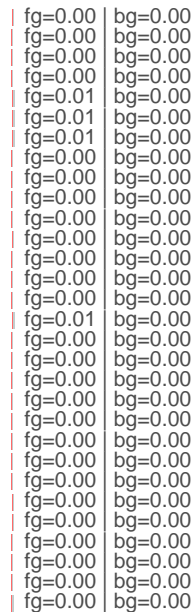


fraction genes in fg and bg



fraction

n=342/347 input genes with annotations