#### Supporting information for:

# The NRON complex controls circadian clock function through regulated PER and CRY nuclear translocation

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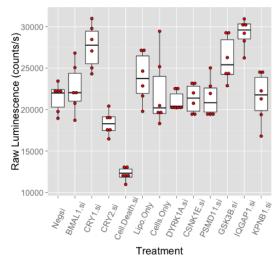
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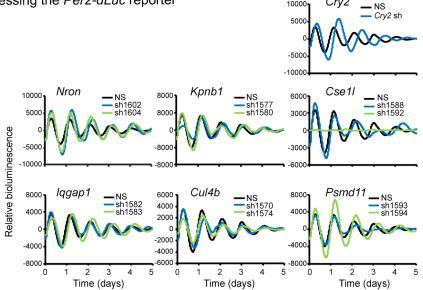
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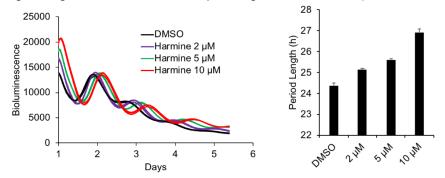
A ATP-lite cell death assay 96 h post transfection



**B** shRNA knockdown of the NRON complex components in MMH-D3 cells expressing the *Per2-dLuc* reporter

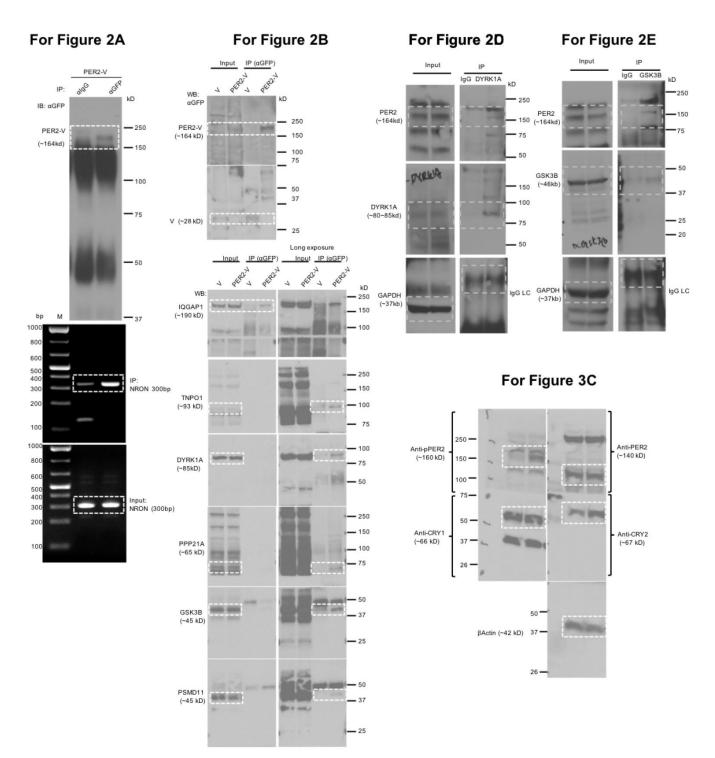


**C** Inhibition of DYRK1A with Harmine causes dose-dependent period lengthening effect in U2OS cells expressing the *Per2-dLuc* reporter

