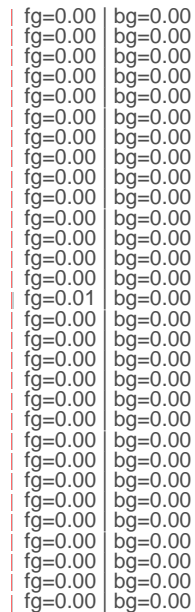


fraction genes in fg and bg



fraction

n=292/274 input genes with annotations