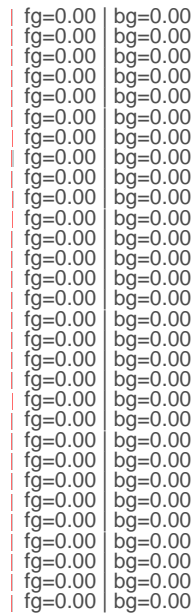


fraction genes in fg and bg



fraction

n=581/549 input genes with annotations