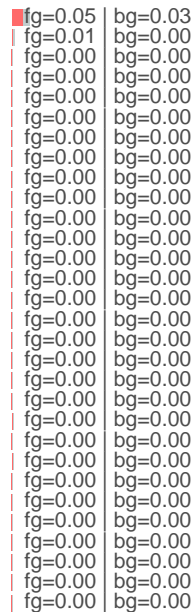


fraction genes in fg and bg



fraction

n=297/296 input genes with annotations