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disease HA total(k) 1.0 1.5 2.0 mTconv mTreg nTconv nTreg (1.78e-07)(7.93e-12)(1.96e-02)(2.76e - 03)1.16 1.22 **-**C fold change 1.11 1.05 1.11 1.11 1.05 0.9 0.95 0.9 0.82 0.9 0.95 0.82 p-value 0.74 0.9 0.86 0.01 nTconv mTconv mTreg nTreg fold change 1.06 1.03 1.11 1.05 1.08 1.04 1.05 1e-04 0.98 0.95 0.97 1e-06 0.95 0.96 1e-08 ADC ADD ADE ADC ADD ADE ADD ADC ADD 1e-10