Glucocorticoids and Stress-Induced Changes in the Expression of PERIOD1 in the Rat Forebrain

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Abstract

The secretion of glucocorticoids in mammals is under circadian control, but glucocorticoids themselves are also implicated in modulating circadian clock gene expression. We have shown that the expression of the circadian clock protein PER1 in the forebrain is modulated by stress, and that this effect is associated with changes in plasma corticosterone levels, suggesting a possible role for glucocorticoids in the mediation of stress-induced changes in the expression of PER1 in the brain. To study this, we assessed the effects of adrenalectomy and of pretreatment with the glucocorticoid receptor antagonist, mifepristone, on the expression of PER1 in select limbic and hypothalamic regions following acute exposure to a neurogenic stressor, restraint, or a systemic stressor, 2-Deoxy-D-glucose (2DG) in rats. Acute restraint suppressed PER1 expression in the oval nucleus of the bed nucleus of the stria terminalis (BNSTov) and the central nucleus of the amygdala (CEAl), whereas 2DG increased PER1 in both regions. Both stressors increased PER1 expression in the paraventricular (PVN) and dorsomedial (DMH) nuclei of the hypothalamus, and the piriform cortex (Pi). Adrenalectomy and pretreatment with mifepristone reversed the effects of both stressors on PER1 expression in the BNSTov and CEAl, and blocked their effects in the DMH. In contrast, both treatments enhanced the effects of restraint and 2DG on PER1 levels in the PVN. Stress-induced PER1 expression in the Pi was unaffected by either treatment. PER1 expression in the suprachiasmatic nucleus, the master circadian clock, was not altered by either exposure to stress or by the glucocorticoid manipulations. Together, the results demonstrate a key role for glucocorticoid signaling in stress-induced changes in PER1 expression in the brain.

Introduction

The circadian clock drives daily rhythms in behavior and physiology and these rhythms help animals adapt to their cyclic environment. In mammals, circadian rhythms depend on the
coordination of the master clock in the suprachiasmatic nucleus (SCN) with subordinate circadian clocks elsewhere in the brain and in the periphery [1, 2]. This coordination is sensitive to a range of external and systemic stressful conditions that disrupt homeostasis, leading to dysregulation of circadian physiological and behavioral rhythms [3–5]. Studies in rodents suggest that the effect of stress is independent of the SCN clock and point to changes in the expression of clock genes elsewhere in the brain and periphery as a possible mechanism [6–8]. How stress might affect clock gene expression in the brain is not fully understood, but several lines of evidence suggest that glucocorticoid hormones, primary effectors of the neuroendocrine stress system, might play a role [7, 9, 10]. For example, it has been shown that treatment with dexamethasone, a synthetic glucocorticoid, induces circadian gene expression in cultured rat-1 fibroblasts, and can transiently alter the phase of circadian gene expression in the liver, kidneys, and heart [11, 12]. Furthermore, we have shown that depletion of endogenous glucocorticoids via adrenalectomy (ADX) or selective genetic deletion of brain glucocorticoid receptors (GR), blunt the rhythm of expression of the clock protein, PER2 in nuclei of the central extended amygdala, the oval nucleus of the bed nucleus of the stria terminalis (BNSTov) and the lateral part of the central nucleus of the amygdala (CEAl), without affecting the rhythms of the SCN in rats [13–16].

The clock protein PER1 is a core component of the circadian oscillator [1, 17]. It is expressed in the SCN, as well as in many other central structures that play diverse roles in behavior and physiology including structures important in stress, autonomic control and emotion regulation [8, 18–20]. In recent work, we and others have shown that stressors known to induce the release of the adrenal glucocorticoid hormone, corticosterone (CORT), rapidly alter the expression of PER1 in various hypothalamic and limbic nuclei and in some peripheral tissues such as the liver, but not in the SCN [8, 9, 18, 20–24]. Furthermore, it has been shown that the effect of stress on Per1 expression in the liver is mediated by a glucocorticoid-responsive element (GRE) present on the promoter region of Per1, directly implicating glucocorticoids and their receptors in the stress-induced modulation of PER1 expression in the periphery [9].

