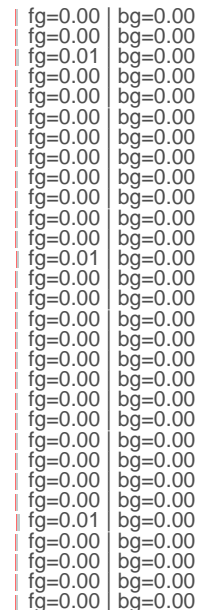


fraction genes in fg and bg



fraction

n=245/207 input genes with annotations