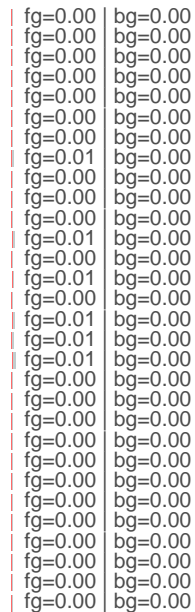


fraction genes in fg and bg



fraction

n=269/246 input genes with annotations