Regulation of urocortin I and its related peptide urocortin II by inflammatory and oxidative stresses in HL-1 cardiomyocytes

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Abstract

Despite our knowledge on the regulation of urocortin (Ucn) I and its related peptides in the heart, the possible involvement of cardiovascular stress substances, such as cytokines or angiotensin II (Ang II), on this regulation remains to be fully elucidated. We therefore evaluated the potential role of cardiovascular stress substances on the regulation of the Ucn–corticotropin-releasing hormone (CRH) receptor system in HL-1 cardiomyocytes using a Ucn I-specific RIA, conventional reverse transcription-PCR (RT-PCR) and quantitative real-time RT-PCR. Ucn I mRNA levels were shown to be up-regulated by lipopolysaccarides (LPS), tumor necrosis factor-α (TNF-α), Ang II, H2O2, and pyrrolidinedithiocarbamate (PDTC). The LPS- and Ang II-induced increase in Ucn I mRNA levels was abolished by tempol. In addition, the secretion of Ucn I from HL-1 cardiomyocytes was stimulated by LPS and TNF-α.

On the contrary, Ucn II mRNA was increased by TNF-α alone and Ang II with tempol, and the TNF-α-induced increase in Ucn II mRNA was abolished by erythromycin and PDTC. These results suggested that Ucn I mRNA may be up-regulated by oxidative stress, whereas Ucn II mRNA may be up-regulated by the activated nuclear factor-κB, i.e. inflammatory stress. CRH-R2 mRNA may be negatively regulated by the increase in expression of Ucn I and/or Ucn II mRNA. In conclusion, the Ucn–CRH receptor system may be regulated by two major forms of cardiac stresses, i.e. oxidative and inflammatory stress, and may play a critical role in cardiac stress adaptation in heart diseases.

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Introduction

Urocortin (murine urocortin: Ucn, human urocortin: UCN) I, Ucn II, and Ucn III, the members of corticotropin-releasing hormone (CRH)-related peptides, have various actions on the cardiovascular system via cAMP-dependent pathways and the extra-cellularly regulated kinase 1/2 pathway (Ikeda et al. 1998, 2002, 2005a, Brar et al. 2004a, Honjo et al. 2006). We previously reported the pathophysiological involvement of UCN I in cardiomyopathies (Ikeda et al. 2003), in which a patient’s plasma concentration of tumor necrosis factor (TNF)-α increased (Matsumori et al. 1994). TNF-α can produce various deleterious effects when expressed in large amounts and it is produced in the heart by both cardiomyocytes and resident macrophages under conditions of cardiac stress (Sarzi-Puttini et al. 2005). In addition, Ucn I was up-regulated by TNF-α in human umbilical vein endothelial cells (Honjo et al. 2006), and, in turn, CRH type 2 receptor (CRH-R2) was down-regulated by lipopolysaccarides (LPS) and TNF-α in the mouse heart (Ikeda et al. 1998, Coste et al. 2001). Because CRH-R2 was also down-regulated by Ucn I in mouse cardiomyocytes (Heldwein et al. 1997, Coste et al. 2001) and in the mouse heart (Asaba et al. 2000), we hypothesize that LPS and TNF-α may stimulate the release of endogenous ligand(s) for CRH-R2, such as CRH, Ucn I, Ucn II, and/or Ucn III, from the heart. Also, we previously demonstrated that LPS and TNF-α stimulated the secretion of Ucn I, but not of CRH from neonatal rat cardiac myocytes and non-cardiac myocytes (Ikeda et al. 2001). These findings suggest that Ucn I and its related peptides may be regulated by inflammatory cytokines, such as LPS and TNF-α, and may play a pathophysiological role in the diseased heart because CRH-R2 signaling in the heart may be very important for adaptation to cardiac stress in pathological conditions (Coste et al. 2000). Ucn I may be regulated in response to oxidative stress, thereby exerting a vasoprotective role against the oxidative stress caused by angiotensin II (Ang II) in HUVECs (Honjo et al. 2006). We hypothesized that Ucn I and its...
related peptides may be regulated by inflammatory and/or oxidative stress(es) in the diseased heart. Therefore, the present study aims to evaluate the regulation of Ucn I and Ucn II in relation to inflammatory and oxidative stress.

