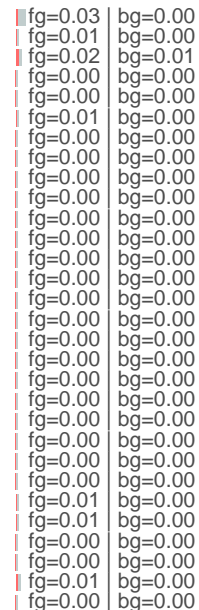


fraction genes in fg and bg



fraction

n=183/187 input genes with annotations