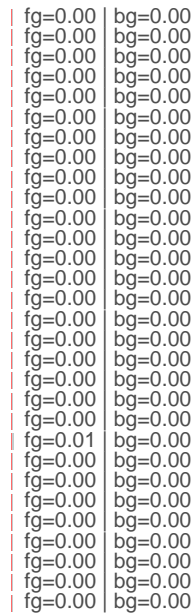


fraction genes in fg and bg



fraction

n=261/238 input genes with annotations