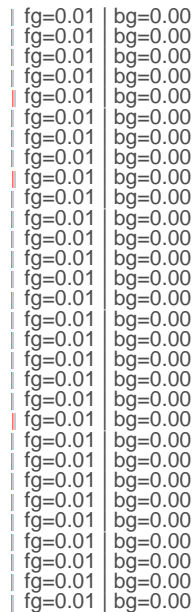


fraction genes in fg and bg



fraction

n=165/134 input genes with annotations