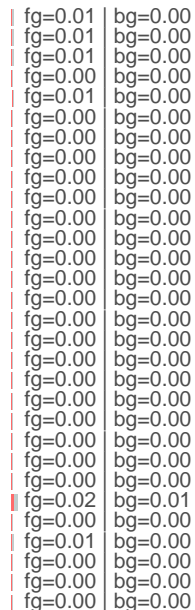


fraction genes in fg and bg



fraction

n=293/285 input genes with annotations