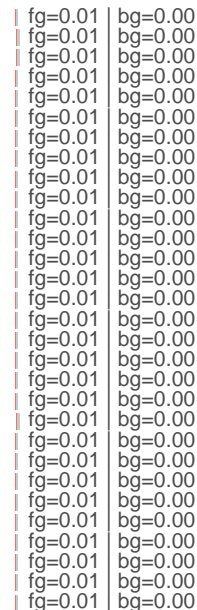


fraction genes in fg and bg



fraction

n=171/121 input genes with annotations