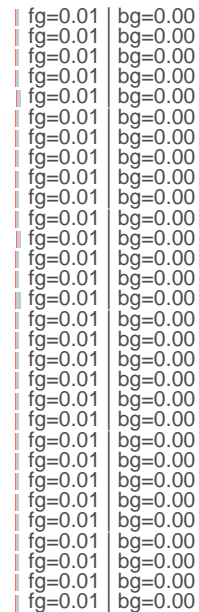


fraction genes in fg and bg



fraction

n=139/118 input genes with annotations