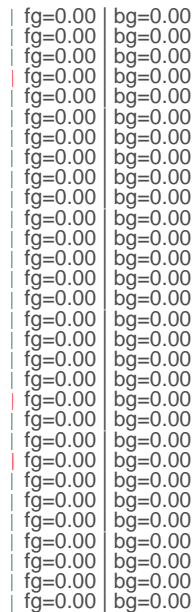


fraction genes in fg and bg



fraction

n=1193/1446 input genes with annotations