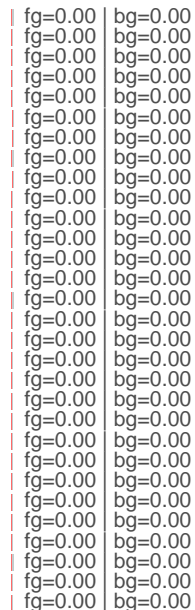


fraction genes in fg and bg



fraction

n=343/279 input genes with annotations