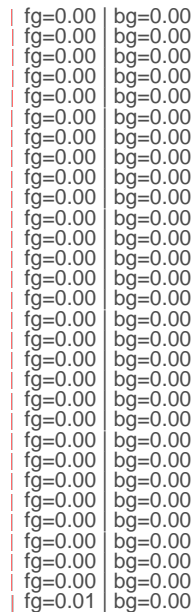


fraction genes in fg and bg



fraction

n=502/482 input genes with annotations