In the present study, we investigated the involvement of endogenous glucocorticoids and GR in the rapid changes in PER1 expression induced by acute exposure to two qualitatively different stressors, restraint and 2-Deoxy-D-glucose (2DG) administration. While the former stressor activates the HPA axis indirectly (through corticolimbic pathways), eliciting acute emotional behavior, the latter represents a direct and invasive challenge that disrupts internal homeostasis without an affective response [25, 26]. We found previously that these two different types of stress, restraint and 2DG, exert opposite effects, decreases and increases, respectively, on PER1 expression in the BNSTov and CEAl while exerting similar effects in other forebrain regions, including the paraventricular nucleus of the hypothalamus (PVN), dorsomedial hypothalamus (DMH), and piriform cortex (Pi) in rats [21]. Here, we found that depletion of endogenous glucocorticoids via ADX or pharmacological blockade of GR (GRX) altered the acute effect of stress on PER1 expression in the forebrain in a stress- and region-specific manner, pointing to a complex interaction between stress type, glucocorticoid signaling and PER1 in the forebrain.

**Materials and Methods**

**Animals and Housing**

Animal handling and housing procedures adhered to the Canadian Council on Animal Care guidelines, and were approved by the Animal Care Committee of Concordia University. Intact, ADX and sham-operated male Wistar rats weighing 125–150 g were purchased from Charles River Laboratories (St. Constant, QC, Canada). As described in detail previously [15,21], rats...
were housed in transparent plastic cages (24cm wide x 20.5cm height x 40cm deep) individually, under a 12:12h light (100 lux at cage bottom)/dark (LD) cycle. Each cage was kept in a temperature-controlled (22°C), ventilated, sound- and light-attenuated isolation box, and was equipped with a running wheel. Running wheel data were recorded (VitalView software, Mini-Mitter, Sunriver, OR, USA) and actograms were analyzed (Actiview Biological Rhythm Analysis software, Mini-Mitter) to ensure that rats were stably entrained to the 12:12h LD cycle. Rats were allowed free access to food and water at all times. During the 2-week entrainment period, ADX rats were given CORT in their drinking water to prevent disruption of normal PER rhythms as previously described [15, 16]. CORT (Sigma-Aldrich, Oakville, ON, Canada) was dissolved in 0.9% saline, at a concentration of 25 mg/l [15]. Sham rats were provided with 0.9% saline only. On the morning of the test day, prior to stress exposure, the drinking fluid of ADX rats was replaced with 0.9% saline only.

Drugs

The non-selective GR antagonist mifepristone (Sigma-Aldrich, Oakville, ON, Canada) was dissolved in 0.9% saline containing 5% dimethyl sulfoxide (Sigma-Aldrich, Oakville, ON, Canada) and 1% Tween 20 (Fischer Scientific, Ottawa, ON, Canada). The suspension was vortexed 1 min prior to drug injection, and as needed throughout the dosing. The drug was administered i.p. at a concentration of 40 mg/kg. A similar dose was shown to attenuate stress-induced changes in a number of behavioral and physiological parameters in rats [27]. Control rats received vehicle only.

Stressors

Restraint. Rats were exposed to a 30 min restraint challenge in custom-designed ventilated Plexiglas tubes (7 mm thick, internal diameter of 75 mm, adjustable in length from 130–180 mm). Control rats were handled only.

2DG. Rats received a subcutaneous injection of 250 mg/kg 2DG (Sigma-Aldrich, Oakville, ON, Canada) in 0.9% saline. Control rats were injected with vehicle only.

Plasma CORT Collection and Analysis

Rats were placed in restraining devices and then tail-clipped using a razor and capillary tube (0.5 ml) for blood sampling prior to perfusion. Blood samples were centrifuged for 10 min at 13,000 rpm, 4°C, and the plasma was extracted and stored at -80°C, as previously described [16,21]. Circulating plasma CORT levels were determined using a CORT Enzyme Immunoassay kit (Enzo Life sciences, Farmingdale, NY, USA).

