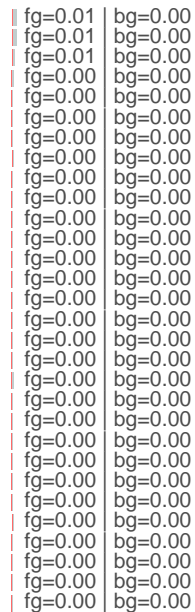


fraction genes in fg and bg



fraction

n=317/273 input genes with annotations