

Angiotensin II and the JNK pathway mediate urotensin II expression in response to hypoxia in rat cardiomyocytes

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Abstract

Cardiomyocyte hypoxia causes cardiac hypertrophy through cardiac-restricted gene expression. Urotensin II (UII) cooperates with activating protein 1 (AP1) to regulate cardiomyocyte growth in response to myocardial injuries. Angiotensin II (AngII) stimulates UII expression, reactive oxygen species (ROS) production, and cardiac hypertrophy. This study aimed to evaluate the expression of UII, ROS, and AngII as well as their genetic transcription after hypoxia treatment in neonatal cardiomyocytes. Cultured neonatal rat cardiomyocytes were subjected to hypoxia for different time periods. UII (Uts2) protein levels increased after 2.5% hypoxia for 4 h with earlier expression of AngII and ROS. Both hypoxia and exogenously added AngII or Dp44mT under normoxia stimulated UII expression, whereas AngII receptor blockers, JNK inhibitors (SP600125), JNK siRNA, or *N*-acetyl-L-cysteine (NAC) suppressed UII expression. The gel shift assay indicated that hypoxia induced an increase in DNA–protein binding between UII and AP1. The luciferase assay confirmed an increase in transcription activity of AP1 to the UII promoter under hypoxia. After hypoxia, an increase in ³H-proline incorporation in the cardiomyocytes and expression of myosin heavy chain protein, indicative of cardiomyocyte hypertrophy, were observed. In addition, hypoxia increased collagen I expression, which was inhibited by SP600125, NAC, and UII siRNA. In summary, hypoxia in cardiomyocytes increases UII and collagen I expression through the induction of AngII, ROS, and the JNK pathway causing cardiomyocyte hypertrophy and fibrosis.

Key Words

- ▶ hypoxia
- ▶ cardiac hypertrophy
- ▶ reactive oxygen species
- ▶ transcriptional activity

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Introduction

Cardiac hypertrophy and remodeling are considered to be compensatory processes in response to increased cardiac workload, caused by mechanical stress, hypertension, neurohumoral stimuli, myocardial injuries, or other

environmental factors (Ni *et al.* 2006). After such events, the heart undergoes compensatory adaptation to maintain adequate circulation, which is regulated by cardiac-restricted genes. These adaptive changes include

cardiac hypertrophy, proliferation of cardiac fibroblasts, and adjustment in the interstitial tissue. However, the persistence of these modifications may ultimately result in heart failure. Thus, cardiac hypertrophy is both a response to stress and an independent risk factor for cardiac morbidity and mortality.

Urotensin II (UII) has been identified as an endogenous peptide with a greater magnitude of vasoconstriction than endothelin 1 (ET-1), making UII the most potent mammalian vasoconstrictor (Ames *et al.* 1999). *Uii* (*Uts2*) mRNA is widely expressed throughout the cardiovascular system (Papadopoulos *et al.* 2008, Tolle & van der Giet 2008). *Uii* has been shown to increase cardiac hypertrophy, cardiac fibrosis, and collagen synthesis both *in vitro* and *in vivo* (Zhu *et al.* 2006, Zhang *et al.* 2007). Reactive oxygen species (ROS) are involved in the UII-induced cardiomyocyte hypertrophy (Liu *et al.* 2009). One study has demonstrated that the generation of ROS is involved in UII-induced cell proliferation, tyrosine phosphorylation of epidermal growth factor receptors (EGFRs), and extracellular signal-regulated kinase (ERK) phosphorylation in rat cardiac fibroblasts (Chen *et al.* 2008). Moreover, it is now well established that UII levels are significantly increased in several cardiovascular diseases (Bousette & Giaid 2006). Thus, UII may be an important determinant of cardiac dysfunction via additional non-hemodynamic effects on the myocardium, similar to angiotensin II (AngII) and ET-1 (Li *et al.* 2005). Specifically, Tzanidis *et al.* (2003) reported that UII stimulates collagen synthesis in cardiac fibroblasts, suggesting that UII may be involved in myocardial fibrogenesis. UII and its receptor (UII-R) are upregulated in patients with end-stage heart failure (Ng *et al.* 2002), ischemia (Pakala 2008), chronic hypoxia (MacLean *et al.* 2000), or post-myocardial infarct rat myocardium (Tzanidis *et al.* 2003). Upregulation of UII-R may worsen cardiac hypertrophy (Segain *et al.* 2007) and cardiac injury under ischemia-reperfusion (Prosser *et al.* 2008). The above findings suggest that UII may contribute to cardiovascular diseases through synergistic interactions with other vasoactive substances.

