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Sustained hemodynamic stress disrupts normal circadian rhythms in calcineurin-dependent signaling and protein phosphorylation in the heart

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Abstract

Rationale—Despite overwhelming evidence of the importance of circadian rhythms in cardiovascular health and disease, little is known regarding the circadian regulation of intracellular signaling pathways controlling cardiac function and remodeling.

Objective—To assess circadian changes in processes dependent on the protein phosphatase calcineurin, relative to changes in phosphorylation of cardiac proteins, in normal, hypertrophic, and failing hearts.

Methods and Results—We found evidence of large circadian oscillations in calcineurin-dependent activities in the left ventricle of healthy C57BL/6 mice. Calcineurin-dependent transcript levels and nuclear occupancy of the nuclear factor of activated T cells (NFAT) regularly fluctuated as much as 20-fold over the course of a day, peaking in the morning when mice enter a period of rest. Phosphorylation of the protein phosphatase 1 inhibitor 1 (I-1), a direct calcineurin substrate, and phospholamban (PLB), an indirect target, oscillated directly out of phase with calcineurin-dependent signaling. Using a surgical model of cardiac pressure overload, we found that, although calcineurin-dependent activities were markedly elevated, the circadian pattern of

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activation was maintained, whereas, oscillations in PLB and I-1 phosphorylation were lost. Changes in the expression of fetal gene markers of heart failure did not mirror the rhythm in calcineurin/NFAT activation suggesting that these may not be direct transcriptional target genes. Cardiac function in mice subjected to pressure overload was significantly lower in the morning than in the evening when assessed by echocardiography.

Conclusions—Normal, opposing circadian oscillations in calcineurin-dependent activities and phosphorylation of proteins that regulate contractility are disrupted in heart failure.

Keywords

Calcineurin; circadian rhythms; heart failure; RCAN1/MCIP1

Introduction

Circadian rhythms are self-sustaining, 24-hour cycles in molecular, biochemical, and behavioral parameters that help an organism prepare for anticipated changes in physiological demand. Many important cardiovascular factors, including metabolism, heart rate, blood pressure, and hormone release, oscillate over a 24-hour period.¹ In humans, the incidence of adverse cardiac events, such as myocardial infarction, ventricular tachycardia, and death from ischemic heart disease, vary according to the time of day.² Despite overwhelming evidence of the importance of circadian rhythms in cardiovascular health and disease, little is known regarding the circadian regulation of intracellular signaling pathways in the heart.

The molecular basis of the circadian clock consists of cell-autonomous, positive and negative transcriptional and posttranscriptional feedback loops.³ The “master clock”, located in the suprachiasmatic nucleus within the hypothalamus, influences the phase of independent molecular clocks found in peripheral organs, including the heart. Many cells and tissues also display circadian fluctuations in cytoplasmic Ca^{2+} levels, although the source of these Ca^{2+} oscillations and their relationship to the transcriptional clock mechanism is not fully understood.⁴

Dysregulation of Ca^{2+} handling is a hallmark of heart disease. Several Ca^{2+} -responsive signaling pathways, including the protein phosphatase calcineurin, have been causally linked to the progression of heart failure.⁵ Sustained activation of calcineurin is sufficient to drive pathological hypertrophic remodeling of the myocardium with subsequent heart failure and premature death.⁶ Calcineurin is activated by an elevation in intracellular Ca^{2+} and binding of a Ca^{2+} /calmodulin complex. In a healthy heart, calcineurin is thought to be inactive and unresponsive to high amplitude, high frequency, waves of Ca^{2+} that drive contraction. Calcineurin is activated in response to stress presumably when either diastolic resting Ca^{2+} or Ca^{2+} in sub-cellular domains exceed a required threshold.

Calcineurin has numerous substrates including the transcription factor nuclear factor of activated T cells (NFAT) through which calcineurin influences long term changes in gene expression associated with pathological cardiac remodeling.⁶ When NFAT is dephosphorylated, it translocates to the nucleus where it binds to and activates calcineurin-responsive genes. Among target genes is the exon 4 isoform of the Regulator of Calcineurin 1 (*Rcan1.4*), previously called *MCIP1*, *DSCR1* or *calcipressin*. RCAN1 proteins are potent inhibitors of calcineurin activity.⁷ Expression of the mouse *Rcan1.4* gene is extremely responsive to changes in calcineurin activity *in vivo*.⁸ thus, altered *Rcan1.4* transcript levels have been used as a sensitive indicator of changes in calcineurin activity in the heart and other tissues.

Both Ca^{2+} handling and cardiac myocyte contractility are regulated by changes in phosphorylation of key proteins.⁹ β -adrenergic stimulation activates the cAMP-dependent protein kinase (PKA), which increases cardiac output by phosphorylating a number of proteins including phospholamban (PLB). This releases inhibition of the sarcoplasmic reticulum Ca^{2+} ATPase (SERCA2), thereby enhancing relaxation rate and contractility. PLB and other regulatory proteins are dephosphorylated by the protein phosphatase 1 (PP1). Phosphorylation of the PP1 inhibitor-1 (I-1) by PKA at threonine 35 (I-1^{Thr35}) prolongs β -adrenergic responses by inhibiting PP1, thus slowing dephosphorylation of PLB.¹⁰ Changes in I-1 levels and/or phosphorylation have been implicated in human heart failure and chronic atrial fibrillation.^{9, 11} I-1 can be phosphorylated at other sites including Ser⁶⁷.¹² The *in vivo* consequence of phosphorylation at these various sites remains controversial, however, calcineurin can dephosphorylate both I-1^{Ser67} and I-1^{Thr35}.^{13, 12} *In vitro* and *in vivo* studies suggest that calcineurin activity can promote dephosphorylation of PLB via regulation of I-1.^{14, 15}

Given the need for the heart to adapt to daily changes in cardiac demand and the potentially antagonistic roles of PKA and calcineurin, we asked whether calcineurin activity and/or PLB phosphorylation change over the course of 24 hours in a healthy heart. We found circadian oscillations in both these parameters that were directly out of phase with each other. We then tested what happens to these rhythms when both β -adrenergic activity and calcineurin activity increase in the pressure stressed myocardium.

Methods

An expanded Methods section is available in the Online Data Supplement at <http://circres.ahajournals.org> that provides expanded details for *in situ* hybridization, immunohistochemistry, chromatin immunoprecipitation, quantitative RT-PCR and western blot analysis.

Animal Procedures

Male C57BL/6 mice were housed and fed under standard laboratory conditions with a strict 12:12 hour light:dark cycle with lights turning on at 6AM, circadian time 0 (CT0), and off at 6PM (CT12). For pressure overload experiments, mice were subjected to thoracic aortic constriction (TAC) or severe TAC (sTAC) for three weeks as described previously.¹⁶ Mice were shifted to constant darkness at the end of the normal light cycle for 24 hours prior to harvesting. Hearts were removed and the ventricles flash frozen within 30 seconds of sacrifice to preserve phosphorylation. A minimum of three animals was analyzed for each time point. A VisualSonics Vevo 770 imaging system was used to assess cardiac function in unanesthetized animals. The *α MHC-RCAN1* mice were described previously¹⁷ and wild type littermates used as controls. Surgically implanted mini-osmotic pump (Alzet, Palo Alto, CA) were used to deliver cyclosporine at a rate of 50 μg per hour per 25 kg of body weight.

