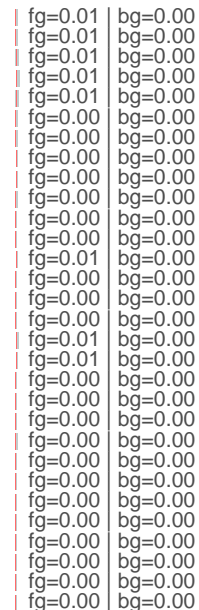


fraction genes in fg and bg



fraction

n=417/445 input genes with annotations