ENSG00000280721_LINC01943

0.2

0.3

disease

HA

MS

0.1

total(k)

mTconv mTreg nTconv nTreg (2.22e-02)(1.60e-11)(1.50e-01)1.65 1.65 fold change 1.49 1.65 1.28 0.61 0.61 0.78 0.37 0.67 0.61 0.37 0.22 p-value HA 1 HA 2 MS 1 HAHAHAHAHAMSM8M8M8M8 4 HAHAM2SMSM2SM3SMS 5 0 0.01 mTconv nTconv mTreg nTreg fold change 1.05 1e-04 1.49 2.72 1.65 1e-06 0.82 0.98 0.61 1e-08 ADC ADD ADE ADD ADE ADD ADD 1e-10