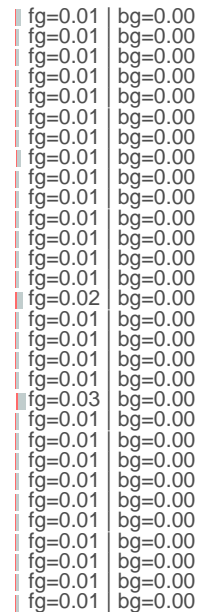


**fraction genes in fg and bg**



fraction

n=137/109 input genes with annotations