**Supplementary tables**

**Table S2.** Slope values, efficiencies, correlation coefficients (R2) and dilutions used for each gene in the rat retroperitoneal tissue.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **Gene** | **Slope** | **Efficiency** | **R2** | **Dilution** |
| *B2m* | -3.65 | 0.88 | 0.99 | 0.1 |
| *Bax* | -3.64 | 0.88 | 0.99 | 0.1 |
| *Bcl2* | -3.23 | 1.03 | 0.97 | 0.1 |
| *Bmal1* | -3.33 | 0.99 | 0.98 | 0.1 |
| *Casp3* | -3.02 | 1.14 | 0.99 | 0.1 |
| *Casp8* | -3.77 | 0.84 | 1.00 | 0.1 |
| *Casp9* | -3.65 | 0.88 | 0.99 | 0.1 |
| *Clock* | -2.91 | 1.20 | 0.99 | 0.1 |
| *Cry1* | -3.28 | 1.01 | 0.99 | 0.1 |
| *Fas* | -3.09 | 1.05 | 0.99 | 0.1 |
| *Nr1d1* | -3.20 | 1.05 | 0.96 | 0.1 |
| *Per2* | -3.14 | 1.07 | 0.99 | 0.1 |
| *Tnfrsf1a* | -3.40 | 0.97 | 0.99 | 0.1 |