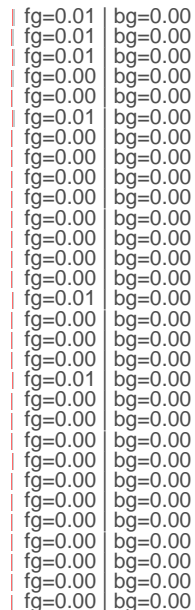


fraction genes in fg and bg



fraction

n=442/400 input genes with annotations