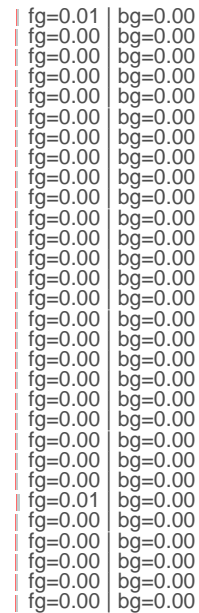


fraction genes in fg and bg



fraction

n=225/187 input genes with annotations