Tissue Preparation and Immunohistochemistry

As described in detail previously [14,21], rats were anesthetized with sodium pentobarbital (100 mg/kg, i.p.) and perfused transcardially with 300 ml of cold 0.9% saline (4°C, 0.9% NaCl in distilled water), followed by 300 ml of cold 4% paraformaldehyde (4°C, 4% paraformaldehyde in 0.1 M phosphate buffer). Brains were gently extracted then fixed in 4% paraformaldehyde at 4°C, for approximately 24 h. A vibratome was used to slice each brain into a series of 50 μm serial coronal sections. These sections were then placed in Watson’s Cryoprotectant and stored at −20°C. PER1 immunohistochemistry was performed using an affinity-purified rabbit polyclonal antibody, raised against PER1 (1:24,000—R1177, EMD-Millipore), as previously described in depth [19,21]. Brain sections were incubated for 40 h in a primary solution that consisted of 2% normal goat serum, PER1 polyclonal rabbit antibodies, and 5% milk buffer in a
Triton Trizma-buffered saline solution (0.3% Triton, 50 mM Trizma buffer, 0.9% saline), at 4°C [21]. Following, all sections were incubated in a secondary solution, composed of biotinylated anti-rabbit IgG raised in goat (1:200, Vector Laboratories, Burlington, ON, Canada). Lastly, as described previously [19,21], the sections were incubated in a tertiary Avidin-Biotin-Peroxidase solution (Vectastain Elite ABC Kit, Vector Laboratories), thoroughly washed in a 0.5% 3,3-diaminobenzidine (DAB) solution, and then stained using a 0.5% DAB, 0.01% H$_2$O$_2$ and 600μL of 8% NiCl$_2$ solution in 150mL.

Microscopy and Data Analysis

DAB-stained sections were gently placed onto gel-coated slides, dehydrated in alcohol and Citrisolv (Fisher), then coverslipped and left to dry. A light microscope (Leica, DMR) and a Swanson rat brain atlas (Swanson, 2004, Brain Maps: Structure of Rat Brain) were used to identify the sections. As previously described in detail [21], images of the BNSTov, CEAl, PVN, DMH, Pi and SCN were captured with a Sony XC-77 video camera, Scion LG-3 frame grabber (Scion Corporation, Frederick, MD, USA), and Image SXM software (http://www.ImageSXM.org.uk1 v1.95, S.D. Barret). Using counts from six unilateral images showing the highest number of labeled nuclei, the mean number of PER1 immunoreactive (IR) cells per region was then calculated for each animal, as previously described [13,21]. Capturing of images and counting of cells were performed blind. Data were analyzed using a two-way analysis of variance (ANOVA), with a confidence level [α] set at 0.05. Differences between stressed and control groups were revealed using Bonferroni or Dunnett’s post-hoc analyses. In both the ADX and GRX experiments, data from control groups for restraint and 2DG were tested and determined to have no significant differences (data not shown), and accordingly, pooled together.

Experimental Protocol

**Adrenalectomy experiment.** On the test day, the ADX and sham-operated rats (n = 4 per group) were exposed to restraint or given an injection of 2DG, at zeitgeber time 2 (ZT2, ZT0 indicates time of lights on). Control rats for restraint were handled only and those for 2DG were injected with vehicle. Blood was collected 1 h after stress onset (ZT3) and rats were then immediately anesthetized and perfused. The time points selected were based on our previous findings that demonstrated maximal changes in PER1 expression 1 h after exposure to stress, during the light phase of the LD cycle, a time when levels of circulating glucocorticoids are known to be at their lowest [21].

**Mifepristone experiment.** Groups of rats (n = 4 per group) were treated with the GR antagonist, mifepristone, or vehicle, 1 h prior to the onset of the stressors, at ZT1. Rats were then exposed to either restraint or 2DG at ZT2, and subsequently treated at ZT3 as described above.

