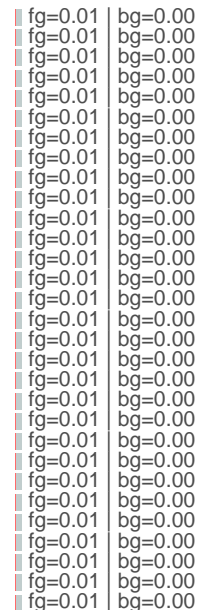


fraction genes in fg and bg



fraction

n=66/42 input genes with annotations