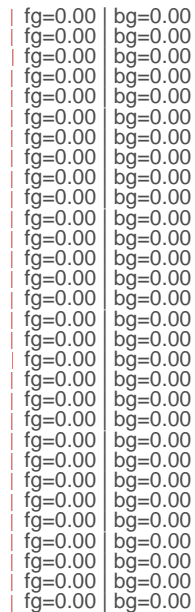


fraction genes in fg and bg



fraction

n=935/982 input genes with annotations