Results

Changes in RCAN1.4 Protein and mRNA Levels Display Circadian Rhythmicity

Biochemical assays of calcineurin activity are limited to measuring the potential activity of the entire cellular pool of calcineurin, rather than the fraction of the pool that was active *in vivo*. We therefore used multiple indirect methods to assess calcineurin activity. Initially, we quantified changes in both protein and mRNA levels of the *Rcan1.4* gene, a direct target of calcineurin/NFAT. Male C57BL/6 mice were entrained to a 12:12 light:dark cycle then shifted to constant darkness at circadian time 12 (CT12) the day before samples were harvested for analysis. RCAN1.4 protein levels were highest in the heart at the beginning of

the day (CT1-CT3) and lowest at the end of the day (CT11-CT13) (Figure 1A). In comparison, there were no significant changes in either the level of the exon 1 isoform of RCAN1 (RCAN1.1) or tubulin. A similar circadian pattern in RCAN1.4 protein levels was found in the hearts of 129/Sv and C3H/He inbred lines demonstrating that the oscillations were not strain dependent (data not shown). These findings are consistent with genome-wide microarray analysis identifying *Rcan1* as having a circadian pattern of mRNA expression in the mouse heart.¹⁸ We found a twenty-fold oscillation in *Rcan1.4* mRNA levels with a peak at CT23-CT1 and a trough at CT11 (Figure 1B) directly preceding the peak and trough in RCAN1.4 protein levels. In contrast, there were no significant circadian changes in the transcript levels of either *Rcan1.1* or *I-1* (Figure 1C and 1D). Thus, circadian regulation of *Rcan1* expression is unique to the *Rcan1.4* isoform and controlled at the level of transcript abundance. Transcription of the circadian clock gene *Period 2* (*Per2*) oscillated with 24-hour periodicity (Figure 1E) verifying the presence of a functional clock in these samples.

Calcineurin-dependent Signaling is Most Active in a Mouse Heart as the Animal Enters a Period of Decreased Physical Activity

Immunohistochemical analysis for NFATc1 in the left ventricle revealed nuclear staining at 6 AM (CT0) (Figure 2A) but not at 6 PM (CT12) (Figure 2B). Although only a modest number of nuclei stained positive for NFATc1 even at the peak of activity, these positive nuclei were always embedded within sarcomere-positive cells and never observed in non-myocyte nuclei (Figure 2C and 2D).

NFAT binding to the *Rcan1.4* promoter was assessed by chromatin immunoprecipitation studies. A six-fold increase in NFAT occupancy of the *Rcan1.4* promoter was detected at CT0 compared to at CT12 (Figure 2E) verifying that the circadian expression pattern of *Rcan1.4* was driven by changes in NFAT nuclear translocation. An *in situ* hybridization specific for *Rcan1.4* indicated that transcription was elevated uniformly across the wall of the myocardium (Figure 2F and 2G). Peak *Rcan1.4* transcript levels were blunted in the hearts of mice with cardiomyocyte-specific expression of a transgene encoding RCAN1 to inhibit calcineurin (Figure 3A) or treated with the calcineurin inhibitor cyclosporine (Figure 3B). Taken as a whole, these results suggest that activation of the calcineurin/NFAT signaling pathway occurs throughout the left ventricular myocardium and is greatest when the animal is entering its rest phase and cardiac demand decreases.

Phosphorylation of I-1 and PLB Oscillates out of Phase with Calcineurin Activity

Immunoblot analyses were conducted to assess phosphorylation of the calcineurin substrate I-1 at Thr³⁵ and Ser⁶⁷. Unfortunately, we were not able to detect phosphorylation of Thr³⁵ in either heart extracts or forskolin-treated cells transfected with an I-1 expression construct (data not shown). There were, however, pronounced circadian changes in Ser⁶⁷ phosphorylation. Phospho-I-1^{Ser67} was lowest in the morning (CT1-8), increased notably at CT11 as the animals became active, and peaked at CT14 directly opposed to circadian changes in calcineurin activity (Figure 3C and 3D). In contrast, total I-1 protein (Figure 3C) and transcript levels (Figure 1D) did not change.

Cardiac contractility and β -adrenergic drive are both higher at night in the hearts of nocturnal rodents.^{19, 20} Antibodies specific for phospho-PLB^{Ser16} showed a peak at CT14, coincident with the peak in I-1^{Ser67} phosphorylation (Figure 3C). The change in phosphorylation was even more pronounced when protein extracts were run such that PLB was maintained as a pentameric complex. Phospho-PLB^{Ser16} and phospho-PLB^{Thr17} were both elevated at CT14 compared to CT2 (Figure 4A, 4B, 4D, and 4E). This was evident in the slower electrophoretic migration of total PLB complexes from CT14 lysates (Figure 4C).

Thus, the overall phosphorylation state of PLB was elevated during the period when calcineurin-dependent activities were lowest.

Although we were not able to detect phosphorylation of I-1 at Thr³⁵, we predict that phosphorylation of this site by PKA should parallel PLB^{Ser16} phosphorylation. The kinase responsible for I-1^{Ser67} phosphorylation in the heart has not yet been identified definitively. To test whether I-1^{Ser67} is phosphorylated in response to β -adrenergic stimulation, we injected wild type mice with 200ng of isoproterenol. Phospho-I-1^{Ser67} was maximal 5 minutes after injection and dissipated within an hour (Figure 4F). Phospho-PLB^{Ser16} followed a similar time course (Figure 4G) suggesting that, in response to β -adrenergic stimulation, the kinetics of I-1^{Ser67} phosphorylation and dephosphorylation may be similar to that of PLB^{Ser16}.

It is likely that the flux through many kinase/phosphatase signaling pathways display circadian rhythmicity in the heart. Glycogen synthase kinase 3-beta (GSK3 β) can antagonize calcineurin activity by rephosphorylating NFAT.²¹ We found a mild increase in GSK3 β phosphorylation indicating inactivation of GSK3 β at CT14 (Figure 3C and 3E), consistent with insulin activation of Akt kinase when the mouse begins to feed. Therefore, circadian changes in GSK3 β activity are not responsible for the observed changes in NFAT localization. We also assayed for changes in *Nfatc1* and *Nfatc3* transcript levels which did not change significantly (data not shown), and thus, are unlikely to account for the magnitude of the changes observed in *Rcan1.4* mRNA and NFAT nuclear localization. Taken as a whole, we document an inverse correlation between calcineurin-dependent activities and the phosphorylation of proteins that either inhibit PP1 or promote contractility.

