

RESEARCH PAPER

Urocortin induced expression of COX-2 and ICAM-1 via corticotrophin-releasing factor type 2 receptor in rat aortic endothelial cells

Rongjian Zhang*, Youhua Xu*, Hong Fu, Juejin Wang, Lai Jin and Shengnan Li

Department of Pharmacology, Nanjing Medical University, Nanjing, China

Background and purpose: Our previous study showed that urocortin (Ucn1) exacerbates the hypercoagulable state and vasculitis in a rat model of sodium laurate-induced thromboangiitis obliterans. Furthermore, the inflammatory molecules COX-2 and ICAM-1 may participate in this effect. In the present study, the effects of Ucn1 on COX-2 and ICAM-1 expression in lipopolysaccharide (LPS)-induced rat aortic endothelial cells (RAECs) were investigated and the mechanisms involved explored.

Experimental approach: RAECs were isolated from adult male Wistar rats, and identified at the first passage. Experiments were performed on cells, from primary culture, at passages 5–8. The expression of COX-2 and ICAM-1 at both mRNA and protein levels was determined by semi-quantitative RT-PCR and Western blot analysis. Levels of PGE₂ and soluble ICAM-1 (sICAM-1) in culture medium were measured by enzyme-linked immunosorbent assay. Furthermore, the phosphorylation status of p38MAPK, ERK1/2, JNK, Akt and NF-κB was analysed by Western blot; nuclear translocation of NF-κB was observed by immunofluorescence.

Key results: Ucn1 augmented LPS-induced expression of COX-2 and ICAM-1 in RAECs in a time- and concentration-dependent manner. Ucn1 increased PGE₂ and sICAM-1 levels. These effects were abolished by the CRF₂ receptor antagonist, antisauvagine-30, but not by the CRF₁ receptor antagonist, NBI-27914. Moreover, Ucn2 activated p38MAPK and augmented NF-κB nuclear translocation and phosphorylation, whereas ERK1/2, JNK and Akt pathways were not involved in this process.

Conclusions and implications: These findings suggest that Ucn1 exerts pro-inflammatory effects by augmenting LPS-induced expression of COX-2 and ICAM-1 in RAECs via CRF₂ receptors and the activation of p38MAPK and NF-κB.

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Abbreviations: CRF, corticotrophin-releasing factor; CRF₁ receptor, corticotrophin-releasing factor type 1 receptor; CRF₂ receptor, corticotrophin-releasing factor type 2 receptor; ECs, endothelial cells; RAECs, rat aortic endothelial cells; TAO, thromboangiitis obliterans; Ucn, urocortin

Introduction

Vascular endothelial cells (ECs) play a pivotal role in modulating local and systemic inflammation. ECs express chemokines that initiate the activation and recruitment of circulating leukocytes at sites of tissue inflammation (Berman *et al.*, 1993; Bannerman and Goldblum, 1997). Among these chemokines,

COX-2 and ICAM-1 are two important modulators, with the former participating in a series of inflammatory diseases and the latter playing an important role in the adhesion process of inflammatory cells. The bacterial endotoxin, lipopolysaccharide (LPS), an essential component of the surface of Gram-negative bacteria (Raetz, 1990), has potent pro-inflammatory properties by acting on many cell types including ECs (Berman *et al.*, 1993; Munshi *et al.*, 2002). LPS-activated ECs may be changed with up-regulation of COX-2 and ICAM-1 expression (Chen *et al.*, 2001b; Heo *et al.*, 2008), pro-coagulant activity, enhanced endothelial permeability and abundant pro-inflammatory mediators' secretion (Berman *et al.*, 1993; Bannerman and Goldblum, 1997; Bierhaus *et al.*

Correspondence: Shengnan Li, Department of Pharmacology, Nanjing Medical University, 140 Hanzhong Road, Nanjing 210029, China. E-mail: snli@njmu.edu.cn

*These authors contributed equally to this work.

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2000), which may lead to an imbalance in the immune system. Thus, finding a way to minimize COX-2 and ICAM-1 expression may have a dramatic impact on the treatment of inflammatory diseases.

Urocortin (Ucn), a 40 aa corticotrophin-releasing factor (CRF) family peptide that was first identified in rat midbrain (Vaughan *et al.*, 1995), has been demonstrated to be widely expressed in peripheral tissues including the cardiovascular system, spleen, skin, lymphocytes, macrophages and mast cells (Fekete and Zorrilla, 2007). The effects of Ucn1 are mediated by two high-affinity receptors, the CRF₁ receptor and the CRF₂ receptor, with the former mainly distributed centrally and the latter primarily expressed peripherally (Fekete and Zorrilla, 2007). Besides its cardiovascular protective property (Okosi *et al.*, 1998; Oki and Sasano, 2004; Yang *et al.*, 2006), Ucn1 is now considered to be a potent immunomodulator which participates in various immune responses (Fekete and Zorrilla, 2007). Our previous studies have demonstrated that Ucn1 participates in the development of allergic asthma in the rat and can trigger mast cell infiltration and activation via CRF receptors (Wu *et al.*, 2006; 2008). Singh *et al.* (1999) found that Ucn1 is potent at inducing mast cell degranulation and triggering vascular permeability via CRF receptors. Moreover, our data showed that CRF plays a significant role in promoting the development of atherosclerosis (Wu *et al.*, 2009). All of these findings suggest that immune-derived Ucn1 may participate in the pathophysiology of many inflammatory conditions as a pro-inflammatory mediator.

Previous studies have indicated that endogenous Ucn1 has an inhibitory role in the growth of ECs and blood vessels (Yang *et al.*, 2008), and it may trigger ECs to change into a procoagulant state (Grignani and Maiolo, 2000) by stimulating interleukin-1b (IL-1b) and interleukin-6 (IL-6) secretion *in vitro* (Kohno *et al.*, 2001). Recently, it was found that activation of CRF₂ receptors suppresses vascularization (Bale *et al.*, 2002; Hao *et al.*, 2008). Thus, Ucn1 may have a dramatic impact on the normal function of ECs.

