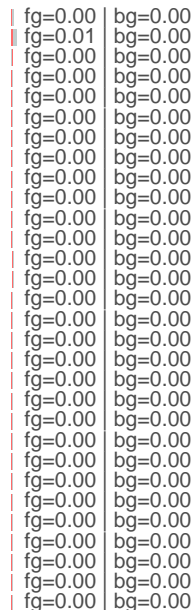


fraction genes in fg and bg



fraction

n=482/440 input genes with annotations