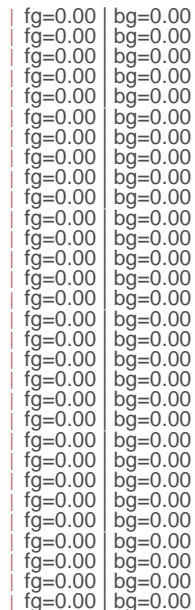


fraction genes in fg and bg



fraction

n=588/581 input genes with annotations