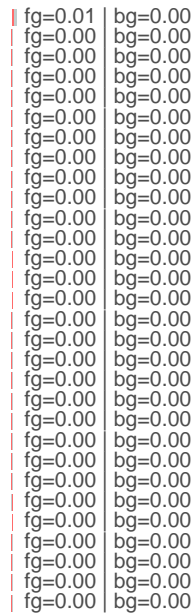


fraction genes in fg and bg



fraction

n=811/1029 input genes with annotations