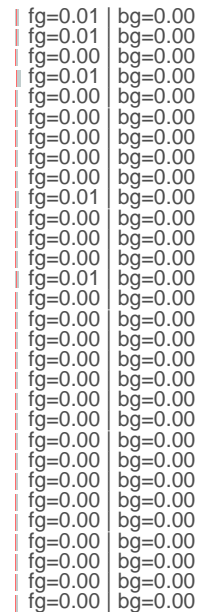


fraction genes in fg and bg



fraction

n=200/159 input genes with annotations