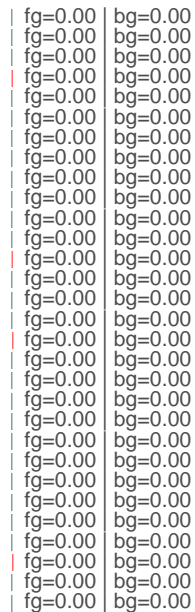


fraction genes in fg and bg



fraction

n=732/736 input genes with annotations