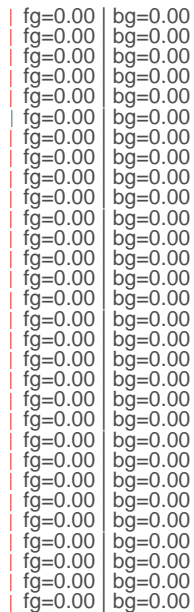


fraction genes in fg and bg



fraction

n=795/805 input genes with annotations