Kageyama *et al.* (2006) demonstrated that in rat aortic smooth muscle cells, Ucn1 induces the expression of COX-2 in a time- and concentration-dependent manner (Kageyama *et al.*, 2006). Also, CRF has been shown to enhance the interferon- γ -stimulated expression of ICAM-1 on human skin keratinocytes (Quevedo *et al.*, 2001). Our recent data demonstrated that Ucn1 promotes the development of vasculitis in the rat thromboangiitis obliterans (TAO) model (original article accepted by *British Journal of Pharmacology*). Because COX-2 and ICAM-1 are associated with the development of vasculitis (de Gaetano *et al.*, 2003; Witkowska, 2005), our data also suggest that this effect of Ucn1 on vasculitis might be exerted through an effect on the expression of these two factors. The present study was performed to examine the effects of Ucn1 on COX-2 and ICAM-1 expression in LPS-induced rat aortic endothelial cells (RAECs) and explore the mechanisms involved. In the present study, it was first demonstrated that Ucn increased the expression of both COX-2 and ICAM-1 in a time- and concentration-dependent manner in LPS-activated RAECs. Furthermore, the p38MAPK and NF- κ B pathways were involved in this effect, which was exerted via the CRF₂ receptor.

Methods

Culture of RAECs

RAECs were isolated from male Wistar rats (Shanghai Laboratory Animal Center, Shanghai, China) according to the method described previously (McGuire and Orkin, 1987; Yang *et al.*, 2006). The animal operation procedure was approved by the ethics review board of Nanjing Medical University. Cells were cultured in RPMI 1640 supplemented with 20% fetal bovine serum, 1% penicillin-streptomycin, at 37°C in a 95% air/5% CO₂ incubator. Our previous studies were carried out with passages 5–8 RAECs, and it was found, using an F-VIII marker, that these cells maintain their properties (Yang *et al.*, 2006). Thus, the present experiments were performed on cells, from primary culture, at passages 5–8. The identity of the RAECs was confirmed by immunofluorescence staining by the use of rabbit anti-rat factor VIII antibody and Cy-3 conjugated goat anti-rabbit IgG.

Immunofluorescence staining

RAECs were cultured on coverslips placed in tissue culture dishes. Following different treatments, the cells were washed with phosphate-buffered solution and fixed with fresh 4% paraformaldehyde for 30 min. Subsequently, the cells were permeabilized with 0.5% Triton X-100 for 15 min on ice and blocked in 5% BSA for 30 min at room temperature. After being blocked, the cells were incubated with factor VIII antibody (1:100) or NF- κ B p65 antibody (1:25) overnight in a humid chamber at 4°C. Then, the cells were incubated with a secondary antibody conjugated to Cy-3 for 30 min in the dark. After being washed three times, the cells were mounted on a slide. The slides were visualized using a fluorescence microscope.

RNA isolation and semi-quantitative RT-PCR analysis

Total RNAs were extracted from RAECs, using TRIzol according to the manufacturer's protocol. For cDNA synthesis, Moloney murine leukaemia virus (MMLV) was applied as the reverse transcriptase. For PCR reaction, Taq DNA polymerase was used in the reaction system. Primers for COX-2 (Ohnaka *et al.*, 2000), ICAM-1 (Taal *et al.*, 2000) and GAPDH (Baigent and Lowry, 2000) were synthesized from published sequences as shown in Table 1. The products were visualized by electrophoresis in 2.0% agarose gel containing 0.5 μ g·mL⁻¹ ethidium bromide. Specific genes were verified by their predicted sizes. GAPDH was set as the internal control.

Western blot analysis

The protein samples were separated on a 10% sodium dodecyl sulphate-polyacrylamide gel and electrophoretically transferred to PVDF membranes in Tris-glycine transfer buffer. Then, membranes were blocked in 5% (w/v) instant non-fat dried milk for 2 h at room temperature, and incubated with primary antibodies corresponding to COX-2 (1:1000), ICAM-1 (1:500), β -actin (1:250), p38MAPK (1:1000), phospho-p38MAPK (1:1000), ERK1/2 (1:1000), phospho-

Table 1 A summary of the RT-PCR primer sequences used to amplify GAPDH, COX-2 and ICAM-1

		Sequences	Product size (bp)	Annealing T (C)
GAPDH	Sense	TCCCAGAGCTGAACGGGAAGCTCACTG	339	68.1
	Antisense	TGGAGGCCATGTAGGCCATGAGGTCCA		
COX-2	Sense	TTCACCAGACAGATTGCTGGC	530	63.5
	Antisense	AGTCTGGAGTGGGAGGCACTTG		
ICAM-1	Sense	AGAAGGACTGCTTGGGGAA	332	58.1
	Antisense	CCTCTGGCGGTAATAGGTG		

ERK1/2 (1:1000), JNK (1:1000), phospho-JNK (1:1000), Akt (1:1000), phospho-Akt (Ser 473) (1:1000), NF- κ B p65 (1:1000), phospho-NF- κ B p65 (1:1000), respectively, at 4°C overnight. The membranes were subsequently washed with TBST [50 mM Tris-HCl (pH 7.5), 150 mM NaCl, 0.05% Tween 20] and incubated with secondary horseradish peroxidase-conjugated IgG for 2 h at room temperature. Immunoreactive proteins were visualized by LumiGLO (Cell Signaling Technology, Beverly, MA, USA) chemiluminescent reagent and peroxide. The light-emitting bands were detected with X-ray films.

Enzyme-linked immunosorbent assay (ELISA)

Levels of PGE₂ and soluble ICAM-1 (sICAM-1) in culture medium were determined by ELISA assay using PGE₂ and sICAM-1 kits according to the manufacturers' instructions. Kit standards and controls were performed for each assay completed. The final concentrations of PGE₂ and sICAM-1 were calculated by converting the optical density reading using standard curves.

Statistical analysis

Data are expressed as means \pm SEM. The significance for the difference among groups was analysed with SPSS 11.0 (Chicago, IL, USA) by one-way analysis of variance with Student-Newman-Keuls multiple comparison methods. Differences were considered to be statistically significant at a *P* value of <0.05.