In response to hypoxia, the heart also generates ROS, resulting in cardiac hypertrophy and expression of fetal cardiac genes (Inamoto *et al.* 2006, Kontaraji *et al.* 2007, Yamashita *et al.* 2007) through various signaling cascades (Seta & Millhorn 2004). A previous study has shown that the density of binding sites for UII in sarcolemma of the myocardium increased in rats exposed to chronic hypoxia (Zhang *et al.* 2002). Interestingly, UII-R upregulation in the myocardium of hypoxic rats was accompanied by ventricular hypertrophy (Huang *et al.* 2006). Hypoxia can

also stimulate cardiomyocytes to secrete cytokines and growth factors (Inamoto *et al.* 2006). However, the effect of hypoxia on UII expression in cardiomyocytes is not yet fully understood. AngII has been shown to induce cardiac hypertrophy by increasing oxidative stress and UII expression (Lamarre & Tallarida 2008). Earlier studies also indicated that AngII stimulated both UII expression and genetic transcription in cardiomyocytes to cause cardiac hypertrophy (Song *et al.* 2012). However, the expression of AngII and UII after hypoxia and the relationship between them in regulating genetic transcription in cardiomyocytes under hypoxia have not been well defined. Transcriptional regulation of UII-R by nuclear receptor- κ B and hypoxia-inducible factor-1 α has been reported (Segain *et al.* 2007, Gould *et al.* 2010). Moreover, the JNK pathway has been reported to mediate the oxidative stress induced by UII (Djordjevic *et al.* 2005). Our recent study documented that hypoxia-induced UII expression in cardiac fibroblasts is mediated by AngII and through the ROS and JNK pathways (Shyu *et al.* 2012). Thus, UII is a mediator of AngII-induced cardiac fibrosis under hypoxia.

This study was designed to investigate the molecular mechanisms and signal pathways mediating UII expression in cardiomyocytes under hypoxia and AngII stimulation. We hypothesized that activating protein 1 (AP1), a well-characterized downstream target of JNK, may play a role in the transcriptional regulation of UII. Our results indicated that, following hypoxia, cardiomyocyte hypertrophy, and cardiac-restricted gene expression were modulated in response to UII expression, specifically through the JNK pathway. AngII was expressed in response to hypoxia to enhance UII expression and cardiomyocyte gene transcription, as previously demonstrated in a cardiac fibroblast model. We also identified the signal transduction pathway that modulates these mechanisms after hypoxic injury.

Materials and methods

Primary culture of left ventricular cardiomyocytes

Cardiomyocytes were obtained from 2- to 3-day-old Wistar rats as described previously (Shyu *et al.* 2001). Cultured myocytes were >95% pure as revealed by observation of contractile characteristics with a light microscope and were stained with anti-desmin antibody (Dako Cytomation, Glostrup, Denmark). The culture medium consisted of DMEM/F12 supplemented with 20% knockout serum replacement, 1% non-essential amino

acids, 2 mM L-glutamine, 0.1% β-mercaptoethanol, and 0.1% of a commercially available penicillin/streptomycin stock solution (Invitrogen; 100 U/ml penicillin and 100 µg/ml streptomycin). The cells were transferred to serum-free medium (Ham's F-12-DMEM, 1:1) and maintained for another 2 days. The enriched myocytes were then subjected to hypoxia. The rats were handled according to the Guide for the Care and Use of Laboratory Animals published by the US National Institute of Health (NIH Publication No. 85-23, revised 2011). The study was reviewed and approved by the Institutional Animal Care and Use Committee of Shin Kong Wu Ho-Su Memorial Hospital.