Materials and methods

Cell culture

HL-1 cardiomyocytes, an immortalized cardiomyocyte cell line derived from mouse atria (Claycomb et al. 1998), were cultured as previously described (Ikeda et al. 2005b) with slight modification. Briefly, cells were plated in 10 cm culture dishes pre-coated with 0.02% Bacto gelatin (Becton, Dickinson, and Company, Franklin Lakes, NJ, USA) and 5 μg/ml fibronectin (Sigma–Aldrich Inc., St Louis, MO, USA) or in fibronectin-pre-coated 10 cm culture dishes (BD BioCoat, Becton, Dickinson, and Company) and cultured until cells reached confluence in Claycomb Medium (SAFC Biosciences, Lenexa, KS, USA) supplemented with 10% fetal bovine serum (FBS, Sigma–Aldrich Inc.), 0.1 mM norepinephrine (Sigma–Aldrich Inc.), 1% L-GlutaMAX (Invitrogen Corp.), and antibiotics (100 u/ml penicillin G and 10 μg/ml streptomycin). The cells were then removed by trypsinization and passed into another 10 cm culture dish, 6 cm culture dish, 12-well plate (3×10^5 cells/well), or a 6-well plate (6×10^5 cells/well) pre-coated with fibronectin/gelatin. Cells were cultured in FBS and norepinephrine-free Claycomb Medium supplemented with 1% L-GlutaMAX and antibiotics for 24 h prior to each experiment except when cells were used for conventional reverse transcription-PCR (RT-PCR).

cDNA preparation and conventional RT-PCR

Total RNA was extracted from HL-1 cardiomyocytes (at confluence in a 6 cm culture dish) by the acid–guanidium–phenol–chloroform (AGPC) method with 1ml/dish TRIzol Reagent (Invitrogen Corp.) after contamination of genomic DNA was ruled out by 50 cycle PCR with β-actin primers (shown in Table 1, estimated amplicon size: 510 bp). PCR was performed using Takara taq hot start version (Takara Bio Inc.) and primers specific for CRH-R1, CRH-R2 (Brar et al. 2004a), Ucn I, Ucn II (Li et al. 2003), and Ucn III (GenBank accession No. AF361944, designed by the Primer3 software (Rozen & Skaltsky 2000), estimated amplicon size: 254 bp). PCR was performed on cDNA from HL-1 cardiomyocytes, with conditions consisting of initial denaturation at 94 °C for 4 min, 5-step PCR cycles (CRH-R1, CRH-R2, Ucn I, and Ucn II, 35 cycles; Ucn III, 36 cycles; β-actin, 25 cycles: denaturation at 94 °C for 1 min, annealing at the temperatures shown in Table 1 for 1 min, and extension at 72 °C for 1 min), with a final extension at 72 °C for 7 min. As a positive control, cDNA from mouse brain (mouse brain QUICK-Clone cDNA, Clontech Laboratories, Inc., Mountain View, CA, USA) was used. In addition, expression of CRH-R1/R2 and Ucn mRNA in the mouse heart was also evaluated using 0.1 ng cDNA/reaction (Mouse Heart QUICK-Clone cDNA, Clontech Laboratories, Inc.). Primer sequences and annealing temperatures are summarized in Table 1.

Measurement of Ucn I levels secreted from HL-1 cardiomyocytes

To evaluate secretion of Ucn I from HL-1 cardiomyocytes, cells were cultured in 6-well plates and stimulated with LPS (1 ng/ml, Sigma–Aldrich Inc.), TNF-α (10 ng/ml, Sigma–Aldrich Inc.) or Ang II (10^{-7} mol/l, Peptide Institute, Inc., Osaka, Japan) for 3, 6,
12, and 24 h. The culture medium was collected and stored at −80 °C until measurement of Ucn I by Ucn I-specific RIA kit (Phoenix Pharmaceutical, Inc., Belmont, CA, USA). The total amount of protein from the HL-1 cardiomyocytes in each well was extracted by the M-PER Mammalian Protein Extraction Reagent (Pierce Biotechnology, Inc., Rockford, IL, USA) and measured using a bicinchoninic acid Protein Assay Reagent Kit (Pierce Biotechnology, Inc.) according to the manufacturer’s protocol (Smith et al. 1985). Secreted Ucn I was normalized against the total cellular protein of the HL-1 cardiomyocytes in each well. Data are presented as pmol/mg protein.

