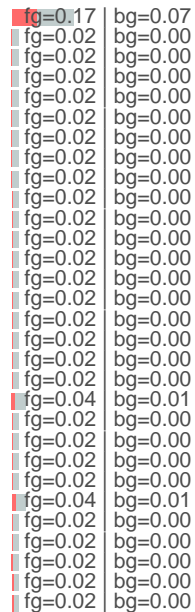


**fraction genes in fg and bg**



fraction

n=42/55 input genes with annotations