Hypoxia settings for cardiomyocyte culture

Hypoxic conditions were achieved by adding medium pre-equilibrated with nitrogen gas to the cells prior to incubation in a Plexiglas chamber purged with water-saturated nitrogen gas by an oxygen controller (PROOX model 110; Biospherix Ltd., Redfield, NY, USA). The partial pressure of oxygen (pO_2) of the culture medium under hypoxic condition was monitored using an ISO₂-dissolved oxygen meter (World Precision Instruments, Inc., Sarasota, FL, USA). Measurements indicated that a steady state in the culture medium was maintained during experiments. Hypoxia culture media (BioSpherix C-chamber) was used with mixed air in and out controlled by a BioSpherix PROOX incubator. Hypoxia settings were as follows: i) 10% O₂, 5% CO₂, and 85% N₂; ii) 5% O₂, 5% CO₂, and 90% N₂; and iii) 2.5% O₂, 5% CO₂, and 92.5% N₂. The use of a lower oxygen concentration (1%) was also tested, but the cardiomyocytes did not survive under such severe hypoxia. Hypoxia was applied for 1, 2, 4, and 6 h and for longer duration. The culture media were not changed during the duration of the experiment.

Antibodies and reagents

Polyclonal anti-UII, anti-UII-R, anti-collagen I, and anti-AngII antibodies were obtained from Santa Cruz Biotechnology. Different inhibitors were used to identify the signaling pathways involved in hypoxia-induced UII expression (Seta & Millhorn 2004). Polyclonal antibodies against JNK and mAbs against phospho-JNK were obtained from Cell Signaling (Beverly, MA, USA). Mouse mAbs against p38 MAPK and ERK were purchased from BD Bioscience Pharmingen (San Diego, CA, USA). PD98059, SB203580, and SP600125 were purchased from Calbiochem (San Diego, CA, USA). Monoclonal anti-α-tubulin

antibody was obtained from Sigma. The role of JNK, p38 MAPK, and ERK in hypoxia-induced UII expression was determined by pretreatment of the myocytes with 25 µM SP600125, 3 µM SB203580, or 50 µM PD98059 for 30 min before inducing hypoxia. SP600125 is a potent, cell-permeable, selective, and reversible inhibitor of JNK. SB203580 is a highly specific and cell-permeable inhibitor of p38 MAPK. PD98059 is a specific and potent inhibitor of the ERK pathway. AngII was purchased from Bachem AG (Torrance, CA, USA). To examine the effect of ARB (AngII type 1 receptor blocker), cardiomyocytes were treated with 100 nM losartan (Merck & Co., Inc.). N-acetyl-L-cysteine (NAC; a free radical scavenger) and Dp44mT (2,2'-dipyridyl-N,N-dimethylsemicarbazone; an ROS generator) were purchased from Calbiochem and used at a concentration of 500 µM and 30 nM respectively.

RNA isolation and real-time quantitative PCR

Total cellular RNA was extracted from cardiomyocytes (~1.5 × 10⁶ cells) using TRI reagent (Molecular Research Center, Cincinnati, OH, USA) according to the manufacturer's instructions. One microgram of total RNA was reverse transcribed using M-MuLV reverse transcriptase (Finnzyme, Vantaa, Finland) in a total volume of 20 µl. The reverse transcriptase products were amplified with the DyNAmo HS SYBR Green qPCR Kit (Finnzyme) in the reaction mix containing DyNAmo SYBR Green Master Mix and primers. Primers were designed for detection of UII gene expression (forward, 5'-CGGACACTGGTCAAATG-3' and reverse, 5'-CCTGGTTCTCGCCAAA-3'). *Gapdh* gene expression was used as an internal control (forward, 5'-GAGAGGCTCTCTGTCGACTAC-3' and reverse, 5'-TAGTGTAGGTTGGGCGCTCAA-3'), and did not change under the above conditions.

Western blot analysis

Cardiomyocytes were washed in PBS and lysed with RIPA buffer containing 1% (v/v) Nonidet P40, 0.5% SDS, and protease inhibitor cocktail containing 10 µg/ml phenylmethylsulphonyl fluoride, 2 µg/ml leupeptin, 2 µg/ml pepstatin A, and 2 µg/ml aprotinin. Cells were disrupted by intermittent sonication. After centrifugation, the amount of protein in the supernatant was measured using BSA as a standard. Cell lysates were then subjected to SDS/PAGE followed by western blotting. Antigen-antibody complexes were detected using with HRP-labeled rabbit anti-mouse immunoglobulin G (IgG) or goat anti-rabbit IgG and an ECL detection system (Pierce, Rockford, IL, USA).

