## ENSG00000102760\_RGCC

0.2 0 0.3

total(k)

mTconv mTreg nTconv nTreg (1.13e-05)(9.70e-01)(5.32e-11)(3.77e - 03)1.49 1.65 , 000 fold change 1.22 0.61 0.82 0.61 0.82 0.67 0.37 0.37 p-value 0 0.01 mTconv nTconv mTreg nTreg fold change 1e-04 1.08 0.95 1e-06 0.9 0.96 1e-08 ADD 4DE ADD ADE 1e-10

0.4

0.5

disease

HA

MS