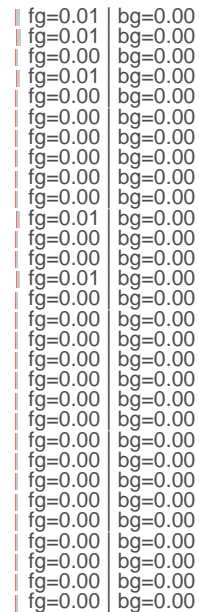


fraction genes in fg and bg



fraction

n=182/157 input genes with annotations