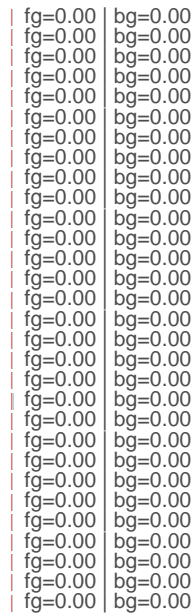


fraction genes in fg and bg



fraction

n=453/381 input genes with annotations