Reagents

Rat urocortin1 (Ucn1), Ucn2, CRF₁ receptor antagonist NBI-27914, CRF₂ receptor antagonist antisauvagine-30, LPS, rabbit anti-rat factor VIII (a stable endothelial antigen) antibody and Cy-3 conjugated goat anti-rabbit IgG were purchased from Sigma (St. Louis, MO, USA). Polyclonal COX-2 antibody and monoclonal ICAM-1 antibody were obtained from Abcam (Cambridge, UK). PGE₂ and sICAM-1 ELISA kits were obtained from R&D (Minneapolis, MN, USA) and Boster (Wuhan, China) respectively. Specific antibodies to p38MAPK, phospho-p38MAPK, ERK1/2, phospho-ERK1/2, JNK, phospho-JNK, Akt, phospho-Akt (Ser 473), NF- κ B p65, phospho-NF- κ B p65, p38MAPK inhibitor SB203580, LumiGLO chemiluminescent reagent and peroxide were provided by Cell Signaling Technology. TRIzol and MMLV

were obtained from Invitrogen (Carlsbad, CA, USA), Taq DNA polymerase from Promega (Madison, WI, USA) and the X-ray films from Kodak (Rochester, NY, USA). The other reagents used were derived from commercial sources. All drug/molecular target nomenclature conforms to *British Journal of Pharmacology's* Guide to Receptors and Channels (Alexander *et al.*, 2008).

Results

Immunofluorescent identification of RAECs

After being isolated, RAECs were identified at the first passage. As shown in Figure 1, the cytoplasm was completely stained red with factor VIII-specific antibody and Cy-3 conjugated secondary antibody. Factor VIII-associated antigen is a cytological marker closely identified with ECs (Karasek, 1989); the cells we obtained were undoubtedly RAECs.

Ucn augmented LPS-induced COX-2 and ICAM-1 expression in RAECs

As depicted in Figure 2, after exposure of RAECs to 10 μ g·mL⁻¹ LPS, 10⁻⁷ M Ucn (Ucn1) induced COX-2 and ICAM-1 expression in a time-dependent manner, with the mRNA and protein levels reaching their peaks after 4 and 8 h respectively. With increased time, their expression was decreased. Hence, in the following experiments, the time points of 4 and 8 h were selected to determine the mRNA and protein levels respectively. Ucn1 augmented LPS-induced COX-2 and ICAM-1 expression in a concentration-dependent manner; a marked effect was observed with 10⁻⁹ M Ucn1, which did not increase at higher concentrations (Figure 3).

COX-2 and ICAM-1 expression was mediated by the CRF₂ receptor

To illustrate the involvement of CRF receptors in Ucn1-induced augmentation of COX-2 and ICAM-1 expression in LPS-activated RAECs, cells were pretreated with Ucn1 alone or together with CRF receptor antagonists (CRF₁ receptor antagonist, NBI-27914, or CRF₂ receptor antagonist, antisauvagine-30). Ucn1 pretreatment augmented LPS-induced COX-2 and ICAM-1 mRNA expression to levels 1.36- and 1.40-fold observed after LPS pretreatment alone, respectively; antisauvagine-30 reversed this augmentation (Figure 4A,B). A similar result was observed on COX-2 and ICAM-1 protein

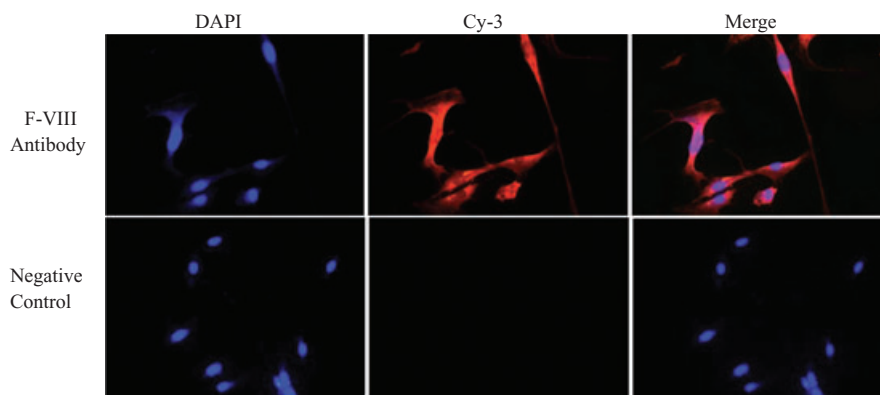


Figure 1 Immunofluorescent identification of rat aortic endothelial cells. Cells cultured on coverslips were incubated with rabbit anti-rat factor VIII (a stable endothelial antigen) antibody and then incubated with Cy-3 conjugated secondary antibody; the nuclei were stained with DAPI. In the negative control, factor VIII antibody was omitted (magnification: 200 \times).

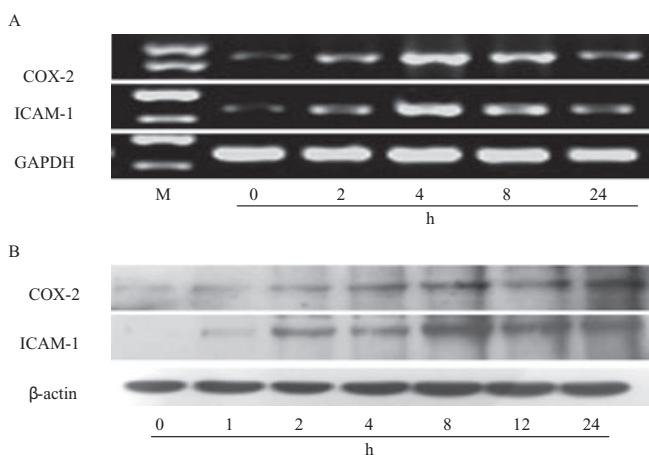


Figure 2 COX-2 and ICAM-1 expression induced by Ucn1 occurred in a time-dependent manner in lipopolysaccharide (LPS)-activated rat aortic endothelial cells. In the presence of LPS ($10 \mu\text{g}\cdot\text{mL}^{-1}$), cells were pretreated with Ucn1 (10^{-7} M) and incubated for the durations shown. (A) COX-2 and ICAM-1 mRNA expression induced by Ucn1 reached their peak at the time point of 4 h. (B) Ucn1 promoted COX-2 and ICAM-1 protein expression with significant effects at the time point of 8 h. Similar results were obtained from more than three independent cultures, and a representative experiment is shown.

