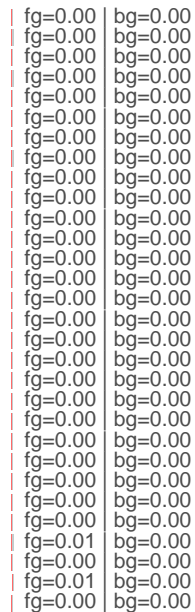


fraction genes in fg and bg



fraction

n=424/375 input genes with annotations