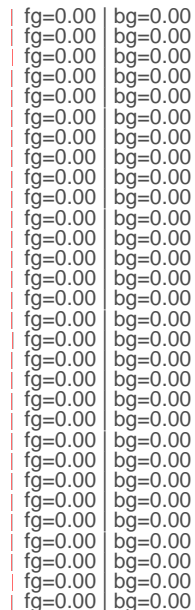


fraction genes in fg and bg



fraction

n=546/515 input genes with annotations