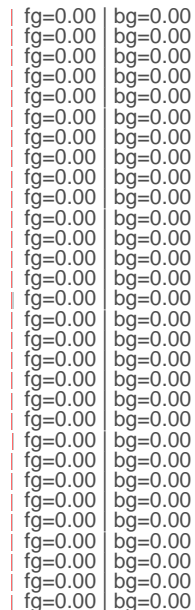


fraction genes in fg and bg



fraction

n=335/266 input genes with annotations