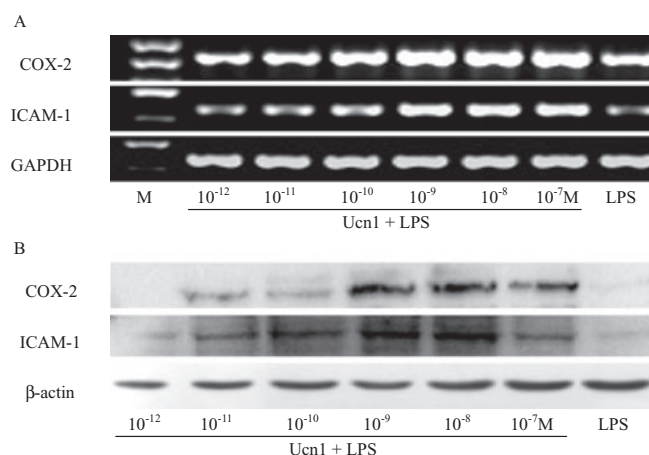


Figure 3 COX-2 and ICAM-1 expression induced by Ucn1 was in a concentration-dependent manner in lipopolysaccharide (LPS)-activated rat aortic endothelial cells. On exposure to LPS ($10 \mu\text{g}\cdot\text{mL}^{-1}$), cells were pretreated with Ucn1 at concentrations ranging from 10^{-12} to 10^{-7} M for 30 min. (A) Cells were incubated for 4 h, and COX-2 and ICAM-1 mRNA expression was determined by RT-PCR. (B) Cells were incubated for 8 h, and COX-2 and ICAM-1 protein level was measured by Western blot. Experiments were done from more than three independent cultures, and a representative experiment is shown.

expression (Figure 4C,D). However, NBI-27914 had no significant effect on COX-2 and ICAM-1 expression.

The concentrations of PGE₂, the main metabolite of COX-2, and sICAM-1 in the supernatant of the culture were measured by ELISA. Similar results were observed for both PGE₂ and sICAM-1 in that levels were increased by the application of Ucn1, and pretreatment with antisauvagine-30 abolished this increase (Figure 4IIA,B).

Interestingly, it was found that the two molecules, COX-2 and ICAM-1, changed in a similar manner. To explore their possible relationship, NS-398, a selective inhibitor of COX-2, was used. As shown in Figure 4IIA, in the presence of both Ucn1 and LPS, PGE₂ production was significantly decreased after NS-398 pretreatment. Furthermore, Ucn1-induced elevation of ICAM-1 was dramatically reduced on blockade of COX-2 in LPS-activated RAECs (Figure 4IIB). These

observations indicate that Ucn1-induced ICAM-1 expression was partially mediated by COX-2.

Ucn2-induced p38MAPK and NF- κ B phosphorylation in LPS-activated RAECs

Mitogen-activated protein kinases (MAPKs) and Akt pathways have been implicated in the activation of the pro-inflammatory process (Bhat *et al.*, 1998; Ojaniemi *et al.*, 2003), and have been shown to be key signalling intermediates downstream of CRF₂ receptor activation (Dermizaki *et al.*, 2002; Sananbenesi *et al.*, 2003; Karteris *et al.*, 2004; Moss *et al.*, 2007; Markovic *et al.*, 2008). To investigate the role of these two pathways in the effect of CRF₂ receptor on LPS-induced COX-2 and ICAM-1 expression, the cells were pretreated with Ucn2, a selective CRF₂ receptor agonist

