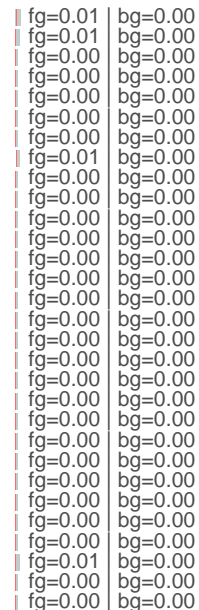


fraction genes in fg and bg



fraction

n=180/136 input genes with annotations