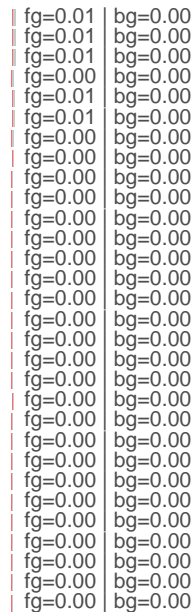


**fraction genes in fg and bg**



fraction

n=292/292 input genes with annotations