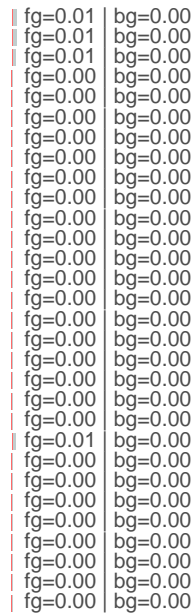


fraction genes in fg and bg



fraction

n=254/237 input genes with annotations