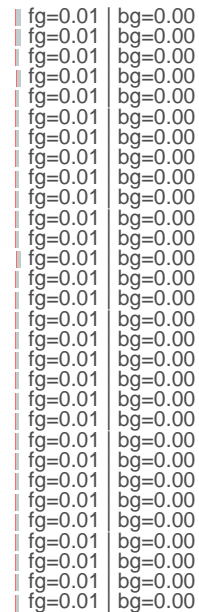


fraction genes in fg and bg



fraction

n=132/97 input genes with annotations