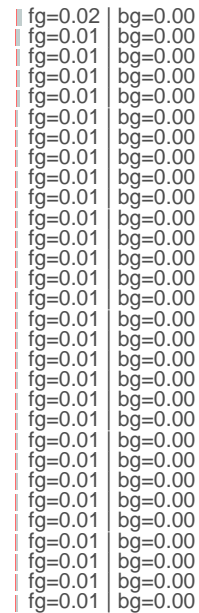


fraction genes in fg and bg



fraction

n=162/134 input genes with annotations