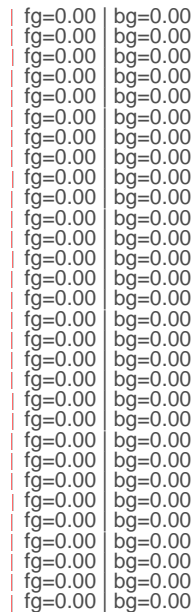


fraction genes in fg and bg



fraction

n=608/584 input genes with annotations