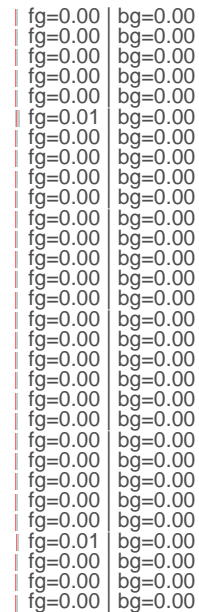


fraction genes in fg and bg



fraction

n=223/189 input genes with annotations