RNA interference

Neonatal cardiomyocytes were transfected with 800 ng JNK- and UII-annealed siRNA (Dharmacon, Lafayette, CO, USA). JNK or UII siRNAs are target-specific 20- to 25-nt siRNAs designed to knock down gene expression. The JNK sense and antisense siRNA sequences were 5'-CGUGGAUUUAUGGUCUGUGdTdT and 5'-CACAGACC-AUAAAUCCACGdTdT respectively. The UII sense and antisense of siRNA sequences were 5'-CACGGACACUGGUGAAAUGUU and 5'-PCAUUUCACCAGUGUCCGUGUU respectively. As a negative control, a non-targeting (control) siRNA (Dharmacon) was used. Neonatal cardiomyocytes were transfected with siRNA oligonucleotides, using Effectene transfection reagent according to the manufacturer's instructions (Qiagen). After incubation at 37 °C, cells were kept under hypoxia and analyzed by western blot.

ROS assay and mitochondria isolation from cultured cardiomyocytes

ROS production was measured using the cell-permeable probe 2',7'-dichlorodihydrofluorescein diacetate, which passively diffuses into the cells where intracellular esterases cleave the acetate groups to form the impermeable DCFH₂, which remains trapped within the cells (McLennan & Degli Esposti 2000). After hypoxia treatment, cells were collected by trypsinization and resuspended in PBS medium. The ROS assay was performed according to the manufacturer's instructions (Invitrogen). Fluorescence microscopy was used to detect the green fluorescence. A mitochondrial isolation kit (Pierce) was used to isolate mitochondria from cultured cardiomyocytes as described previously (Zhang *et al.* 2010). Briefly, cultured cardiomyocyte pellets were gently lysed using a proprietary formulation that results in maximum yield of mitochondria with minimal damage to integrity.

ELISA for AngII

The level of AngII was measured using a quantitative sandwich enzyme immunoassay (SPI-BIO, Massy, France), using a specific anti-AngII antibody (Peninsula Laboratories, St Helens, Merseyside, UK), as described previously (Baker *et al.* 2004). Conditioned media from cultured cardiomyocytes under hypoxia and from control cells (normoxia) were collected for AngII measurement. The lower limit of detection of AngII was 0.5–1.5 pg/ml. Both the intra-assay and the interassay coefficients of variation were <10%.

Electrophoretic mobility shift assay

Nuclear protein concentration from cultured myocytes was determined by the Bradford method following the manufacturer's recommendations (Bio-Rad). Consensus and control oligonucleotides (Research Biolabs, Singapore) were labeled by the polynucleotide kinase incorporation of [γ -³²P] ATP. The consensus oligonucleotide sequence of AP1 was 5'-CGCTTGATGACTCAGCCGGA-3'. The AP1 mutant oligonucleotide sequence was 5'-CGCTTGATGACTTGCCCGGAA-3'. Electrophoretic mobility shift assay (EMSA) was performed as described previously (Chang *et al.* 2003). In each case, mutant or cold oligonucleotide, controls were used to compete with labeled sequences.

Promoter activity assay

A –664 to +44 bp rat UII promoter construct was generated as follows: rat genomic DNA was amplified with forward primer (CACAGCACTTTGCCAGGTTGA) and reverse primer (AGGAGTCCTACGAAGAGCAGG). The amplified product was digested with MluI and BglII restriction enzymes and ligated into pGL3-basic luciferase plasmid vector (Promega) digested with the same enzymes. The UII promoter contains the AP1-conserved sites (CA) at –621 to –620 bp. To produce the mutant, the UII binding sites were mutated using a mutagenesis kit (Stratagene, La Jolla, CA, USA). Site-specific mutations were confirmed by DNA sequencing. Plasmids were transfected into cardiomyocytes using a low-pressure accelerated gene gun (Bioware, New Taipei City, Taiwan).

Determination of protein synthesis

Protein synthesis was examined by measuring ³H-proline incorporation into trichloroacetic acid-precipitated macromolecules. Cultured cardiomyocytes were divided into the following groups: i) the control group, in which cells were cultured in serum-free DMEM and ii) the hypoxia group, in which 2.5% O₂ was added to serum-free medium. Each experiment was repeated six times. Cardiomyocytes were first grown in DMEM with 10% fetal bovine serum (FBS) and 200 mg/l L-glutamine and then seeded in 24-well plates at 1 × 10⁵ cells/well in DMEM + 10% FBS. After synchronization of cardiomyocytes, the medium was changed to DMEM without serum. Cardiomyocytes were incubated under hypoxia and exposed to ³H-proline at a concentration of 1 μCi/well for the last 1 h of the 4-h incubation period. After incubation, the cells

were washed with ice-cold PBS and 10% trichloroacetic acid. Acid-insoluble ^3H -proline was collected on glass fiber filters (Whatman, Kent, UK) and measured using a liquid scintillation counter (LS 6500; Beckman, Fullerton, CA, USA).

