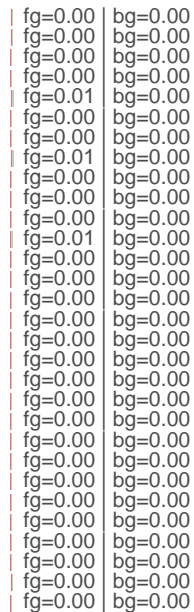


fraction genes in fg and bg



fraction

n=425/426 input genes with annotations