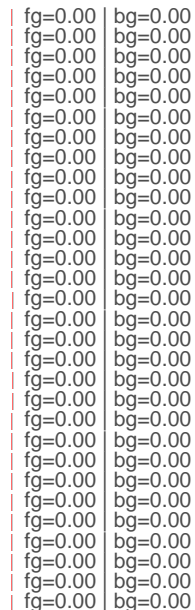


fraction genes in fg and bg



fraction

n=377/324 input genes with annotations