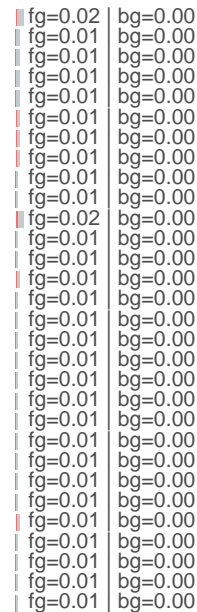


fraction genes in fg and bg



fraction

n=150/155 input genes with annotations