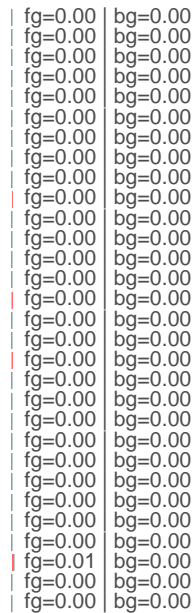


fraction genes in fg and bg



fraction

n=655/627 input genes with annotations