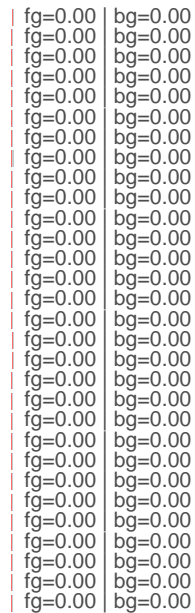


fraction genes in fg and bg



fraction

n=385/350 input genes with annotations