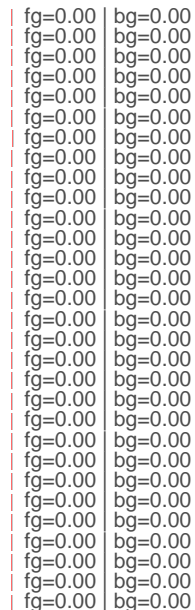


fraction genes in fg and bg



fraction

n=615/593 input genes with annotations