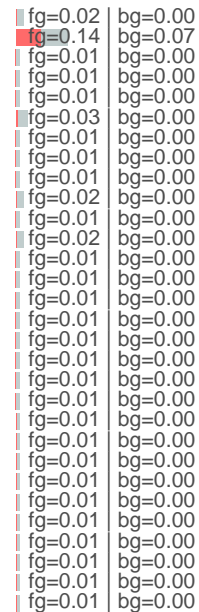


fraction genes in fg and bg



fraction

n=66/70 input genes with annotations