Results

**Adrenalectomy experiment**

As expected, plasma CORT levels were elevated in sham-operated rats 1 h following acute exposure to both the restraint and 2DG stressors, with significantly higher levels seen following restraint than 2DG (main effect of stress: $F_{2,18} = 37.48, p<0.0001$; Fig 1A). These effects were absent in ADX rats (main effect of treatment: $F_{1,18} = 66.8, p<0.0001$; interaction: $F_{2,18} = 28.69, p<0.0001$; Fig 1A). The effect of ADX on the stress-induced changes in PER1 expression is shown in Fig 1B–1G, Fig 2 and statistical analyses in Table 1. As can be seen in Fig 1, in sham-operated rats, restraint stress suppressed PER1 expression in both the BNSTov and CEAl (Fig 1B and 1C) and increased PER1 expression in the PVN, DMH, Pi (Fig 1D–1F). ADX reversed
the suppressive effect of restraint in the BNSTov and CEA1 (Fig 1B and 1C), whereas in contrast, the stimulatory effect of restraint stress on PER1 expression in the PVN, DMH and Pi was not significantly attenuated (Fig 1D–1F). Treatment with the systemic stressor 2DG increased PER1 levels in all regions studied including the BNSTov and CEA1, but not in the SCN.
(Fig 1B–1G). ADX reversed this increase in the CEA and DMH (Fig 1C and 1E). Contrastingly however, ADX significantly enhanced the stimulatory effect of 2DG on PER1 levels in the PVN (Fig 1D). The 2DG-induced increase in PER1 expression in the BNSTov and Pi was unaffected by ADX (Fig 1B and 1F). PER1 expression in the SCN was not affected by either stressor or by ADX (Fig 1G).

Mifepristone experiment

Plasma CORT levels in untreated rats increased following exposure to both restraint stress and 2DG (main effect of stress: \( F_{2,17} = 17.84, p<0.0001; \) Fig 3A). Treatment with the GR antagonist, mifepristone, had no effect on CORT levels in control group rats, but greatly enhanced the stress-induced circulating CORT levels (main effect of treatment: \( F_{1,17} = 34.75, p<0.0001; \)

Table 1. Statistical analysis (ANOVA) of the effect of adrenalectomy (ADX) and stress type (Stressor) on PER1 expression in limbic and hypothalamic brain regions.

| Structure | ADX | Stressor | ADX x Stressor |
|-----------|-----|----------|----------------|
| BNSTov    | \( F(1,18) = 2.26, \text{ ns} \) | \( F(2,18) = 10.75, p<0.001 \) | \( F(2,18) = 5.99, p<0.05 \) |
| CEA       | \( F(1,18) = 0.06, \text{ ns} \) | \( F(2,18) = 7.03, p<0.01 \) | \( F(2,18) = 13.78, p<0.001 \) |
| PVN       | \( F(1,18) = 6.41, p<0.05 \) | \( F(2,18) = 14.37, p<0.001 \) | \( F(2,18) = 1.57, \text{ ns} \) |
| DMH       | \( F(1,18) = 8.84, p<0.01 \) | \( F(2,18) = 5.0, p<0.05 \) | \( F(2,18) = 1.34, \text{ ns} \) |
| Pi        | \( F(1,18) = 0.24, \text{ ns} \) | \( F(2,18) = 18.03, p<0.0001 \) | \( F(2,18) = 1.33, \text{ ns} \) |
| SCN       | \( F(1,15) = 0.17, \text{ ns} \) | \( F(2,15) = 1.4, \text{ ns} \) | \( F(2,15) = 0.86, \text{ ns} \) |

\text{ns not significant}

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The effect of GR blockade on stress-induced PER1 expression is shown in Fig 3B–3G, Fig 4 and statistical analyses in Table 2. As with ADX, GR blockade prevented the restraint-induced suppression of PER1 expression in the BNSTov and CEA1, as well as the 2DG-induced increase in PER1 levels in these nuclei (Fig 3B and 3C). In addition, it attenuated the stimulatory effect of both stressors on PER1 levels in the DMH (Fig 3E). As with ADX, GR blockade accentuated the stress-induced increase in PER1 expression in
the PVN (Fig 3D). Mifepristone treatment had no effect on the restraint or 2DG-induced increase in PER1 expression in the Pi (Fig 3F). PER1 levels in the SCN of control and mifepristone-treated animals were again unaffected (Fig 3G).

