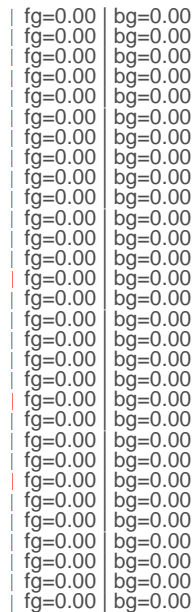


fraction genes in fg and bg



fraction

n=552/504 input genes with annotations