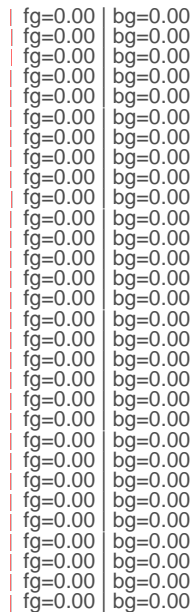


fraction genes in fg and bg



fraction

n=703/667 input genes with annotations