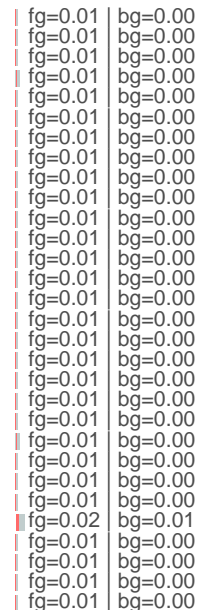


fraction genes in fg and bg



fraction

n=130/130 input genes with annotations