### Figure 1
Expression of CRH-R1, CRH-R2, Ucn I, Ucn II, Ucn III, and β-actin mRNA in HL-1 cardiomyocytes by RT-PCR. CRH-R2, Ucn I, and Ucn II mRNAs, but not CRH-R1 and Ucn III mRNAs, were expressed in HL-1 cardiomyocytes. HL-1 cardiomyocytes cDNA, MH, mouse heart QUICK-clone cDNA, MB, mouse brain QUICK-clone cDNA.

### Figure 2
The influence on the secretion of Ucn I from HL-1 cardiomyocytes by cytokines LPS (A) and TNF-α (B) and Ang II (C). Secretion of Ucn I was significantly increased by LPS, TNF-α, and Ang II. Data were represented as mean ± S.D. *P<0.01 versus vehicle, **P<0.05 versus vehicle.

### Table 2 Primers for quantitative real-time reverse transcription-PCR

<table>
<thead>
<tr>
<th>Primer sequences</th>
<th>CRH-R2 Forward</th>
<th>CRH-R2 Reverse</th>
<th>Ucn I Forward</th>
<th>Ucn I Reverse</th>
<th>Ucn II Forward</th>
<th>Ucn II Reverse</th>
<th>GAPDH Forward</th>
<th>GAPDH Reverse</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>5'-CTTCAACTCTTTCTGCACTTTC-3'</td>
<td>5'-GCTGTGGAAGCTGATCCTTG-3'</td>
<td>5'-CGCGGCTCTTTCTGCTGTTAG-3'</td>
<td>5'-ATGGACAGTGAGGGTGTC-3'</td>
<td>5'-CCCTTACCTAGACCCGTG-3'</td>
<td>5'-CGAGCCTGTTCCAGTAAGATCC-3'</td>
<td>5'-CAACCGGCTCATCATCTCCG-3'</td>
<td>5'-CTCGTGGTTCACCCCATCACCAC-3'</td>
</tr>
</tbody>
</table>
cardiomyocytes were then stimulated by LPS (0.01–10 ng/ml), TNF-α (0.1–100 ng/ml), and Ang II (10⁻⁹, 10⁻⁷, and 10⁻⁶ mol/l) for 3 h with or without a nuclear factor (NF)-κB inhibitor (erythromycin (10⁻⁵ mol/l, Sigma–Aldrich Inc.; Desaki et al. 2004) or ammonium pyrrolidinedithiocarbamate (PDTC, 10⁻⁴ mol/l, Sigma–Aldrich Inc.; Sherman et al. 1993, Boyd et al. 2006). To evaluate the involvement of oxidative stress on the expression of Ucn I and Ucn II mRNAs, we used H₂O₂ (Saito et al. 2001) at 0.5, 0.75 (Saito et al. 2001), and 1 mmol/l and the superoxide dismutase mimetic, 4-hydroxy-2, 2, 6, 6-tetramethylpiperidine-1-oxyl (tempol, 10⁻⁴ mol/l, Merck KGaA (Calbiochem), Darmstadt, Germany; Sakai et al. 2007). After washing twice with cold PBS, the cells were lysed by TRIzol reagent (1 ml/well) and stored at −80 °C until RNA extraction. Total RNA was extracted by the AGPC method and treated with DNase I until genomic DNA was no longer detected by 50 cycle PCR as described. Then cDNA was synthesized by primescript RT reagent kit (Takara Bio, Inc.) with oligo dT primers. Real-time PCR was performed on a Thermal Cycler Dice (Takara Bio, Inc) with SYBR premix Ex Taq/Taq II (Takara Bio, Inc.) according to the manufacturer’s protocols. The specific primers against mouse CRH-R2, Ucn I, Ucn II, and glyceraldehyde-3-phosphate-dehydrogenase (GAPDH) are summarized in Table 2. The real-time PCR was performed with conditions consisting of initial denaturation at 94 °C for 10 s, 2-step PCR cycles (50 cycles, denaturation at 95 °C for 5 s, annealing and detection step at 60 °C for 30 s), and the confirmation step of quality of PCR products by the dissociation curve (denaturation at 95 °C for 15 s followed by step-up elevation of temperature from 60 to 95 °C by 0.5 °C for every 30 s). Quantitative data were analyzed by the 2⁻ΔΔCt method (Livak & Schmittgen 2001), using the 2nd derivative curve of the amplification plots (Thermal Cycler Dice Real Time System TP800 software version 2.00B, Takara Bio, Inc.) and normalized against GAPDH cycle thresholds.

