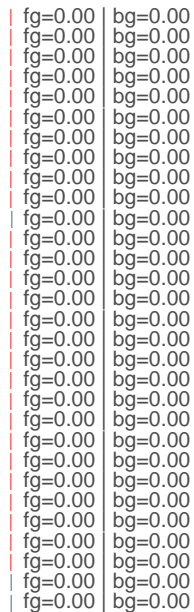


fraction genes in fg and bg



fraction

n=757/771 input genes with annotations