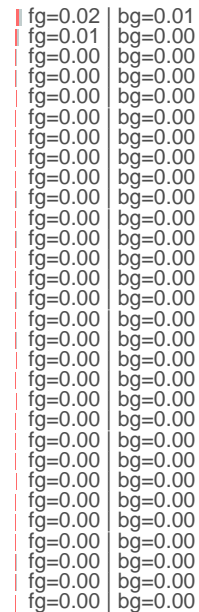


fraction genes in fg and bg



fraction

n=1576/1770 input genes with annotations