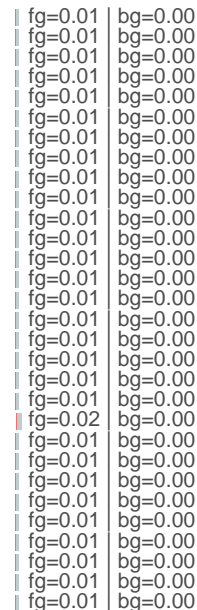


fraction genes in fg and bg



fraction

n=112/87 input genes with annotations