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disease HA total(k) 1.5 2.0 2.5 nTconv mTconv mTreg nTreg (1.78e-07)(7.93e-12)(1.96e-02)(2.76e - 03).22 1.11 1.22 fold change 1.22 1.11 1.11 0.9 0.82 0.9 0.9 0.82 0.82 p-value 0.82 0.67 0.01 mTconv mTreg nTconv nTreg fold change 1.11 1.05 1e-04 1.11 1.05 1.05 1.04 0.95 0.96 1e-06 0.95 0.95 0.92 1e-08 ADC ADD 4DE ADD ADE ADD ADC ADD 1e-10