## ENSG00000189067\_LITAF

total(k)

disease

HA

mTconv mTreg nTconv nTreg (9.38e-01)(6.65e-04)(7.31e-01)(2.27e - 09)1.22 1.22 fold change 1.11 1.05 0.95 0.82 0.82 0.9 0.9 0.67 p-value nTconv mTconv mTreg nTreg fold change 0.01 1.06 1.06 1.03 1.11 1.03 1.04 1e-04 0.95 0.97 0.97 0.96 1e-06 ADC ADD 4DE ADD ADE ADD ADD 1e-08

1.0

1.5

2.0