Calcineurin Activity is Elevated but Maintains Rhythmicity in Hypertrophic and Failing Hearts

To determine whether cardiac hypertrophy or failure alters the temporal relationship between changes in calcineurin activity and protein phosphorylation, mice were subjected to either TAC surgery to induce stable compensated hypertrophy or severe sTAC to induce decompensated heart failure.¹⁶ Three weeks after surgery hearts from TAC mice showed a 29% increase in heart weight to body weight ratio compared to sham operated controls (Figure 5A). Cardiac function was preserved in these mice as assessed by echocardiography (data not shown) and lung weight (Figure 5B). In sTAC mice there was a 78% increase in heart weight and a doubling of lung weight compared to sham controls indicating that these mice had progressed to heart failure. Cardiac function was significantly reduced after sTAC (Figure 5C). Remarkably, the decline in function was more pronounced when echocardiography was performed in the morning (AM) between 5AM and 7AM than if function was assessed in the evening (PM) between 5PM and 7PM at the end of the light period, regardless of whether the initial echo was performed in the morning or evening. The time of day differences in function in the sTAC mice were primarily systolic (Figure 5D), whereas, the decline in diastolic function was similar regardless of the time of day (Figure 5E). In the control sham mice there was also a trend toward improved function if the mice were echoed in the evening, but the difference did not reach statistical significance.

In both TAC and sTAC hearts *Rcan1.4* mRNA levels were elevated at all times of the day compared to controls, however, circadian rhythmicity was maintained in both the hypertrophic and failing hearts (Figure 6A), suggesting that, although calcineurin activity increased greatly, it retained a circadian pattern of activation. Maximal *Rcan1.4* expression was only slightly higher in failing hearts compared to hypertrophic hearts. However, trough expression at CT12 was much higher in the sTAC hearts than in the TAC hearts. Our results suggest that although calcineurin activity remains circadian, there is a progressive elevation

in trough activity and a dampening of the fold change between peak and trough activity with increasing stress.

Rcan1.1 transcript levels remained constant throughout (Figure 6B). *Per2* expression retained rhythmicity (Figure 6C) indicating maintenance of a cardiac-specific circadian clock, although *Per2* oscillation was dampened slightly in sTAC hearts, consistent with previous reports.²² Expression of the fetal genes beta myosin heavy chain (*βMHC*) and atrial natriuretic factor (*ANF*) was elevated in TAC and sTAC hearts compared to sham controls indicative of cardiac stress and failure (Figure 6D and 6E). Expression of the high-affinity glucose transporter (*Glut4*) was depressed in TAC and sTAC hearts compared to sham controls also consistent with a failure phenotype^{22, 23} (Figure 6F). Importantly, there were no pronounced circadian oscillations in *ANF*, *βMHC*, and *Glut4* expression in control hearts suggesting that daily activation of calcineurin in healthy hearts is not sufficient to alter expression of these genes associated with heart failure. Whether the circadian changes in *βMHC* and *ANF* expression observed in the TAC and sTAC hearts are linked to changes in calcineurin activity or other signaling pathways is not yet known.

The Circadian Pattern of PLB and I-1 Phosphorylation is Disrupted

Western blot analysis of RCAN1.4 protein in control, TAC and sTAC hearts at CT2 and CT14 reflect the changes seen in *Rcan1.4* mRNA (Figure 7A and 7B). Within each experimental group RCAN1.4 protein was more abundant at CT2 than at CT14. Within a given time point RCAN1.4 protein was consistently higher in the TAC and sTAC hearts than in control hearts.

The normal circadian pattern of I-1 and PLB phosphorylation was lost in the TAC and sTAC hearts. Phosphorylation of PLB was elevated throughout the day in both TAC and sTAC hearts (Figure 7C, Figure 7D and Online Figure I). This is most easily seen by comparing the delay in migration of the PLB pentamer in the TAC and sTAC samples to the migration of PLB in the control sham samples at CT2. Importantly, phosphorylation of I-1^{Ser67} could not be detected in either TAC or sTAC hearts at any time of day (Figure 7E and 7F), demonstrating that there is a loss of phosphorylation in addition to a loss of circadian changes. This was not due to loss of total I-1 protein levels as these were similar in control, TAC and sTAC hearts.

Discussion

These studies provide several important new insights into the dynamics and consequences of calcineurin signaling in the heart. First, we present evidence that there is a circadian rhythm in calcineurin activity in a normal, healthy mouse heart. This assertion is supported by five different assessments of calcineurin activity: RCAN1.4 protein levels, *Rcan1.4* mRNA levels, NFATc1 nuclear translocation, NFATc1 occupancy of the *Rcan1.4* promoter, and phosphorylation levels of the calcineurin substrate I-1^{Ser67}. Second, although activation of calcineurin has been viewed primarily as a stress response driving pathological hypertrophic remodeling, these daily increases in calcineurin activity are not associated with transcriptional changes indicative of pathological remodeling. Third, calcineurin activity oscillates out of phase with phosphorylation of proteins that promote cardiac contractility. Fourth, cardiac function in failing hearts shows diurnal variation. Finally, in failing hearts, calcineurin activity is elevated above normal peak activity throughout the day. Although a pronounced circadian oscillation in calcineurin-dependent activities persists, the amplitude of circadian variation declines correlating to a decline in cardiac function. Circadian rhythmicity in the phosphorylation of the regulatory proteins I-1 and PLB is lost in hypertrophic and failing hearts. Based on our findings we propose that circadian oscillations provide temporal separation of kinase and phosphatase activities that help to regulate

changes in cardiac function in response to physiological demand and that this temporal relationship is disrupted when the heart is placed under sustained hemodynamic load.

In nocturnal animals, the light to dark transition at CT12 is anticipated by a rapid increase in physical activity, blood pressure, and β -adrenergic drive,²⁴ all factors shown to increase calcineurin activity in cardiomyocytes. We therefore expected *Rcan1.4* expression to increase in parallel with these physiological parameters at dusk. Instead, there was a gradual increase in *Rcan1.4* expression throughout the course of the night suggesting a progressive activation of calcineurin that peaked at dawn when β -adrenergic drive is minimal in a mouse. Figure 8A depicts a model of how calcineurin activity could crosstalk with cardiac function via dephosphorylation of I-1, thus promoting reversal of phosphorylation-driven increases in cardiac contractility as discussed in the introduction. The findings we present here document that in a normal heart, circadian control provides temporal separation of these opposing processes as schematized in Figure 8B.

The peak in NFAT activity and *Rcan1.4* transcription at dawn (CT0) implies that calcineurin activity and cytoplasmic Ca^{2+} levels are also maximal at this time of day. Conversely, Ca^{2+} levels are likely lowest at dusk (CT12) when these indicators of calcineurin activity are lowest. Oscillating out of phase with calcineurin-dependent activities is phosphorylation of I-1 and PLB in response to increased β -adrenergic drive as the animal wakes.