**Discussion**

In mammals, the SCN clock governs the rhythmic secretion of glucocorticoid hormones [28]. In turn, glucocorticoids have been shown to play a role in the modulation of clock gene expression in subordinate clocks downstream of the SCN [7, 11, 29, 30]. In the present study we found, that depletion of endogenous glucocorticoids via ADX or treatment with the GR blocker, mifepristone altered the changes in PER1 expression induced by a neurogenic stressor, restraint, or by a systemic stressor, 2DG. These effects, however, were contingent upon the type of stress and brain region studied, where the central extended amygdala appears to be differentially sensitive to qualitative differences between systemic and processive stressors, as previously shown (21). This

**Table 2. Statistical analysis (ANOVA) of the effect of pharmacological blockade of glucocorticoid receptors (GRX) and stress type (Stressor) on PER1 expression in limbic and hypothalamic brain regions.**

| Structure | GRX         | Stressor  | GRX x Stressor |
|-----------|-------------|-----------|----------------|
| BNSTov    | $F_{(1,18)} = 0.22$, ns | $F_{(2,18)} = 7.79$, $p<0.01$ | $F_{(2,18)} = 10.65$, $p<0.001$ |
| CEA       | $F_{(1,18)} = 0.84$, ns | $F_{(2,18)} = 3.62$, $p<0.05$ | $F_{(2,18)} = 8.44$, $p<0.01$ |
| PVN       | $F_{(1,18)} = 24.78$, $p<0.0001$ | $F_{(2,18)} = 33.05$, $p<0.0001$ | $F_{(2,18)} = 2.26$, ns |
| DMH       | $F_{(1,17)} = 39.89$, $p<0.0001$ | $F_{(2,17)} = 7.55$, $p<0.01$ | $F_{(2,17)} = 8.7$, $p<0.01$ |
| Pi        | $F_{(1,18)} = 0.97$, ns | $F_{(2,18)} = 27.41$, $p<0.0001$ | $F_{(2,18)} = 1.95$, ns |
| SCN       | $F_{(1,17)} = 1.12$, ns | $F_{(2,17)} = 0.16$, ns | $F_{(2,17)} = 1.34$, ns |

*ns not significant*
points to a complex role for glucocorticoid signaling in stress-induced changes in PER1 expression in the brain.

The attenuation of endogenous glucocorticoid signaling, via ADX or GR blockade, reversed the suppressive effect of restraint and also the stimulatory effect of 2DG on PER1 expression in the BNSTov and CEAl, regions that we have previously shown to exhibit rhythms of PER2 expression that are uniquely dependent on circadian glucocorticoid signaling. Specifically, we found that PER2 rhythms in these two regions were blunted by ADX [13, 14] and could be restored by giving CORT via the drinking water, a regimen that mimics endogenous CORT rhythms [7, 15, 31]. Furthermore, we found that the genetic deletion of GR restricted to brain tissues of mice blunted PER2 rhythms in the BNSTov and CEAl without affecting rhythms in other brain regions, including the SCN [31]. Thus, the present findings show that PER expression in the BNSTov and CEAl is sensitive not only to daily rhythms of glucocorticoid signaling, but also to acute stress-induced changes in circulating levels of glucocorticoid hormones. Our current work examines PER1 levels at only 1 time point, in the context of stress-induced protein expression almost immediately post-stress exposure. This notion builds upon our previous work (21) that explores the pleiotropic role of PER1 in regulating endogenous circadian oscillations, as well as perhaps the integrative arm of the stress system with that of the circadian one. In order to further address the former role and capture any modulatory effect of changes in glucocorticoids on local oscillations, it would be wise to assess stress-induced PER1, as well as PER2, levels through several time points across a 24 h cycle.

