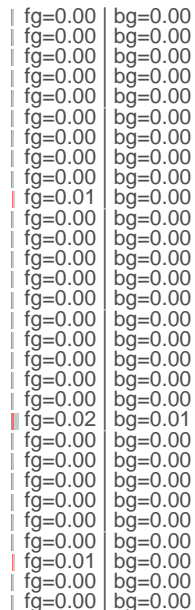


fraction genes in fg and bg



fraction

n=215/213 input genes with annotations