Phosphorylation of PLB, releases inhibition of SERCA2 increasing the rate of Ca^{2+} uptake, consequently lowering resting cytoplasmic Ca^{2+} levels, and potentially helping to maintain calcineurin in an inactive state. Eventually, cardiomyocytes become desensitized to prolonged β -adrenergic stimulation leading to a decrease in I-1 and PLB phosphorylation by PKA. Restored inhibition of SERCA2 by PLB would cause a gradual rise in cytoplasmic Ca^{2+} consistent with a gradual increase in calcineurin activity during the second half of the night. We postulate that the peak in calcineurin activity at the dark to light transition (CT0) helps maintain PLB and other key proteins in an unphosphorylated state as the mouse transitions to a time of lower cardiac demand. Conversely, a trough in calcineurin activity around dusk (CT12) would allow maximal β -adrenergic responses when the animal enters its waking hours. Consistent with this, fractional shortening of sTAC mice was higher in the evening than in the morning (Figure 5C) potentially reflecting circadian differences in the adrenergic response of the unanaesthetized mice to handling. Likewise, a recent study demonstrated circadian differences in the response of isolated adult cardiomyocytes to β -adrenergic stimulation.²⁵

Although in this study we used changes in *Rcan1.4* transcript levels primarily as an indication of changes in calcineurin activity, the RCAN1 protein itself likely contributes to shaping the dynamics of calcineurin-dependent signaling. It is interesting to note that cardiac damage from ischemia-reperfusion (I/R) is greater in mice lacking *Rcan1* than in wild type hearts.²⁶ Furthermore, I/R damage in wild type mice has been shown to be greater when the procedure is performed at CT12 than when I/R is performed at CT0.²⁷ This time of greater susceptibility corresponds to the time when we find RCAN1.4 protein levels in the heart are lowest.

An important short-coming of the current study is our inability to monitor changes in I-1^{Thr35} phosphorylation and the as yet incomplete understanding of the cumulative effect of changes in I-1 phosphorylation at other sites including Ser⁶⁷. Taking these limitations into account, the model in Figure 8B suggests that daily oscillations in Ca^{2+} handling and calcineurin activity form interdependent positive and negative feedback loops typical of circadian rhythms that result in a separation between times of day when kinase activities predominate and times of day when phosphatase activities predominate. Clearly this simple model does not take into account many additional factors that can influence calcineurin

activity, PP1 activity, and cardiac contractility. For instance, the density of β -adrenergic receptors, adenylyl cyclase activity, and phosphodiesterase activity have all been shown to undergo circadian cycling both in neurons and the heart.²⁸

Sustained hemodynamic stress elicits increases in both calcineurin activity and β -adrenergic drive. The model in Figure 8A suggests that these are opposing processes. Our data indicates that in hypertrophy and failure the normal temporal separation of calcineurin activity and phosphorylation of contractile proteins is disrupted. PLB phosphorylation was elevated throughout the day even during the animal's sedentary period. In contrast, I-1^{Ser67} phosphorylation was completely lost, consistent with increased calcineurin activity. A corresponding loss of I-1^{Thr35} phosphorylation would release PP1 inhibition and result in uncoupling of circadian changes in calcineurin activity from regulation of PLB phosphorylation (Figure 8C). Initially, release of I-1 inhibition would be compensatory, helping to reverse hyper-phosphorylation of regulatory proteins, however, sustained loss of a circadian pattern of PP1 inhibition could ultimately contribute to declining cardiac function. Consistent with our observations, patients with heart failure demonstrate an increase in PP1 activity coincident with a decrease in I-1 phosphorylation.^{29, 30}

The present studies provide a deeper understanding of the dynamics of calcineurin regulation in the heart and draw attention to the need to control for normal, underlying circadian changes in the activity of intracellular signaling pathways. In a human heart, it is likely that calcineurin activity cycles with a reverse phase compared to a mouse heart although as yet this is not known. In humans there is disproportionate ischemic activity, arrhythmic activity and acute cardiovascular events in the first few hours after waking.³¹ Interestingly this would correspond to the time of day when calcineurin activity would be lowest allowing maximal β -adrenergic responsiveness in a normal heart. Elevation of the trough in calcineurin activity in heart failure would compromise the ability to respond appropriately. There is a growing appreciation for the potential therapeutic benefit of timing drug delivery to correlate with maximal biological need.³² Our findings highlight the importance of remaining mindful of inherent circadian oscillations in the cardiovascular system during both the study and treatment of heart disease.

Novelty and Significance

What is known?

Circadian rhythms are important for maintaining cardiovascular health.

Activation of the protein phosphatase calcineurin is known to promote cardiac hypertrophy and heart failure.

What new information does this article contribute?

In healthy hearts, there is an apparent circadian rhythm in calcineurin activity that oscillates out of phase with phosphorylation of proteins, such as phospholamban (PLB) and inhibitor I (I-1), that regulate cardiac contractility.

In heart failure, calcineurin activity increases but continues to cycle, whereas cycling of PLB and I-1 phosphorylation is lost.

Summary

Despite overwhelming evidence that circadian rhythms are important in cardiovascular health and disease, little is known regarding circadian regulation of intracellular signaling pathways that control cardiac function and remodeling. Activation of calcineurin is known to promote pathological cardiac remodeling. Here we present evidence that there is a circadian rhythm in calcineurin activity in normal, healthy hearts that is not

associated with transcriptional changes indicative of pathological remodeling. Calcineurin activity oscillates out of phase with phosphorylation of proteins, such as PLB and I-1 that promote cardiac contractility. In heart failure, calcineurin activity increases but continues to cycle, whereas cycling of PLB and I-1 phosphorylation is lost. We propose a model in which daily oscillations in Ca^{2+} handling and calcineurin activity form interdependent positive and negative feedback loops typical of circadian rhythms that result in separation between times of day when kinase activities predominate and times of day when phosphatase activities predominate. These studies provide a deeper understanding of the dynamics of calcineurin regulation in the heart and draw attention to the importance of normal, underlying circadian changes in regulating the activity of intercellular signaling pathways.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Non-standard Abbreviations and Acronyms

ANF	atrial natriuretic factor
βMHC	beta myosin heavy chain
CT	circadian time
Glut4	high-affinity glucose transporter
GSK3b	glycogen synthase kinase 3 beta
I-1	PP1 inhibitor-1
I/R	ischemia-reperfusion
NFAT	nuclear factor of activated T cells
Per2	period 2
PKA	cAMP-dependent protein kinase
PLB	phospholamban
PP1	protein phosphatase 1
RCAN1	<u>R</u> egulator of <u>C</u> alcineurin 1
SERCA2	sarcoplasmic reticulum Ca^{2+} ATPase
sTAC	severe transverse aortic constriction
TAC	transverse aortic constriction
TG	transgenic
WT	wild type

