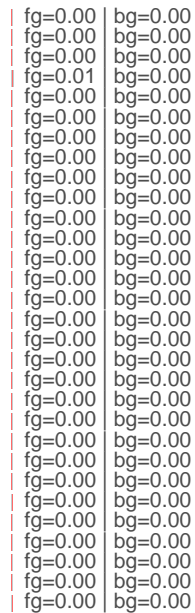


fraction genes in fg and bg



fraction

n=514/477 input genes with annotations