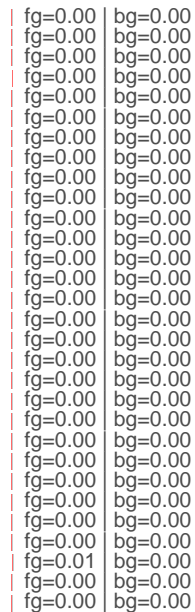


**fraction genes in fg and bg**



fraction

n=682/677 input genes with annotations