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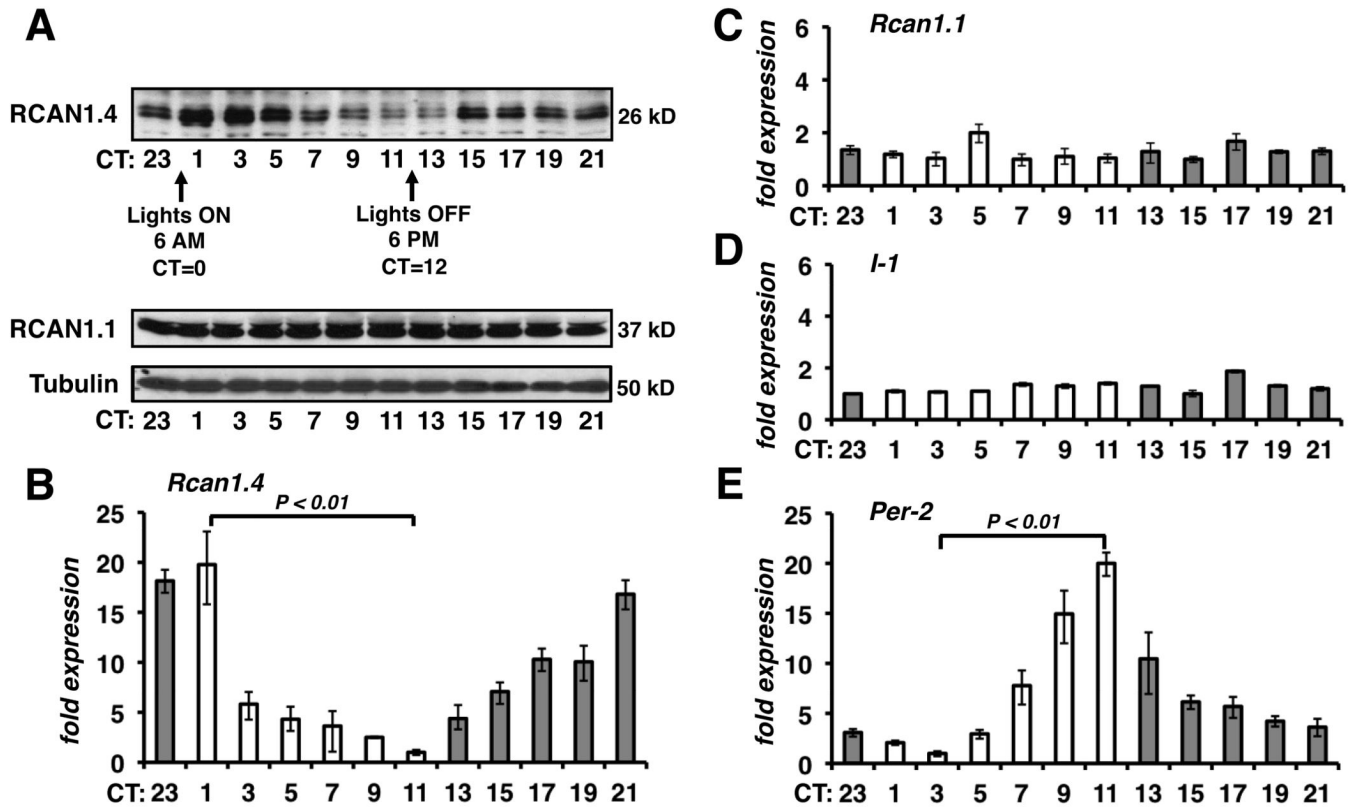


Figure 1.

Rcan1.4 mRNA and protein levels oscillate with 24-hour periodicity in the hearts of healthy mice. Samples were pooled from three mice for each time point. (A) Immunoblot analysis of RCAN1.4, RCAN1.1, and tubulin are shown. Real-time RT-PCR for *Rcan1.4* (B), *Rcan1.1* (C), *I-1* (D), and *Per-2* (E) mRNA levels were normalized to *18S rRNA*. Trough values for each gene were set at a value of 1. ($n=3$ each time point in B-E)

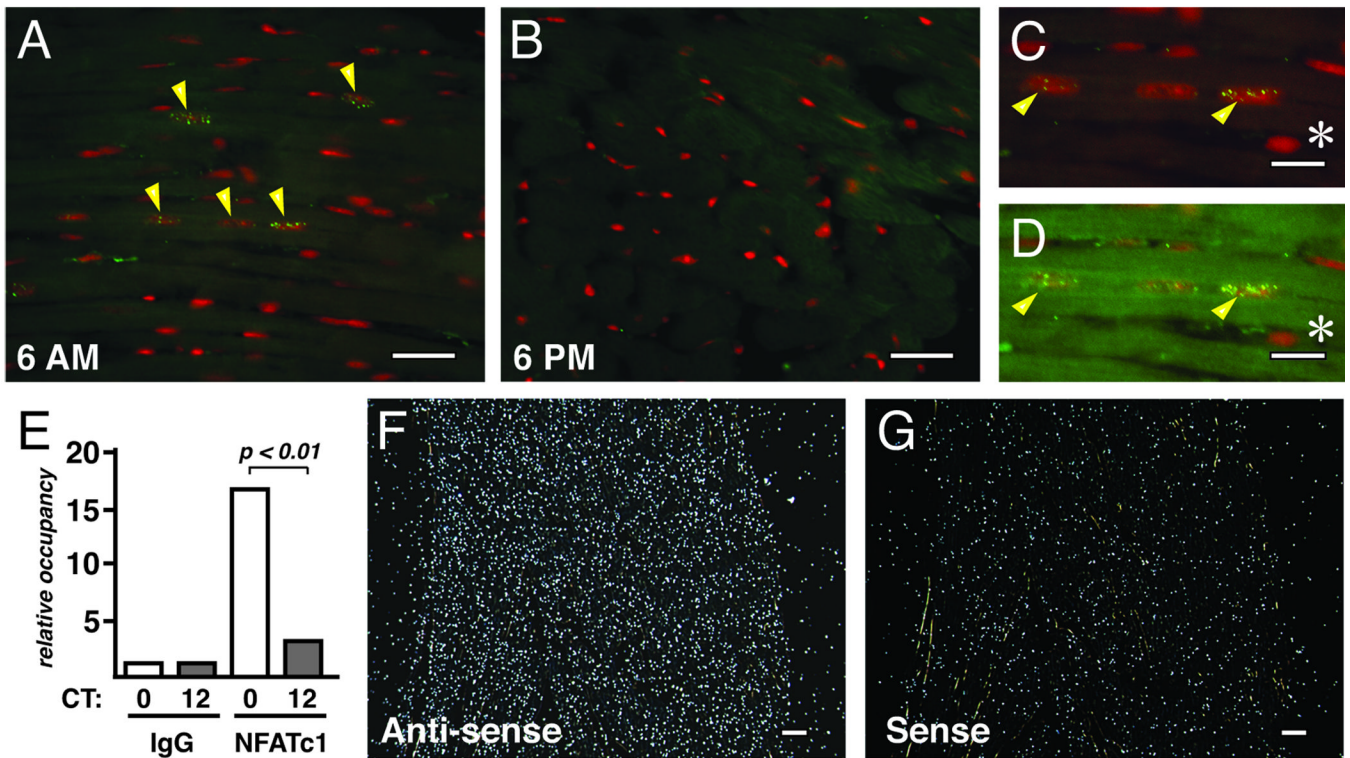


Figure 2.

