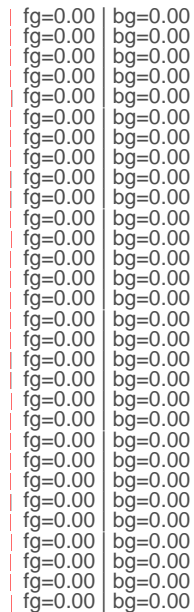


fraction genes in fg and bg



fraction

n=1823/2447 input genes with annotations