Statistical analysis

Statistical analysis was performed using ANOVA followed by a post hoc test for between-group comparison (StatView 5.0, SAS Institute, Inc., Cary, NC, USA).
Values $<0.05$ were considered to indicate statistical significance, and all data were expressed as the mean ± s.d.

Results

Expression of mouse Ucn I and its related peptides in HL-1 cardiomyocytes

RT-PCR revealed that CRH-R2, Ucn I, and Ucn II mRNA were expressed in both HL-1 cardiomyocytes and the mouse heart (Fig. 1). However, CRH-R1 and Ucn III mRNA were not expressed in this cardiomyocyte cell line and the mouse heart (Fig. 1).

Secretion of Ucn I by HL-1 cardiomyocytes stimulated with LPS, TNF-α, and Ang II

Ucn I-like immunoreactivity (Ucn I-LI) increased in the culture medium of HL-1 cardiomyocytes in a time-dependent manner. The secretion of Ucn I, as detected by RIA, significantly increased in response to LPS (3, 6, 12 h: $P<0.01$ versus vehicle, Fig. 2A) and to TNF-α (24 h: $P<0.05$ versus vehicle, Fig. 2B). In addition, secretion of Ucn I-LI from HL-1 cardiomyocytes was also significantly increased by Ang II (3, 6, 12 h: $P<0.01$ versus vehicle, Fig. 2C).

Real-time RT-PCR detection of mouse Ucn I, Ucn II, and CRH-R2 mRNA

Real-time RT-PCR demonstrated that both LPS and TNF-α significantly increased Ucn I mRNA expression at 3 h (Fig. 3A and B). At 3 h, expression of Ucn II mRNA was significantly increased by TNF-α, but not by LPS (Fig. 4A). Expression of Ucn I mRNA increased in a dose-dependent manner in response to LPS at 3 h (Fig. 2B). On the contrary, expression of Ucn II mRNA was not significantly affected by LPS at 3 h (Fig. 4A). TNF-α, on the other hand, increased expression of both Ucn I and Ucn II mRNA dose dependently (Fig. 5B and F). Furthermore, Ang II and H2O2 significantly stimulate the expression of Ucn I mRNA compared with vehicle at 3 h (Fig. 3C and D). This increase in Ucn I mRNA occurs in a dose-dependent manner at 3 h (Fig. 5C and D). But Ang II suppressed the expression of Ucn I mRNA at 6 and 12 h (Fig. 3C). On the contrary, Ang II did not affect expression of Ucn II mRNA (Fig. 4B). Expression of CRH-R2 mRNA seemed to be decreased by the combined effects of the Ucn I and Ucn II induced by LPS and TNF-α at 3 and 12 h (Fig. 5B and F). Erythromycin ameliorated the LPS-induced increase in Ucn I mRNA. However, erythromycin did not affect the TNF-α-induced increase in Ucn I mRNA (Fig. 7A and B). Erythromycin and PDTC seemed to be additive to that of LPS but the difference did not reach statistical significance (Fig. 7C). The TNF-α-induced increase in Ucn II mRNA was abolished by erythromycin and PDTC (Fig. 7E). Furthermore, the Ang II-induced increase in Ucn II mRNA was abolished by tempol, whereas Ucn II mRNA was inversely regulated by Ang II and tempol (Fig. 8C and E). In addition, tempol abolished the PDTC-induced increase in Ucn I mRNA (Fig. 8D).

Discussion

In the present study, we demonstrate that HL-1 cardiomyocytes express Ucn I, Ucn II, and CRH-R2, indicating that this cardiomyocyte cell line is a useful
model for investigating the Ucn–CRH receptor system (Ikeda et al. 2005a). Although CRH-R1 is involved in anxiety and neuroendocrine stress and CRH-R1 and Ucn III were reported to be expressed in the human heart (Kimura et al. 2002, Takahashi et al. 2004), neither CRH-R1 nor Ucn III mRNA was expressed in the HL-1 mouse atrial cardiomyocyte cell line. These data are supported by previous studies using rat and mouse cardiomyocytes (Brar et al. 2004b, Ikeda et al. 2005a).

