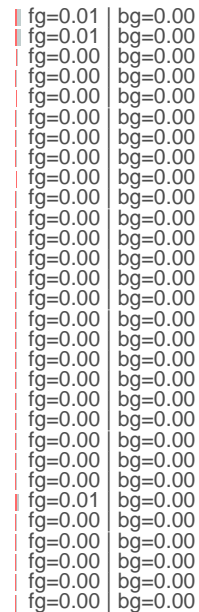


fraction genes in fg and bg



fraction

n=780/802 input genes with annotations