## ENSG00000165272\_AQP3

total(k)

disease

HA

mTconv mTreg nTconv nTreg (1.98e - 09)(1.27e-02)(4.96e-01)(7.62e-01)1.22 1.11 φ¢ fold change 1.11 404 0.82 0.82 0.9 0.9 0.67 0.67 0.82 p-value mTconv nTconv mTreg nTreg fold change 0.01 1.05 1.06 1.03 1.04 1.02 1.05 1e-04 0.98 0.95 0.97 1e-06 ADD ADE ADD ADE

1.5

2.0

1e-08