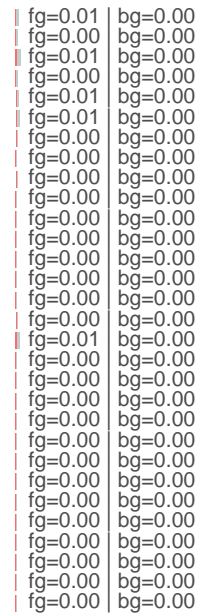


**fraction genes in fg and bg**



fraction

n=425/436 input genes with annotations