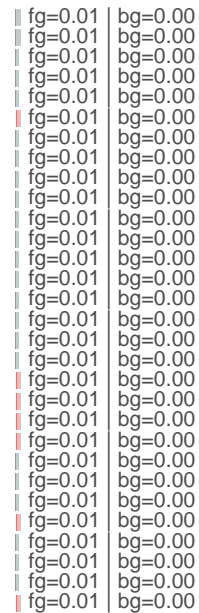


fraction genes in fg and bg



fraction

n=123/110 input genes with annotations