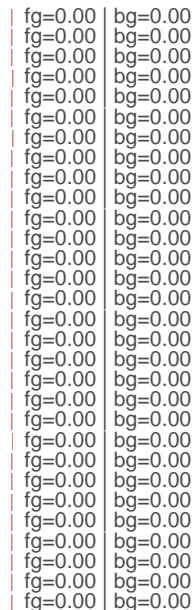


fraction genes in fg and bg



fraction

n=601/612 input genes with annotations