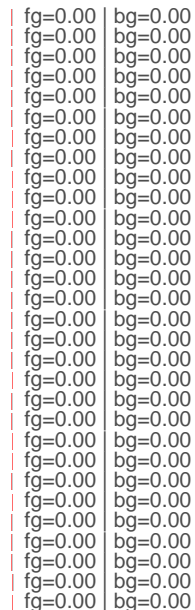


fraction genes in fg and bg



fraction

n=966/1041 input genes with annotations