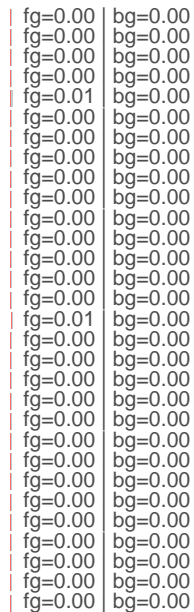


fraction genes in fg and bg



fraction

n=319/263 input genes with annotations