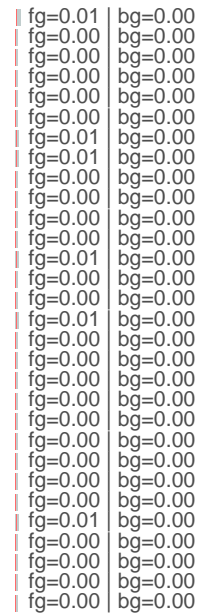


fraction genes in fg and bg



fraction

n=191/152 input genes with annotations