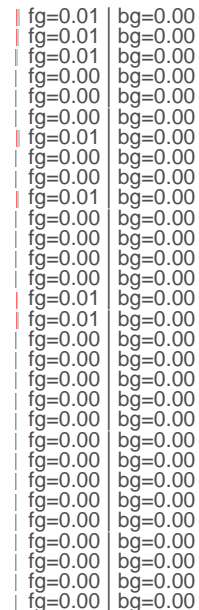


fraction genes in fg and bg



fraction

n=279/293 input genes with annotations