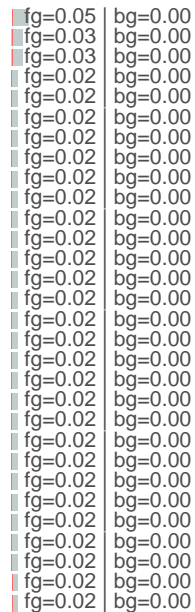


fraction genes in fg and bg



fraction

n=58/49 input genes with annotations