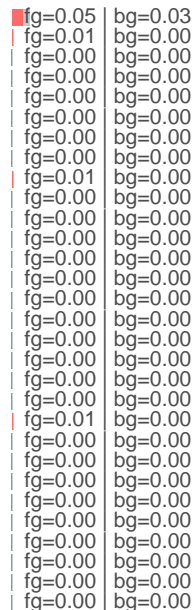


fraction genes in fg and bg



fraction

n=288/267 input genes with annotations