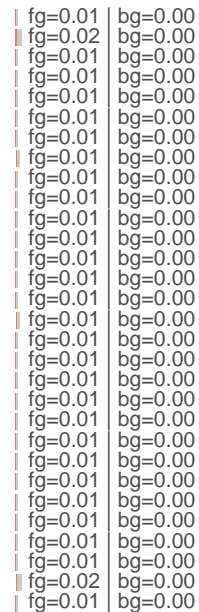


fraction genes in fg and bg



fraction

n=160/125 input genes with annotations