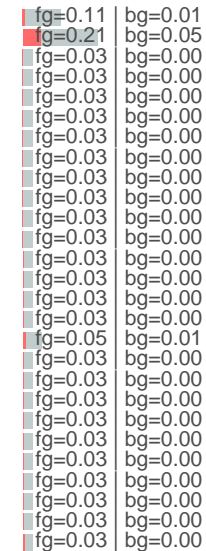


fraction genes in fg and bg



fraction

n=27/45 input genes with annotations