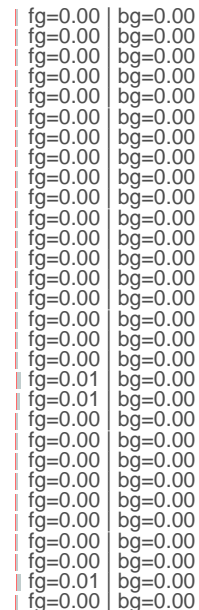


fraction genes in fg and bg



fraction

n=182/153 input genes with annotations