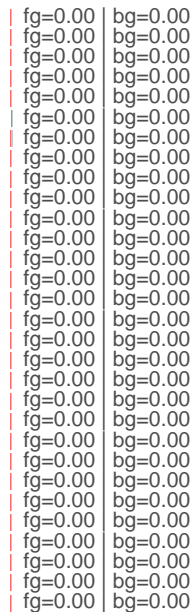


fraction genes in fg and bg



fraction

n=588/571 input genes with annotations