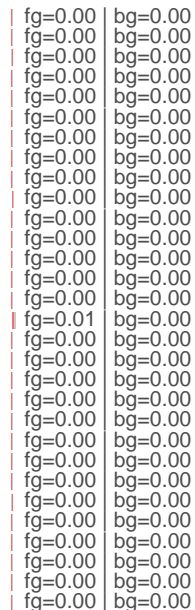


fraction genes in fg and bg



fraction

n=454/435 input genes with annotations