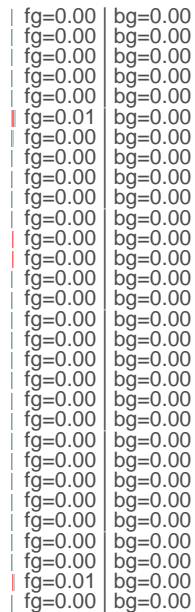


fraction genes in fg and bg



fraction

n=347/328 input genes with annotations