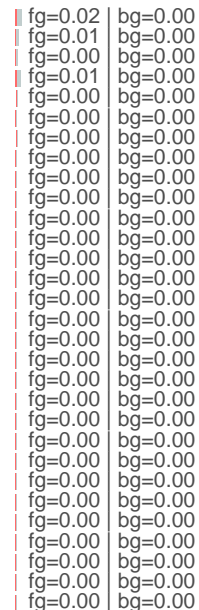


fraction genes in fg and bg



fraction

n=299/309 input genes with annotations