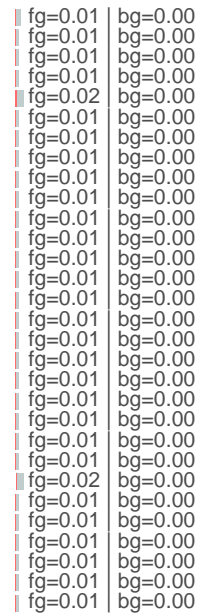


fraction genes in fg and bg



fraction

n=126/106 input genes with annotations