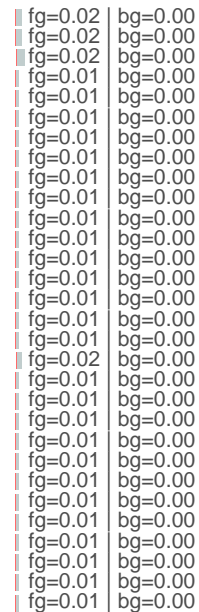


fraction genes in fg and bg



fraction

n=117/105 input genes with annotations