Therefore, Ucns act through CRH-R2, the main mediator of the stress response in cardiomyocytes (Coste et al. 2000, Pañeda et al. 2005), rather than through CRH-R1. This suggests that the autocrine stress adaptation system is mediated by endogenous Ucn I, Ucn II, and CRH-R2 in HL-1 cardiomyocytes and by exogenous CRH-R2 ligands.

Previous studies demonstrated an increase in the expression of Ucn I in target regions of inflammation in diseases involving TNF-α (Uzuki et al. 2001, Ikeda et al. 2003, Saruta et al. 2004). In addition, a recent study showed that the expression of Ucn II mRNA was increased in mucosal samples of patients with inflammatory bowel disease, and in human intestinal xenografts following exposure to Clostridium difficile toxin A (Moss et al. 2007), an activator of NF-κB (Jefferson et al. 1999), indicating that the regulation of Ucn II may be involved in the NF-κB pathway. Our present results showed that LPS, Ang II, and H₂O₂
individually enhanced the expression of Ucn I mRNA in as well as the secretion of Ucn I from HL-1 cardiomyocytes, and that these effects were abolished by tempol. This increase in Ucn I mRNA by LPS and Ang II was abolished by antioxidants such as erythromycin (Miyachi et al. 1986) and tempol. Although HL-1 cardiomyocytes expressed toll-like receptor (TLR; Boyd et al. 2006), the present results showed that the actions of LPS may not be the action via the TLR signaling because of the absence of LPS-binding protein, which enhances biological action of LPS (Hailman et al. 1994), in stimulation medium and abolishment of the action of LPS on expression of Ucn I mRNA by antioxidant. These results, taken in combination with the demonstration that LPS and Ang II cause oxidative stress (Honjo et al. 2006, Yuan et al. 2009), strongly suggest that Ucn I may be positively regulated by oxidative stress. Ang II markedly increased the secretion of Ucn I up to 12 h, with a temporal increase in the expression of Ucn I mRNA followed by

![Figure 6](https://www.endocrinology-journals.org)
negatively regulate each other. Taken together, the present results suggest that Ucn I mRNA and Ucn II mRNA are differentially regulated, i.e. NF-κB may, at least in part, be involved in the regulation of Ucn II mRNA, while oxidative stress may up-regulate Ucn I mRNA in HL-1 cardiomyocytes. These data also suggest that Ucn I and Ucn II may play a pathophysiological role in cardiac disease, especially in heart failure, in which the levels of vasoactive substances such as cytokines and Ang II (Matsumori et al. 1994, Ikeda et al. 2008) are increased.

In conclusion, the present results demonstrate that HL-1 cardiomyocytes contain the Ucn–CRH receptor system, which may be regulated by two major forms of cardiac stress, oxidative and inflammatory stress. In heart disease, such differential regulation of Ucn I and Ucn II could play a critical role in stress adaptation via the Ucn–CRH receptor system.
Declaration of interest

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

Funding

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References


Figure 8 The effects of tempol on the regulation of Ucn I mRNA induced by LPS (A), TNF-α (B), Ang II (C), and ammonium pyrrolidinedithiocarbamate (PDTC) (D) and on Ucn II mRNA induced by Ang II (E) at 3 h. Tempol abolished the LPS- and Ang II-induced expression of Ucn I mRNA, indicating that Ucn I mRNA may be dominantly regulated by oxidative stress. On the contrary, co-incubation with Ang II and tempol resulted in an increase in Ucn II mRNA expression. Data are represented as the mean ± S.D. LPS, LPS 1 ng/ml, TNF, TNF-α 10 ng/ml, Ang II, Ang II 10⁻⁷ mol/l, PDTC, PDTC 10⁻⁴ mol/l, Tem, Tempol 10⁻⁴ mol/l, (A) *P<0.01 versus vehicle, **P<0.01 versus LPS, (B) *P<0.01 versus vehicle, **P<0.05 versus vehicle, #P<0.05 versus TNF, $P<0.01 versus vehicle, (C) *P<0.05 versus vehicle, **P<0.05 versus Ang II, (D) *P<0.05 versus vehicle, **P<0.05 versus PDTC, (E) *P<0.01 versus vehicle, **P<0.01 versus Ang II, #P<0.05 versus vehicle.


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