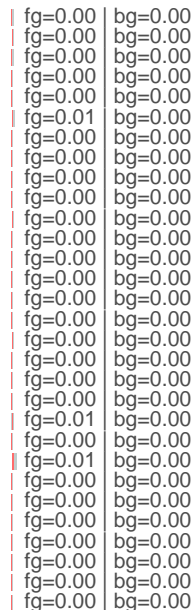


fraction genes in fg and bg



fraction

n=392/379 input genes with annotations