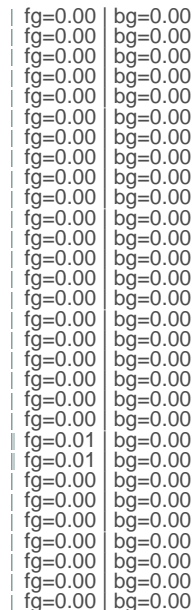


fraction genes in fg and bg



fraction

n=253/212 input genes with annotations