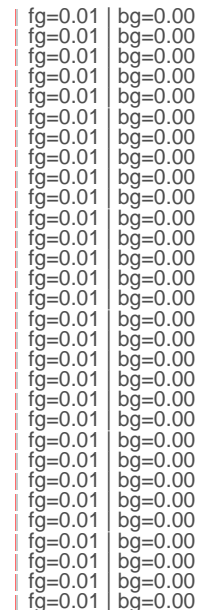


fraction genes in fg and bg



fraction

n=154/143 input genes with annotations