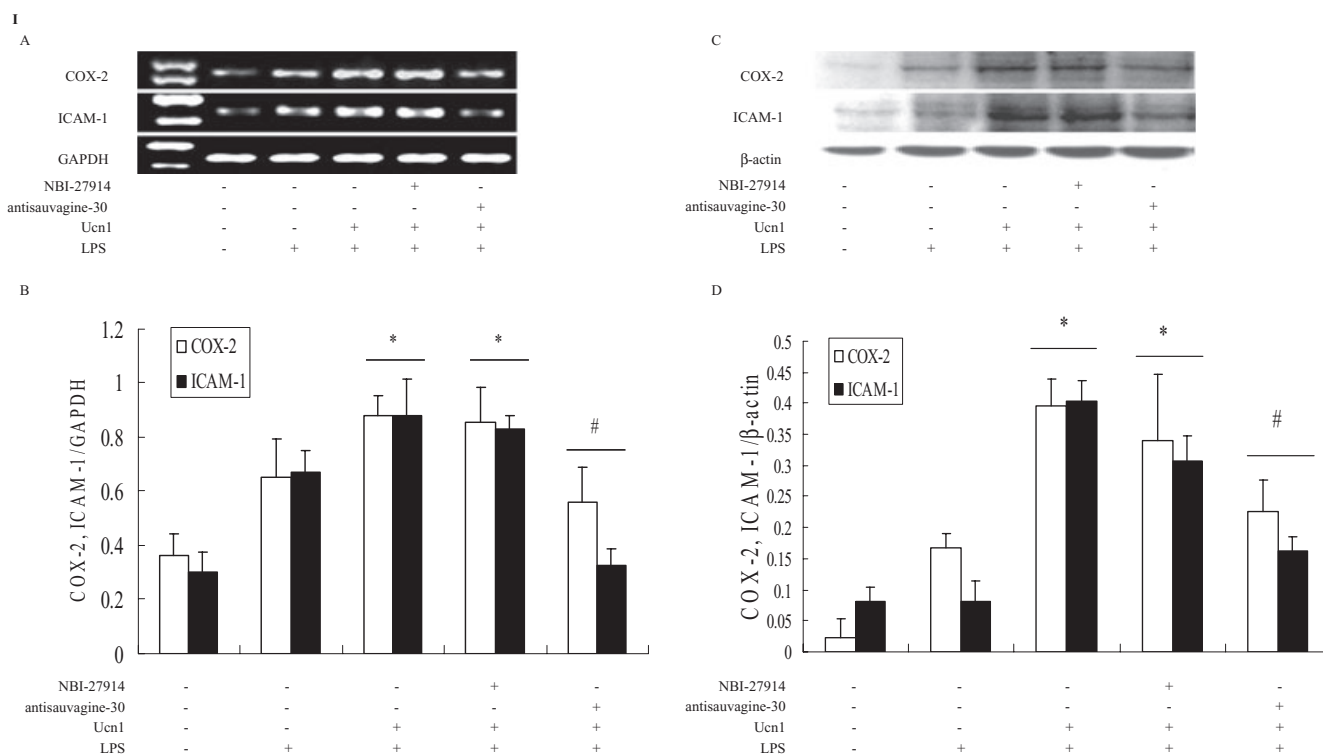


Figure 4 Effect of Ucn1 on COX-2 and ICAM-1 expression in rat aortic endothelial cells. Cells were treated with vehicle, Ucn1 (10^{-9} M) alone or together with NBI-27914 (10^{-8} M) or antisauvagine-30 (10^{-8} M), or NS-398 (10^{-5} M) in the presence of lipopolysaccharide (LPS) ($10 \mu\text{g}\cdot\text{mL}^{-1}$) as indicated. (I) Effect of Ucn1 on COX-2 and ICAM-1 expression. Cells were incubated for 4 h (mRNA) or 8 h (protein) to determine COX-2 and ICAM-1 levels by RT-PCR (IA and B) or Western blot (IC and D). (II) Effect of Ucn1 on PGE₂ and soluble ICAM-1 (sICAM-1) production measured by enzyme-linked immunosorbent assay. Cells were incubated for 24 h, and the culture supernatant was obtained to determine PGE₂ (IIA) or sICAM-1 (IIB) levels. Data given are the means \pm SEM of values taken from three independent cultures. * $P < 0.05$, versus LPS group; # $P < 0.05$, versus Ucn1 + LPS group. Similar results were obtained from more than three independent cultures, and a representative experiment is shown.

(Fekete and Zorrilla, 2007). Subsequently, the phosphorylation status of p38MAPK, ERK1/2, JNK and Akt were analysed by Western blot.

As depicted in Figure 5A, in the presence of $10 \mu\text{g}\cdot\text{mL}^{-1}$ LPS, 10^{-9} M Ucn2 induced a transient phosphorylation of p38MAPK with peak activation at 15 min. Furthermore, LPS-induced significant phosphorylation of p38MAPK and Ucn2 application dramatically augmented this phosphorylation (Figure 5B,C). This phosphorylation of p38MAPK was completely blocked in the presence of 10^{-5} M SB203580, a p38MAPK inhibitor. However, the phosphorylation of ERK1/2, JNK and Akt did not appear to be altered by Ucn2 treatment (Figure 5D–F).

NF- κ B is known to participate in the inflammatory process (Ghosh and Hayden, 2008). To determine whether it is involved in Ucn2-induced inflammation, nuclear translocation of NF- κ B was observed by immunofluorescence. As shown in Figure 6A, NF- κ B was visualized primarily in the cytoplasm in untreated control cells, while LPS treatment for 1 h induced NF- κ B translocation from the cytoplasm into the nucleus. What is more important, compared with LPS treatment, Ucn2 noticeably augmented NF- κ B nuclear translocation as after Ucn2 treatment, most NF- κ B was visualized in the nucleus of the cells. Furthermore, the phosphorylation status of NF- κ B was determined by Western blot analysis. As shown in Figure 6B,C, LPS pretreatment significantly increased

NF- κ B phosphorylation; Ucn2 application further enhanced the phosphorylation of NF- κ B. Combined with the observation in Figure 6A, we came to the conclusion that Ucn2 can increase LPS-induced phosphorylation of NF- κ B in RAECs.

Discussion and conclusions

Ucn (Ucn1), a CRF family peptide (Vaughan *et al.*, 1995), has been demonstrated to stimulate the release of pro-inflammatory mediators under inflammatory conditions (Kohno *et al.*, 2001; Saruta *et al.*, 2004). In this study, we found that Ucn1 could increase LPS-induced COX-2 and ICAM-1 expression in RAECs via the CRF₂ receptor. Furthermore, p38MAPK and NF- κ B pathways participated in this process.

It has been well established that, in addition to the indirect anti-inflammatory effect via ACTH/cortisol (Elenkov and Chrousos, 1999; Elenkov *et al.*, 1999), the CRF family peptides play a direct pro-inflammatory role in the regulation of the inflammatory process (Elenkov and Chrousos, 1999; Elenkov *et al.*, 1999). Moreover, the pro-inflammatory effect of CRF family peptides is partially attributed to their stimulation of immune cells, such as mast cells and macrophages (Theoharides *et al.*, 1995; 1998; Agelaki *et al.*, 2002; Tsatsanis *et al.*, 2007). It has been demonstrated that these immune cells

