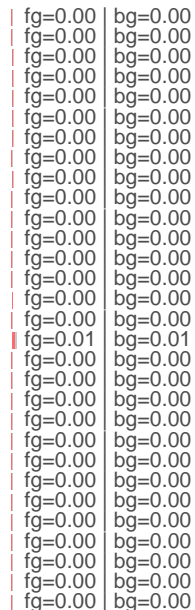


**fraction genes in fg and bg**



fraction

n=761/765 input genes with annotations