More NFATc1 is located in the nucleus and bound to chromatin at CT0 than at CT12. Left ventricular free-wall harvested at 6 AM (CT0) (**A**, **C**, and **D**) and 6 PM (CT12) (**B**) were stained with a FITC-labeled NFATc1 antibody (yellow/green) and propidium iodide (PI) (red). Cardiac myocytes have a high level of autofluorescence due to the abundance of mitochondrial flavins and flavoproteins, which emit in a broad band overlapping the FITC-NFATc1 signal. In image (**C**) the gain on the green channel has been turned down to obtain a clear outline of the nuclear PI signal. In (**D**) the intensity of the green overlay has been restored so that the autofluorescence of the sarcomere now obscures the myocyte-localized NFATc1-positive nuclei marked with arrows. A non-myocyte nucleus is marked with an * and is not obscured. NFATc1 occupancy of the *RCAN1.4* promoter was determined by chromatin immunoprecipitation from ventricular lysates using either pre-immune IgG or NFATc1 antibodies (**E**) ($n=4$ each time point). *In situ* hybridization was carried out using an *Rcan1.4*-specific probe on the left ventricular wall harvested at 6 AM (CT12). Anti-sense probe (**F**) and sense probe (**G**) are shown. White bars denote 20 μ m (**A**, **B**, **F** and **G**) or 10 μ m (**C** and **D**).

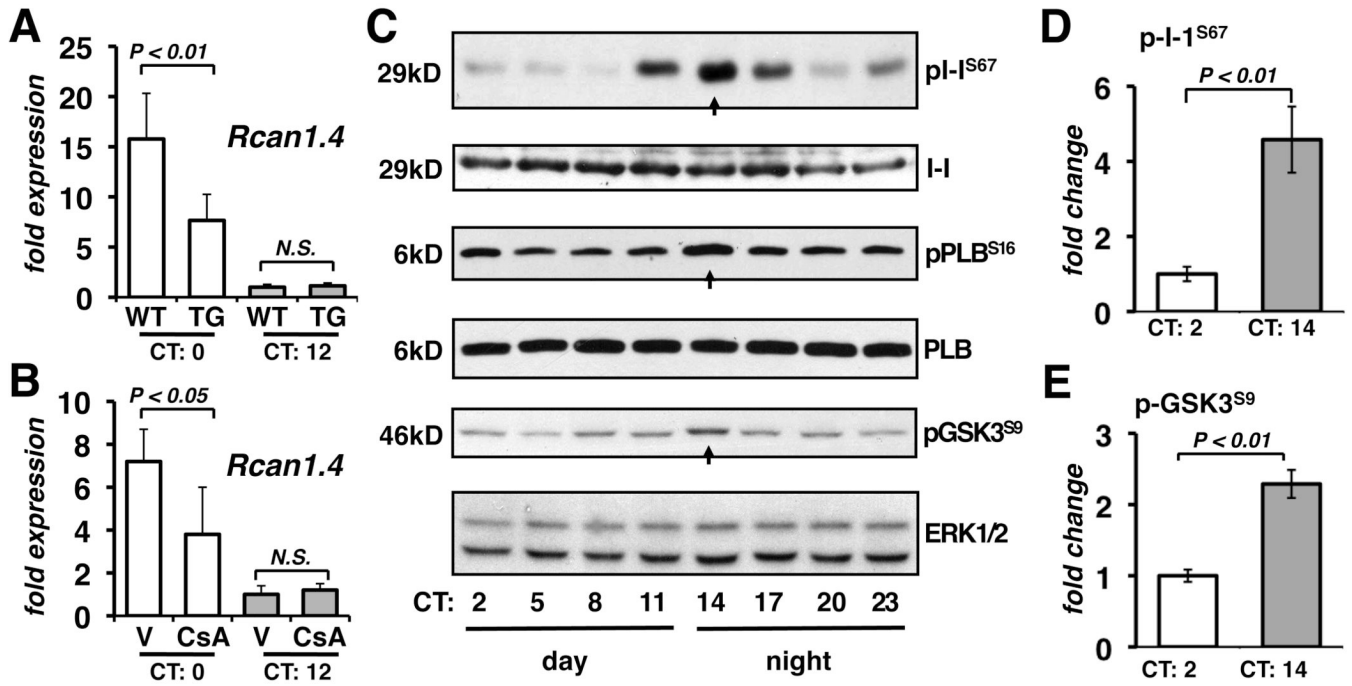


Figure 3.

Calcineurin-dependent transcription oscillates out of phase with protein phosphorylation. *Rcan1.4* transcript levels were quantified in the hearts of wild type (WT) and α MHC-*Rcan1* (TG) mice (A) or wild type mice receiving either vehicle (V) or cyclosporine A (CsA) via a mini-osmotic pump (B) ($n=4$ each). Ventricular lysates collected from three different wild type hearts at the times indicated were pooled and probed for phospho-I-1Ser67, total I-1, phospho-PLBSer16, total PLB, phospho-GSK3Ser9, and total ERK1/2 (C). Signals from three experiments were quantified by densitometry (D and E).

