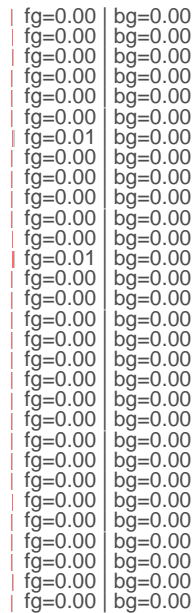


fraction genes in fg and bg



fraction

n=523/522 input genes with annotations