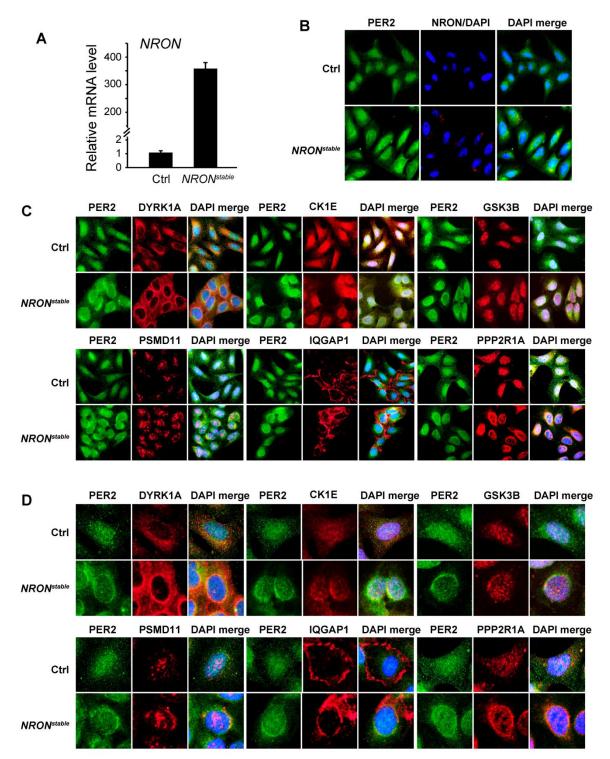


**Figure S1.** Effects of RNAi knockdown of the NRON complex components on circadian oscillations in cellular clock models. **(A)** Cytotoxicity of selected siRNA pools using the ATP-lite cell viability assay 96 h post-

transfection (n= 6 replicates per treatment). The Cell Death siRNA (Cell.Death.si) pool had ~50% cell viability compared to the negative siRNA control (Negsi) at this time point. **(B)** shRNA knockdown effects of the NRON complex components in MMH-D3 hepatocyte cells expressing the *Per2-dLuc* reporter. *Nron, Kpnb1, Cse1l, Iqgap1, Cul4b* and *Psmd11* are the NRON complex components. NS (non-specific), negative control. *Cry2*, positive control. **(C)** Pharmacological inhibition of DRYK1A lengthens period length in U2OS cells expressing the *Per2-dLuc* reporter. Representative bioluminescence records (left) and quantified period length (right) are shown for DMSO and different dosages of Harmine.



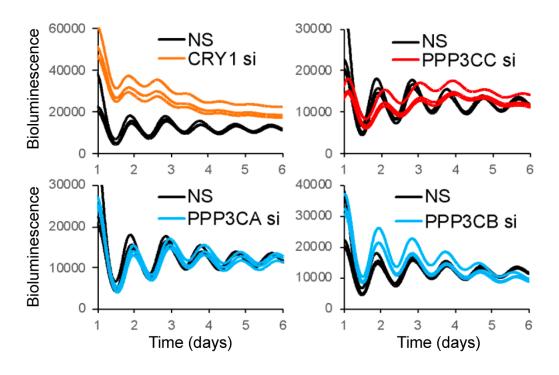
**Figure S2.** Original blot data for Figure 2 and Figure 3C. The clipped blot images are indicated with white dashed rectangles.



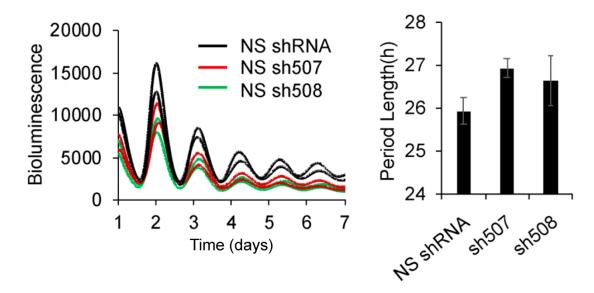
**Figure S3.** PER2 colocalizes with NRON complex components at the perinuclear region. **(A)** Quantitative RT-PCR analysis of NRON expression in control U2OS cells (Ctrl) and cells stably overexpressing *NRON* (*NRON*<sup>Stable</sup>). Data are mean ± SE of triplicate samples. **(B)** Combined RNA fluorescence in situ hybridization (RNA FISH) and immunofluorescence analysis of subcellular localization of *NRON* and PER2. Fluorescence microscopy filter sets: FITC (green, PER), TRITC (red, *NRON*), and DAPI (blue, nucleus). **(C)** Immunofluorescence analysis of subcellular localization of PER2 and NRON complex components.

