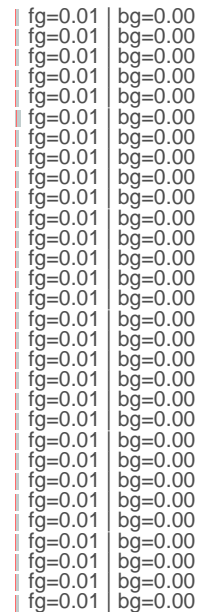


fraction genes in fg and bg



fraction

n=122/112 input genes with annotations