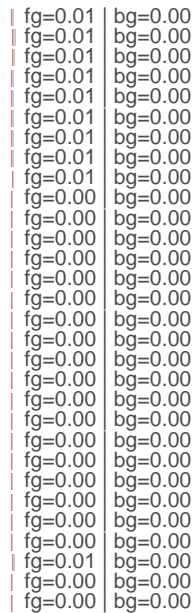


fraction genes in fg and bg



fraction

n=314/282 input genes with annotations