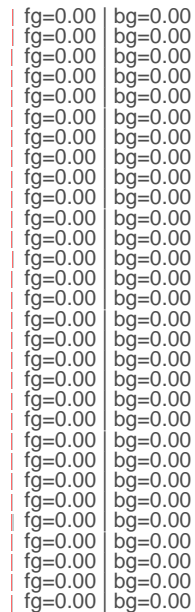


fraction genes in fg and bg



fraction

n=366/341 input genes with annotations