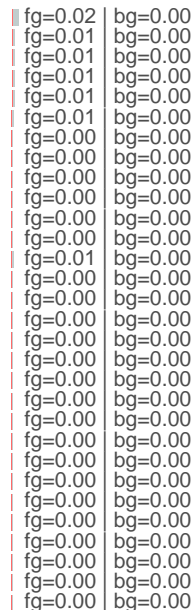


fraction genes in fg and bg



fraction

n=306/262 input genes with annotations