Fluorescence microscopy filter sets: FITC (green, PER2), TRITC (red, NRON complex components as indicated), and DAPI (blue; nucleus). **(D)** Representative single cell images from (C).

A siRNA knockdown of the catalytic subunits of calcineurin (*PPP3CA, CB* and *CC*) in human U2OS cells expressing the *Per2-dLuc* reporter

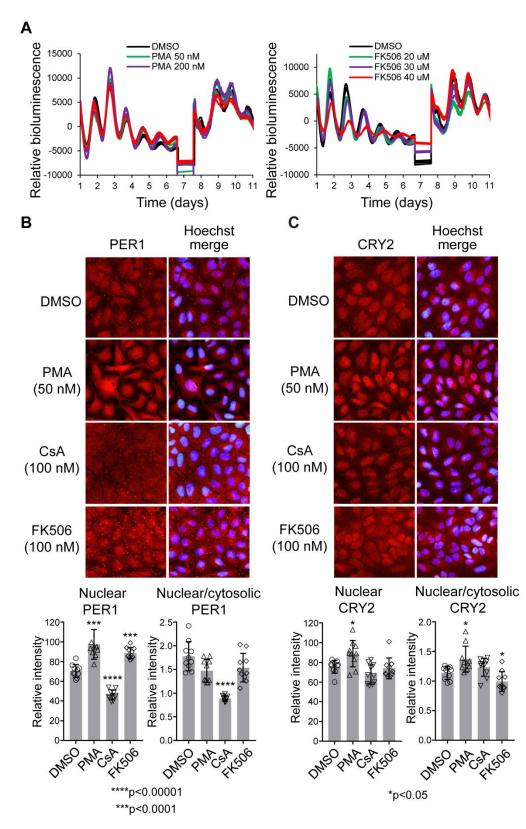


**B** shRNA knockdown of the catalytic subunit of calcineurin (*Ppp3cc*) in mouse MMH-D3 cells expressing the *Per2-dLuc* reporter



**Figure S4.** Effects of knockdown of the catalytic subunit (*PPP3CC*) of calcineurin (CaN) on period length. (A) siRNA knockdown of the catalytic subunits of CaN (*PPP3CA, PPP3CB* and *PPP3CC*) in human U2OS cell expressing the *Per2-dLuc* reporter. NS, non-specific. *CRY1* serves as a positive control. (B) Lentiviral shRNA knockdown of *Ppp3cc* caused long period length in mouse MMH-D3 cells expressing the *Per2-dLuc* reporter.

Two independent shRNA constructs against *Ppp3cc* were used. NS, non-specific. Representative bioluminescence records (left) and quantified period length data (right) are shown.



**Figure S5.** Effects of pharmacological perturbation of the NRON/NFAT pathway on circadian rhythms and nuclear localization of PER1 and CRY2 in U2 OS cells. PMA, activator. CsA and FK506, inhibitors. **(A)** Representative bioluminescence records of circadian rhythms in U2OS cells expressing the *Bmal1-Luc* 

reporter upon PMA (left) or FK506 (right) treatment. Cellular toxicity was not shown in varying doses of PMA (50~200 nM) and FK506 (20~40  $\mu$ M) as bioluminescence rhythms of the the drug-treated cells were reactivated as robustly as DMSO-treated controls when resynchronized. (**B-C**) PER1 and CRY2 proteins were detected by immunostaining using anti-PER1 and anti-CRY2 antibodies. Representative images (upper) are shown with quantitative assessment of nuclear/cytoplasmic ratio (lower) for the effect on PER1 and CRY2 nuclear abundance after 6 h of treatment. Data are presented as mean ± SD (n = 10); \*p<0.05, \*\*\*p<0.00001 by two-tailed student's t-test.