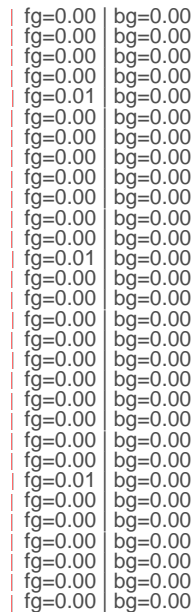


fraction genes in fg and bg



fraction

n=424/385 input genes with annotations