The neuronal subpopulations in which these glucocorticoid-mediated changes in PER1 occur remain to be determined. It has been previously established that systemic stressors increase c-Fos expression in GABA neurons containing enkephalin in the BNSTov and CEAl [32, 33], and that neurogenic stressor, such as restraint, suppress c-Fos in these nuclei [34]. In addition, glucocorticoids have been shown to modulate the expression of enkephalin in these regions [35, 36], suggesting that these are the neurons where stress and glucocorticoids hormones could interact to modulate PER expression. The fact, however, that the two different stressors, restraint, a neurogenic stressor, and 2DG, a systemic stressor, exert opposite effects on PER1 and also c-Fos expression in both regions [21, 34] clearly implies that other factors are interacting with glucocorticoids to mediate the effects of these different stressors on PER1 expression in BNSTov and CEAl.

Contrary to the opposite effects of the two stressors in the BNSTov and CEAl, in the DMH, both restraint and 2DG increased PER1 expression and these effects were prevented by ADX and by GR blockade, pointing to a permissive role for glucocorticoids on PER1 expression in this region. Curiously, in the PVN, a nucleus that receives major afferent inputs from the DMH [37–39], both ADX and GR blockade enhanced the effect of the two stressors on PER1 expression, pointing to an inhibitory role for glucocorticoids on PER1 in this region. Similar inhibitory effects of glucocorticoids on neural activity and on expression of different peptides (corticotropin-releasing hormone, adrenocorticotropic hormone, vasopressin, glutamate, gamma-aminobutyric acid) in the PVN have been previously reported [40–43]. Neither ADX nor GR blockade had any effect on stress-induced PER1 expression in the Pi, indicating that the regulation of PER1 expression in this region by stress is independent of glucocorticoid signaling. Finally, the lack of effects of either stressor or of glucocorticoid manipulations on PER1 levels in the SCN was to be expected and is consistent with previous evidence that the adult SCN is devoid of GR [7, 44], and unlike its downstream oscillators, is immune to the effects of glucocorticoids [11]. Taken together, this set of findings highlights an indirect role of stress-induced changes in circulating glucocorticoids in the regulation of clock gene expression in downstream oscillators of the forebrain.
Blockade of GR mimicked the effect of ADX, underscoring the importance of glucocorticoid signaling via GR in stress-induced changes in PER1 expression in the forebrain. The transient suppression of GR expression within select brain regions of interest, through dsRNA interference, can be used to further validate our findings and evade any consequences of global GR antagonism. There is some evidence that a GRE in the promoter region of the Per1 gene mediates the effects of restraint stress on Per1 transcription in mouse peripheral organs including the liver, kidneys and heart [9]. It is not known whether the Per1 GRE is involved with any of the effects of stress on PER1 in the forebrain and, in particular, the suppressive effect of restraint in the BNSTov and CEAl. We found that acute changes in PER1 expression, resulting from a single stress exposure, were most robust in the short-term, 1 h post stress onset [21]. This suggests that stress-induced, glucocorticoid-dependent changes in brain PER1 expression may be mediated in parallel by rapid nongenomic mechanisms, perhaps via the activation of membrane-bound GR [41, 45, 46][41, 47].

In summary, the results of the present study show that acute exposure to categorically different stressors can induce rapid changes in PER1 expression, in a region-specific manner. Furthermore, these acute effects are attributed to changes in glucocorticoid signaling. It is possible that the stress-induced changes in PER1 can affect local clocks in the brain, leading to transient misalignment between the stress-resilient master clock and downstream brain oscillators, and ultimately to disruptions in circadian-controlled processes. Finally, it is important to note that PER1 has been implicated in behaviors such as stress-induced grooming, nociception, sensitization of drug effects, and anxiety-related behaviors in rodents [18, 20, 48–51], indicating that stress-induced changes in PER1 may have additional effects that are independent of the clock mechanism.

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Author Contributions

Conceived and designed the experiments: SAS SA. Performed the experiments: SAS MB SR. Analyzed the data: SAS. Wrote the paper: SAS SA.

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