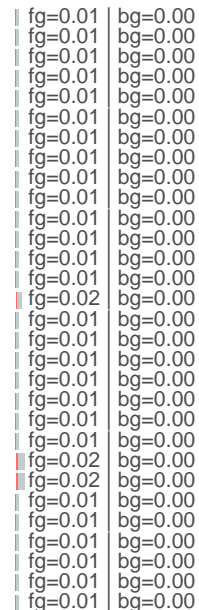


fraction genes in fg and bg



fraction

n=110/110 input genes with annotations