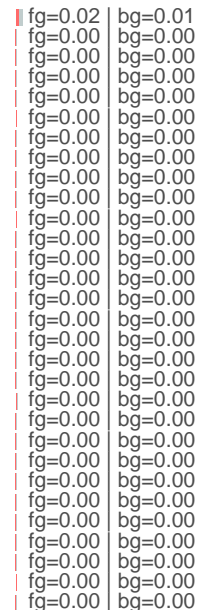


fraction genes in fg and bg



fraction

n=795/797 input genes with annotations