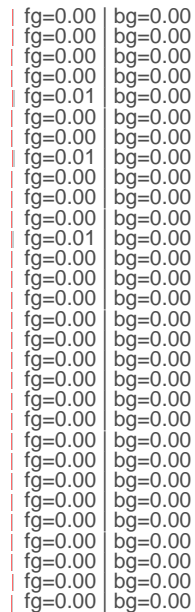


**fraction genes in fg and bg**



fraction

n=425/426 input genes with annotations