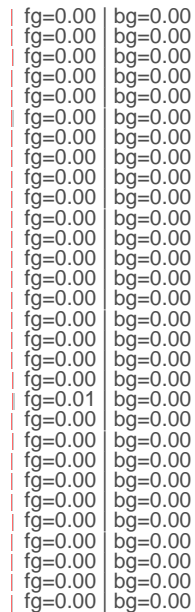


fraction genes in fg and bg



fraction

n=376/337 input genes with annotations