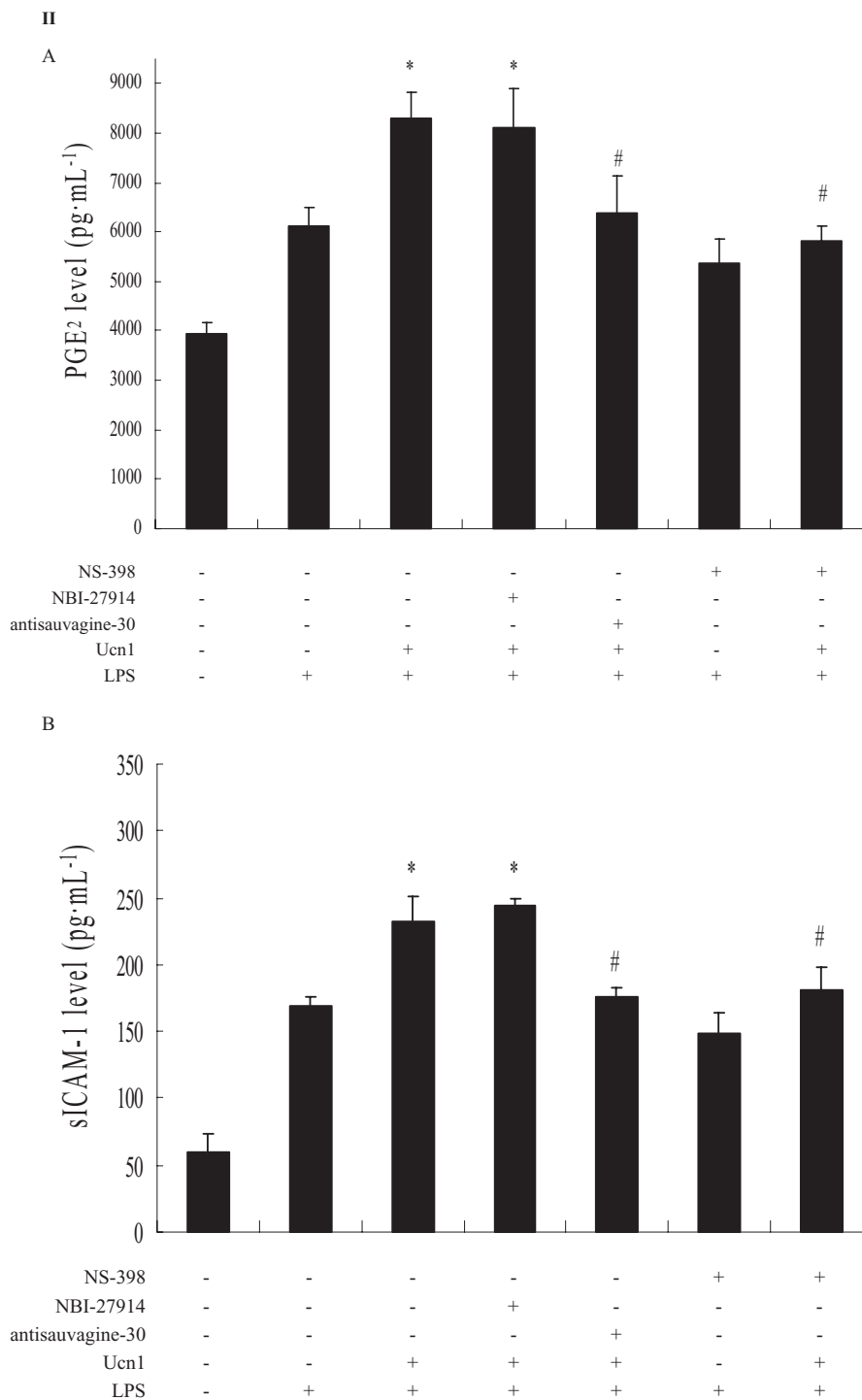


Figure 4 Continued

express CRF₁ and CRF₂ receptors, and can synthesize CRF family peptides (Theoharides *et al.*, 1998; Baigent, 2001). CRF and Ucn1 stimulate the production and release of pro-inflammatory mediators in mast cells (Theoharides *et al.*, 1995; 1998), such as IL-1, IL-6 and TNF- α . Moreover, CRF intensifies the response of macrophages to bacterial LPS by augmenting their synthesis of the pro-inflammatory cytokines, TNF- α and IL-6, at the mRNA level (Agelaki *et al.*, 2002).

Vascular ECs play a pivotal role in modulating local and systemic inflammation. LPS has potent pro-inflammatory properties and acts on many cell types including ECs (Berman *et al.*, 1993; Munshi *et al.*, 2002). There is much evidence indicating that COX-2, the rate-limiting enzyme in the metabolism of arachidonic acid, is involved in inflammatory diseases (Dubois *et al.*, 1998; Cipollone and Fazia, 2006); ICAM-1, as an important cellular adhesion molecule, has also

