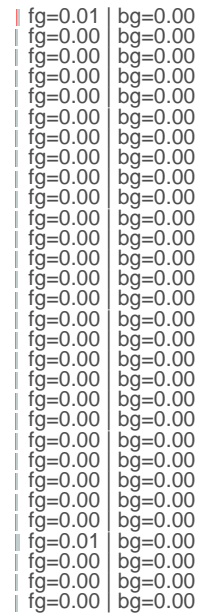


fraction genes in fg and bg



fraction

n=185/155 input genes with annotations