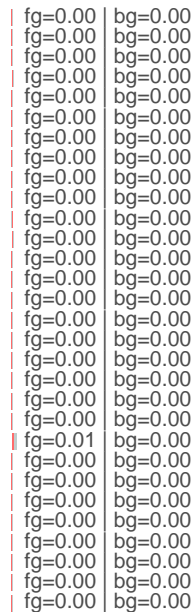


fraction genes in fg and bg



fraction

n=639/699 input genes with annotations