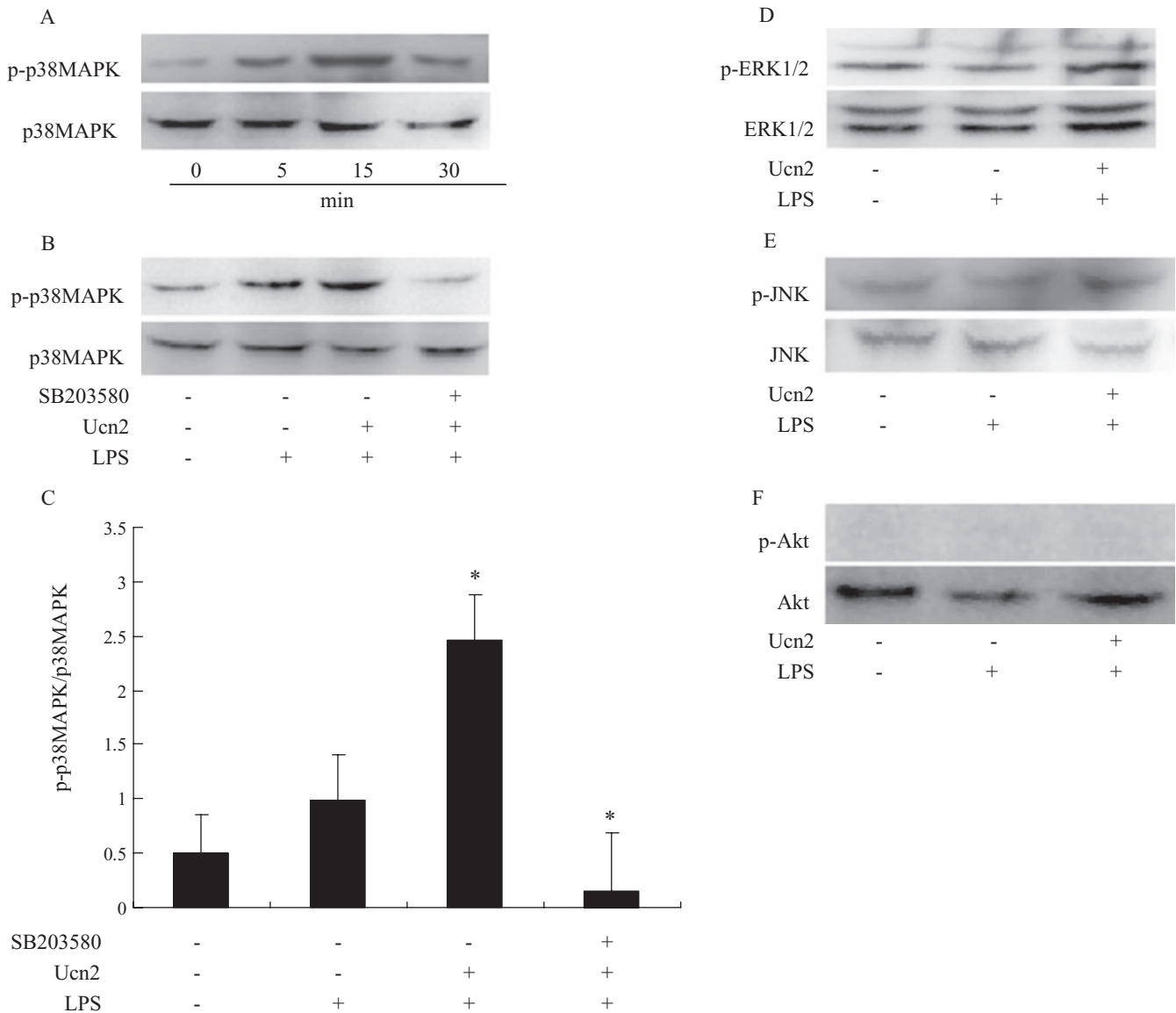


Figure 5 Effect of Ucn2 on mitogen-activated protein kinases (MAPKs) and Akt in lipopolysaccharide (LPS)-induced activation of rat aortic endothelial cells. Cells were treated with vehicle or Ucn2 (10^{-9} M) in the presence of LPS ($10 \mu\text{g}\cdot\text{mL}^{-1}$). (A) For p38MAPK determination, cells were treated with Ucn2 (10^{-9} M) and incubated for the durations shown. Phospho-p38MAPK was obviously observed at the time point of 15 min. The p38MAPK inhibitor SB203580 (10^{-5} M) was used to investigate the effect of p38MAPK phosphorylation (B and C). (D and E) For ERK1/2 and JNK measurement, cells were incubated for 15 min, and phospho-ERK1/2, JNK were analysed by Western blot. (F) Effect of Ucn2 on Akt phosphorylation. Results given are the means \pm SEM of values taken from three independent cultures. * $P < 0.05$, versus LPS group. Similar results were obtained from more than three independent cultures, and a representative experiment is shown.

been shown to have a key role in modulating peripheral inflammatory disease, such as atherosclerosis and TAO (Halacheva *et al.*, 2002; Witkowska, 2005). Several studies have illustrated that LPS-activated ECs demonstrate an up-regulation of COX-2 and ICAM-1 expression (Chen *et al.*, 2001b; Heo *et al.*, 2008). However, the function of Ucn in RAECs and its potential role in modulating COX-2 and ICAM-1 expression have rarely been illustrated.

In the present study, we found that Ucn1 exerted a pro-inflammatory effect by augmenting LPS-induced COX-2 and ICAM-1 expression in RAECs in a time- and concentration-dependent manner via the CRF₂ receptor. Indeed, Ucn1 up-regulated the COX-2 and ICAM-1 mRNA expression to

reach a maximum at 4 h, and the protein levels peaked at 8 h. This effect was via CRF₂ receptors as it could be abolished by the CRF₂ receptor antagonist, antisauvagine-30, but not by the CRF₁ receptor antagonist, NBI-27914. The above effects were further confirmed by measuring the levels of PGE₂ (the main metabolite of COX-2) and sICAM-1 in the culture supernatant of LPS-activated RAECs. The present data are consistent with those from previous studies. For example, it has been demonstrated that CRF-related peptides induce COX-2 expression and PG production in human placental trophoblasts and macrophages (Tsatsanis *et al.*, 2007; Gao *et al.*, 2008). Kageyama *et al.* (2006) showed that Ucn1 and -2 induced the expression of COX-2 in a time- and dose-dependent manner via CRF₂

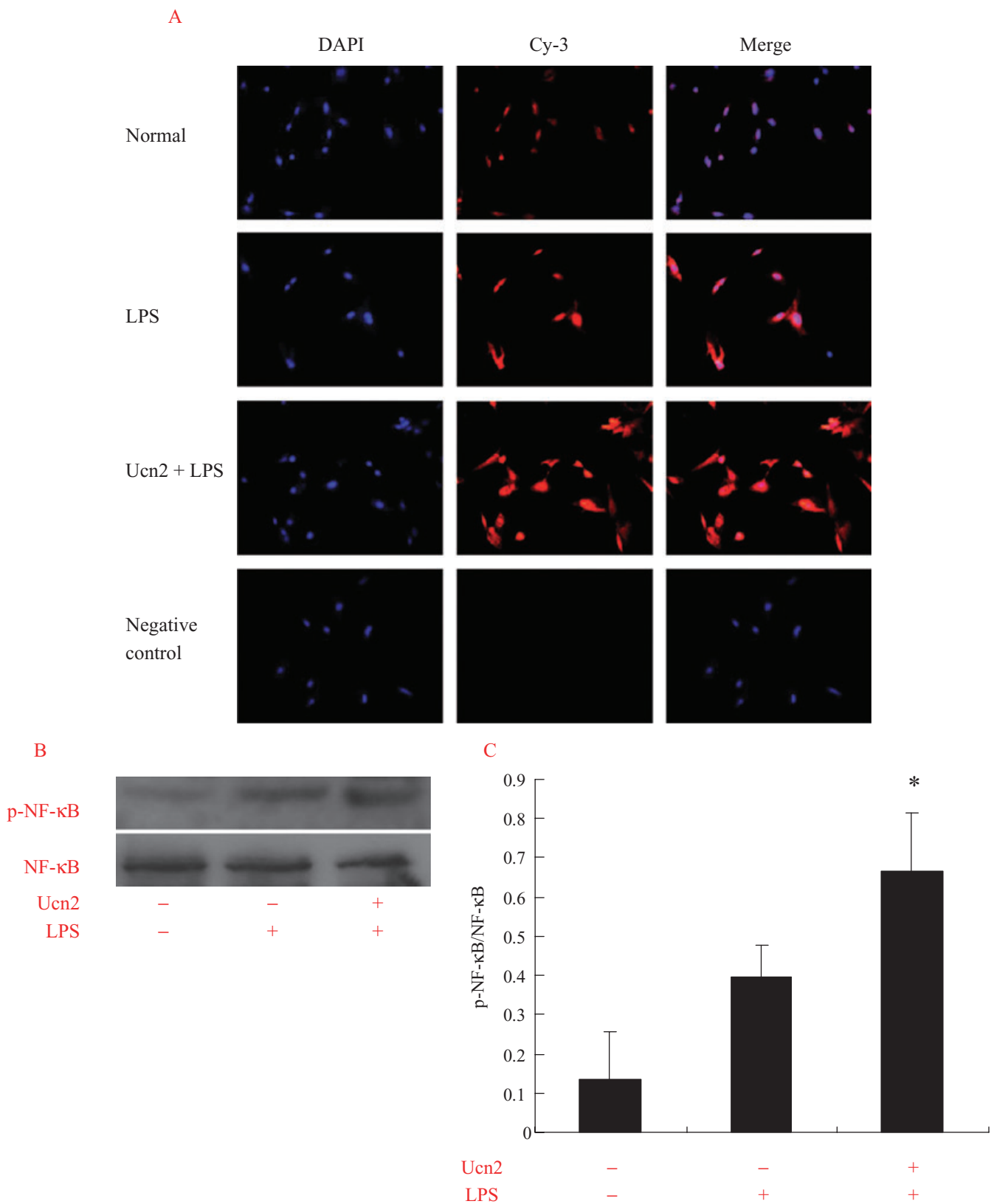


Figure 6 Effect of Ucn2 on the NF- κ B pathway in lipopolysaccharide (LPS)-activated rat aortic endothelial cells. (A) Effect of Ucn2 on nuclear translocation of NF- κ B. Cells cultured on coverslips were incubated with specific NF- κ B p65 antibody and then incubated with Cy-3 conjugated secondary antibody; the nucleus was stained with DAPI. A negative control was obtained by omitting NF- κ B p65 antibody. (Magnification: 200 \times). (B and C) Effect of Ucn2 on NF- κ B phosphorylation determined by Western blot. Results are presented as the means \pm SEM of values taken from three independent cultures. * P < 0.05, versus LPS group. Similar results were obtained from more than three independent cultures, and a representative experiment is shown.

