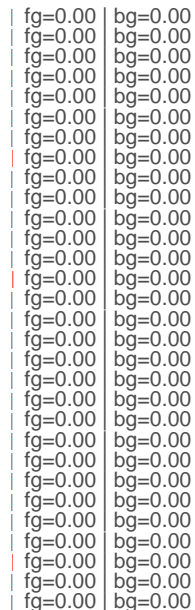


fraction genes in fg and bg



fraction

n=617/599 input genes with annotations