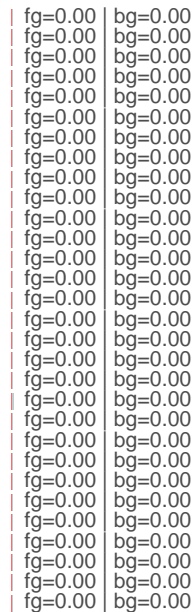


**fraction genes in fg and bg**



fraction

n=453/381 input genes with annotations