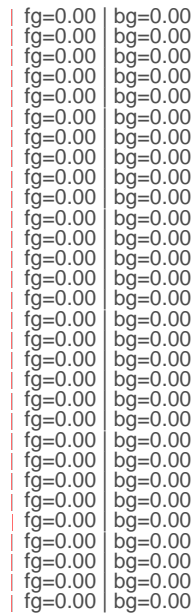


fraction genes in fg and bg



fraction

n=568/609 input genes with annotations