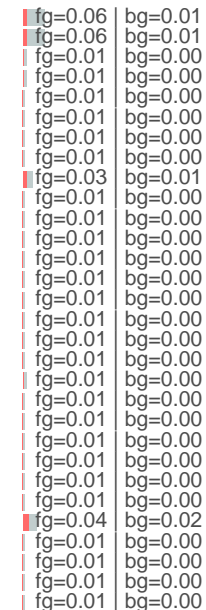


**fraction genes in fg and bg**



fraction

n=126/127 input genes with annotations