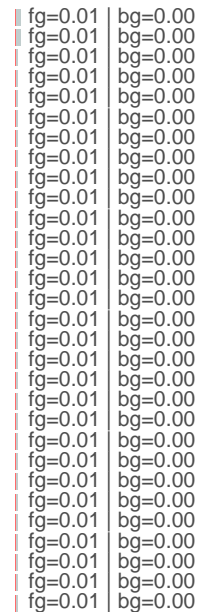


fraction genes in fg and bg



fraction

n=144/118 input genes with annotations