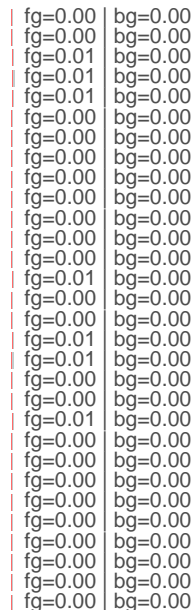


fraction genes in fg and bg



fraction

n=295/260 input genes with annotations