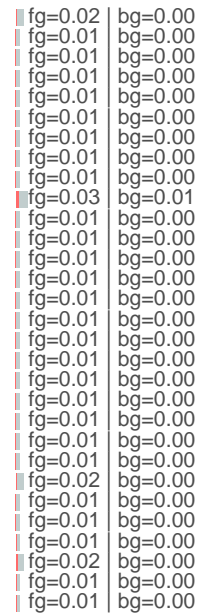


fraction genes in fg and bg



fraction

n=79/87 input genes with annotations