receptors in rat aortic smooth muscle cells. Previously, it was found that CRF could enhance the interferon- γ -stimulated expression of ICAM-1 on human skin keratinocytes (Quevedo *et al.*, 2001). Taken together, our results demonstrate, for the first time, that Ucn can induce COX-2 and ICAM-1 expression in RAECs via the CRF₂ receptor.

Our previous study showed that the application of Ucn1 induced COX-2 and ICAM-1 expression via the CRF₁ receptor in the rat TAO model. This discrepancy with the present results might be attributed to the different distribution of CRF receptor types. In RAECs, the dominant receptor type is the CRF₂ receptor (Fekete and Zorrilla, 2007). Therefore, the effects of Ucn1 on RAECs are mainly implemented via the CRF₂ receptor. Kageyama *et al.* (2006) showed that Ucn1 and-2 could induce the expression of COX-2 via the CRF₂ receptor in rat aortic smooth muscle cells. However, in the rat TAO model, the level of CRF₁ receptors was elevated, while the expression of CRF₂ receptors was not significantly changed. This might be attributed to both the infiltration of immune cells bearing CRF₁ receptors and an effect of inflammatory mediators on CRF₁ receptor expression. On the one hand, the pro-inflammatory action of CRF peptides is partially the result of their effects on immune cells (such as mast cells, macrophages and lymphocytes) (Bamberger *et al.*, 1998; Theoharides *et al.*, 1998; Agelaki *et al.*, 2002), which can synthesize CRF peptides and express their receptors (Theoharides *et al.*, 1998; Baigent, 2001). On the other hand, inflammatory mediators could induce CRF₁ receptor expression (Inada *et al.*, 2009).

The MAPKs pathway is known to play a critical role in cytokine production (Bhat *et al.*, 1998; Means *et al.*, 2000), and Akt is involved in LPS-TLR4-mediated cytokine expression in macrophages and microglia (Jones *et al.*, 2001; Ojaniemi *et al.*, 2003; Kim *et al.* 2004). Recent studies have shown MAPKs, Akt and NF- κ B pathways are key downstream signalling intermediates of CRF₂ receptor activation (Dermitzaki *et al.*, 2002; Sananbenesi *et al.*, 2003; Karteris *et al.*, 2004; Moss *et al.*, 2007; Markovic *et al.*, 2008). In order to further clarify the mechanisms of the effect of Ucn on the expression of COX-2 and ICAM-1 in LPS-induced RAECs activation, the intracellular cell signalling pathways involved were investigated. Because the modulating effect was via the CRF₂ receptor, Ucn2, which binds with high affinity to the CRF₂ receptor (Fekete and Zorrilla, 2007), was used. We found that Ucn2 activated the p38MAPK pathway, and SB203580, the highly specific inhibitor of p38MAPK, abolished this effect. In contrast to previous studies showing that the ERK1/2 and Akt pathways positively regulate CRF peptide-induced expression of pro-inflammatory genes (Kim *et al.*, 2004; Moss *et al.*, 2007), our present study did not provide significant evidence that ERK1/2, JNK and Akt pathways are involved in the effects of Ucn2 on LPS-induced activation of RAECs. These differences may be attributed to the use of different cell types. In addition to the activation of the p38MAPK pathway, Ucn2 induced nuclear translocation and phosphorylation of NF- κ B, which is consistent with previous findings (Moss *et al.*, 2007). Taken together, by showing the positive effect of Ucn2 on LPS-induced p38MAPK and NF- κ B activation in RAECs, the present study revealed part of the mechanism of the CRF₂ receptor in inflammation.

Interestingly, in the present study, we found that the change in the levels of COX-2 and ICAM-1 showed a similar trend. To explore their possible relationship, NS-398, the selective COX-2 inhibitor, was used. It was found that NS-398 reduced the expression of sICAM-1 (Figure 4IA,B). This indicates that ICAM-1 expression is partially mediated by COX-2. Previous studies have shown that COX-2 and ICAM-1 are expressed in a wide variety of cell types, such as HUVECs and mouse brain ECs (Seok *et al.*, 2006; Zhao *et al.*, 2008). Aspirin, the non-selective COX-2 inhibitor, significantly suppressed COX-2 and ICAM-1 expression induced by ox-LDL (Zhao *et al.*, 2008). This is consistent with our present findings. Furthermore, COX-2 and ICAM-1 are now thought to be regulated by NF- κ B, via the expression of inflammatory genes induced by the activation of NF- κ B (Chen, 2006). In human alveolar epithelial cells, TNF- α -induced ICAM-1 expression was mediated through activation of IKK β , phosphorylation of I κ B α at serine and the subsequent activation of NF- κ B (Rahman *et al.*, 1999; Huang *et al.*, 2003a), and COX-2 expression is elevated after activation of IKK α/β and NF- κ B (Chen *et al.*, 2000; Huang *et al.*, 2003b). Degradation of I κ B or blockade of I κ B phosphorylation has been shown to suppress TNF- α -induced expression of COX-2 and ICAM-1 (Chen *et al.*, 1995; 2001a). These findings combined with the present results suggest that an intrinsic relationship exists between the two molecules in that ICAM-1 expression is partially mediated by COX-2, and NF- κ B is involved in this process.

In conclusion, this is the first time that Ucn has been shown to increase LPS-induced COX-2 and ICAM-1 expression, in RAECs, in a time- and concentration-dependent manner. Furthermore, this effect is mediated via the CRF₂ receptor via a mechanism involving both the p38MAPK and NF- κ B pathways.

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Conflicts of interest

None.

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