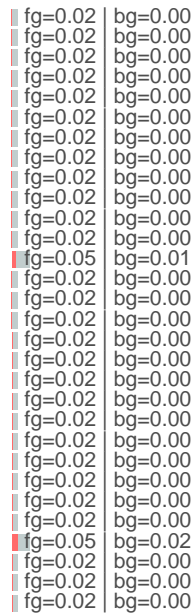


fraction genes in fg and bg



fraction

n=51/49 input genes with annotations