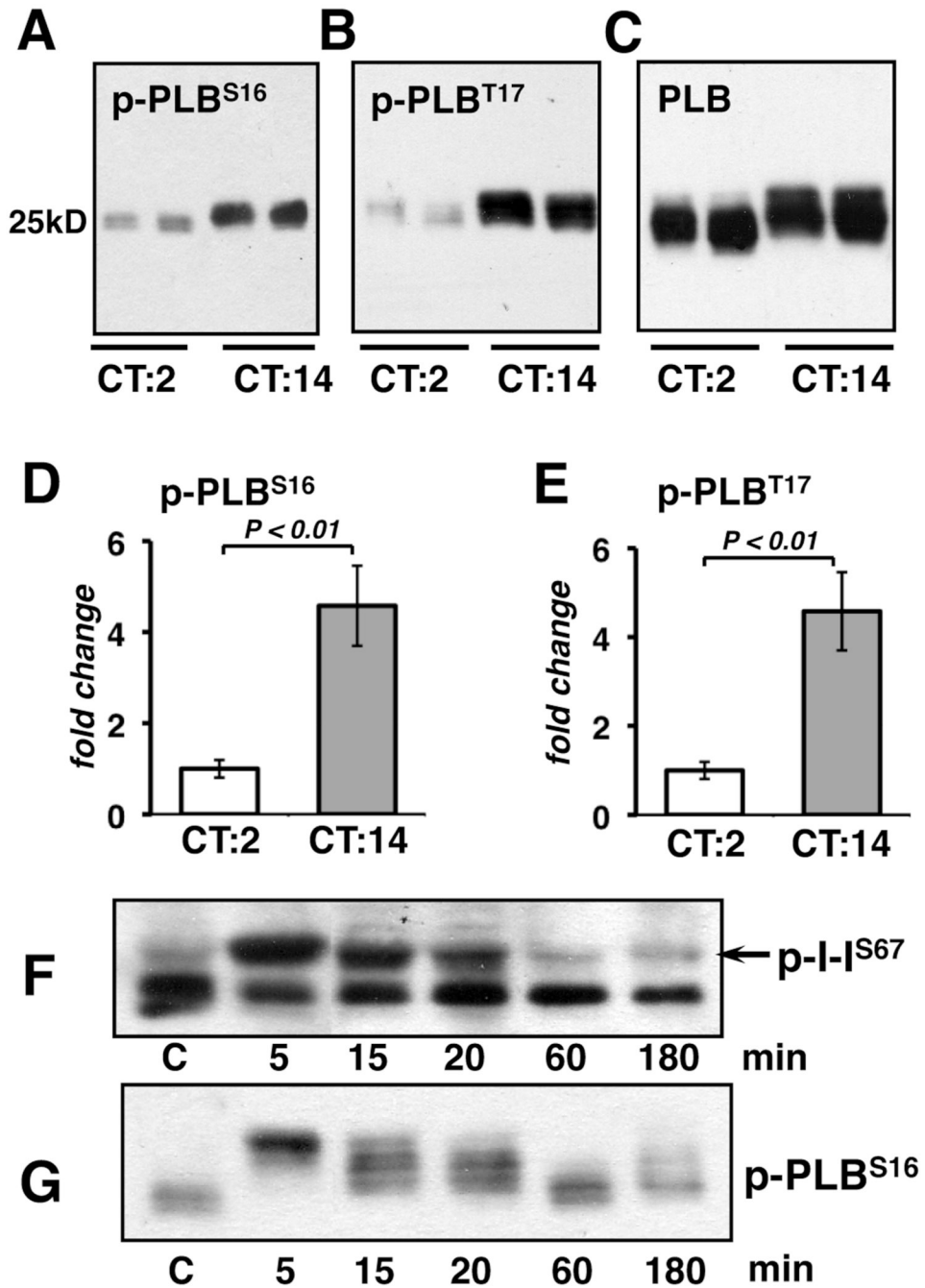


Figure 4.

Ventricular lysates isolated at CT2 and CT14 were probed for phospho-PLBSer16 (**A**: 2 μ g of protein per lane), phospho-PLBThr17 (**B**: 0.5 μ g of protein) or total PLB (**C**: 0.5 μ g of protein) per lane. Phospho-PLBSer16 and phospho-PLBThr17 were quantified by densitometry (**D** and **E**) ($n=6$ each). In (**F** and **G**) wild type mice were injected with 8 mg/kg body weight of isoproterenol at CT10. Hearts were harvested at the indicated time points after injection.

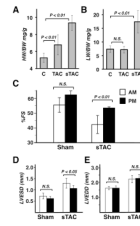


Figure 5.

Changes in cardiac parameters were assessed in control mice (c) and mice subjected to TAC or sTAC. Analysis included: heart weight to body weight ratios (**A**), lung weight to body weight ratios (**B**), as well as echocardiographic analysis to quantify percent fractional shortening (%FS) (**C**), left ventricular end systolic dimension (LVESD) (**D**), and left ventricular end diastolic dimension (LVEDD) (**E**) in Sham and sTAC mice ($n=21$ each in *A* and *B*; $n=10$ each in *C*, *D* and *E*).

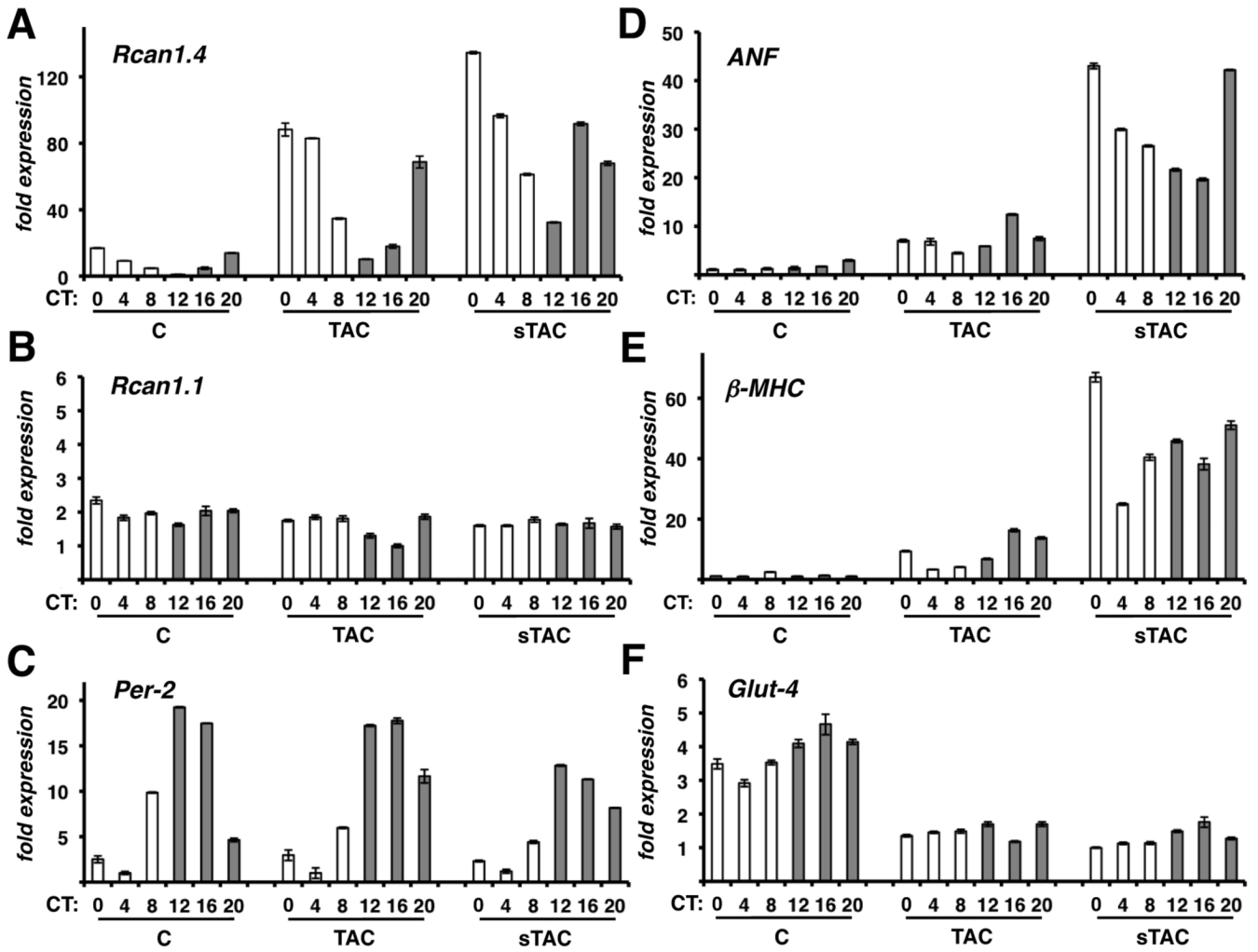


Figure 6.

