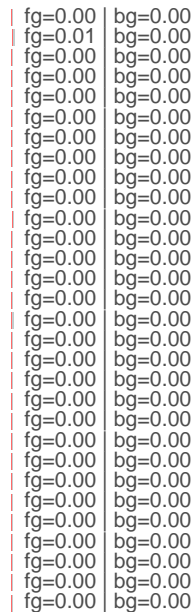


fraction genes in fg and bg



fraction

n=359/316 input genes with annotations