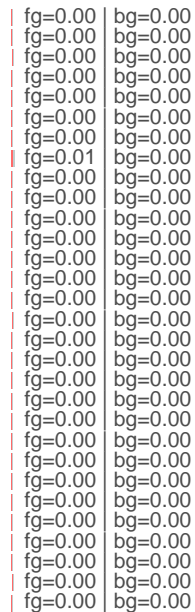


fraction genes in fg and bg



fraction

n=578/588 input genes with annotations