Changes in gene expression was assessed in control mice (c) and mice subjected to TAC or sTAC. Real-time RT-PCR was carried out for *RCAN1.4* mRNA (A), *RCAN1.1* (B), *Per2* (C), *ANF* (D), *βMHC* (E), and *Glut-4* (F). mRNA was normalized to *18S rRNA*. ($n=3$ per time point).

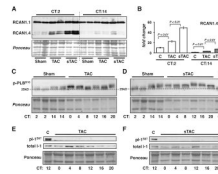


Figure 7.

Ventricular extracts isolated at CT2 and CT14 from sham operated control (c), TAB or sTAB mice were analyzed by western blot for RCAN1 protein (A). Ponceau stain of the corresponding region of the membrane is provided as a loading control. RCAN1.4 protein levels from 4–5 hearts per time point were quantified (B). Phospho-PLBSer16 (p-PLBS16) levels were compared in Sham, TAC (C) and sTAC (D) hearts at the time points indicated. The final lane in D contains extract from an unoperated control wild type mouse. Three times the amount of protein was loaded on the Sham lanes in C and D to facilitate obtaining a clear signal in the Sham lanes. The horizontal dashed line is drawn at mid-level migration of the Sham CT2 pentamer. Phospho-I-1Ser67 (pI-1S67) levels were compared in control Sham (c), TAC (E) and sTAC (F) hearts at the time points indicated. (Each lane in C, D, E and F contains samples from one heart selected randomly from an experiment with a sample size of $n=3$ per time point, per condition. PLB phosphorylation was elevated and I-1 phosphorylation was absent in all TAC and sTAC samples.)

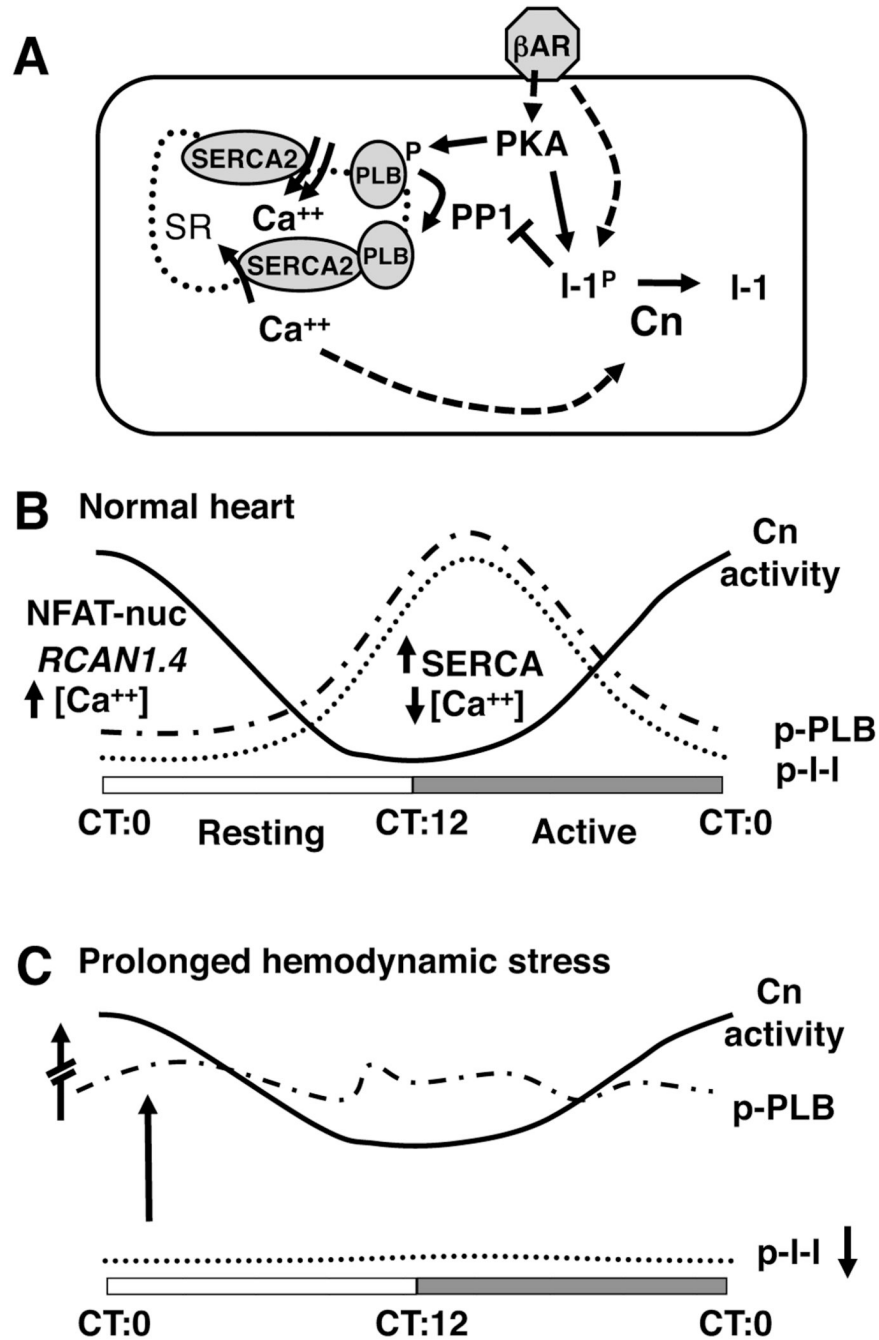


Figure 8. The model in (A) depicts interacting pathways through which PKA, PP1 and calcineurin (Cn) can influence calcium handling and contractility. Please refer to text for details. Solid arrows indicate direct interactions. Dashed arrows indicate multi-step or indirect interactions. Schematic (B) summarizes the temporal relationship between circadian oscillations in calcineurin activity, PKA-mediated phosphorylation of I-1 and PLB in the normal mouse heart. The solid line represents changes in calcineurin activity (Cn) over a 24-hour period. The dotted and broken dashed lines depict opposing changes in phosphorylation of I-1 and PLB. Schematic (C) summarizes changes observed in hypertrophic and failing hearts. Note that calcineurin activity is greatly increased but maintains circadian rhythmicity

in both hypertrophic and failing hearts. In contrast, PLB phosphorylation becomes erratic and elevated throughout the day, whereas, I-1 phosphorylation can no longer be detected.