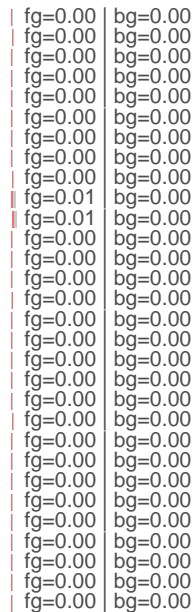


**fraction genes in fg and bg**



fraction

n=448/514 input genes with annotations