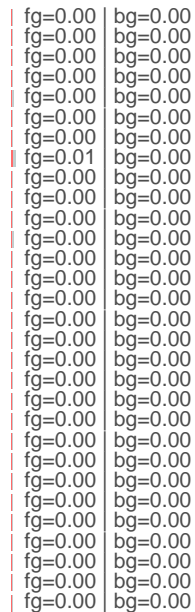


fraction genes in fg and bg



fraction

n=371/355 input genes with annotations