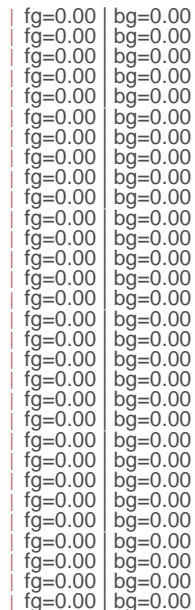


fraction genes in fg and bg



fraction

n=296/279 input genes with annotations