Statistical analysis

The results are expressed as the mean \pm s.d. Statistical significance was evaluated using ANOVA followed by the Tukey–Kramer multiple comparisons test (GraphPad Software, Inc., San Diego, CA, USA). A value of $P < 0.05$ was considered to denote statistical significance.

Results

Hypoxia increases the expression of UII in cultured cardiomyocytes

Different degrees of hypoxia (10, 5, and 2.5%) for different periods of time (1, 2, 4, 6 h, and longer) were used as described in the ‘Materials and methods’ section (Supplementary Fig. 1, see section on supplementary data given at the end of this article). UII expression was notably increased when cardiomyocytes were incubated in 2.5% O_2 hypoxia conditions for 4 h compared with other conditions. Thus, these culture conditions were used for further analysis. There was no significant difference in pH between the different levels of hypoxia. In 2.5% O_2 hypoxia, UII and UII-R protein levels increased gradually and reached a peak after 4 h (Fig. 1A and B). UII mRNA expression also reached its maximal level after 2 h and then decreased gradually (Fig. 1C). UII protein expression was suppressed by UII siRNA (Supplementary Fig. 2).

Hypoxia-induced UII protein expression in cardiomyocytes is mediated by ROS and the JNK pathway

As shown in Fig. 2A and B, ROS production in cardiomyocytes was significantly increased by 2 h of hypoxia exposure. This effect was similar to that of the exogenous addition of Dp44mT. Similarly, ROS production from isolated mitochondria was significantly increased by hypoxia and Dp44mT treatment (Fig. 2C). Treatment with NAC for 30 min prior to subjecting the cardiomyocytes to hypoxia significantly blocked the hypoxia-induced ROS production. In the control group (normoxia), very few cardiomyocytes presented an increase in ROS production. In summary, hypoxia induces an increase in ROS production by cardiomyocytes and in mitochondria isolated from cardiomyocytes, suggesting

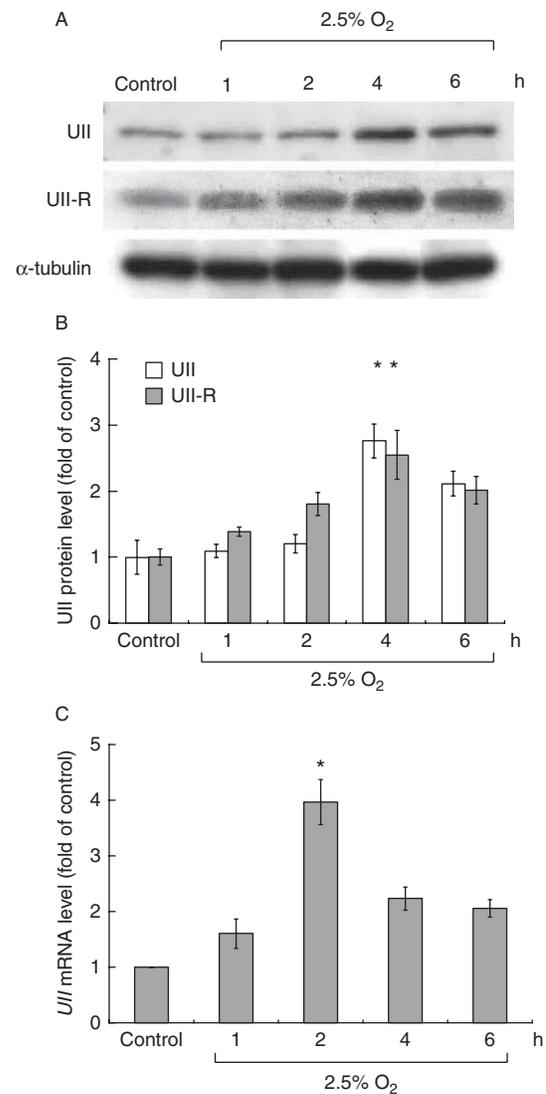
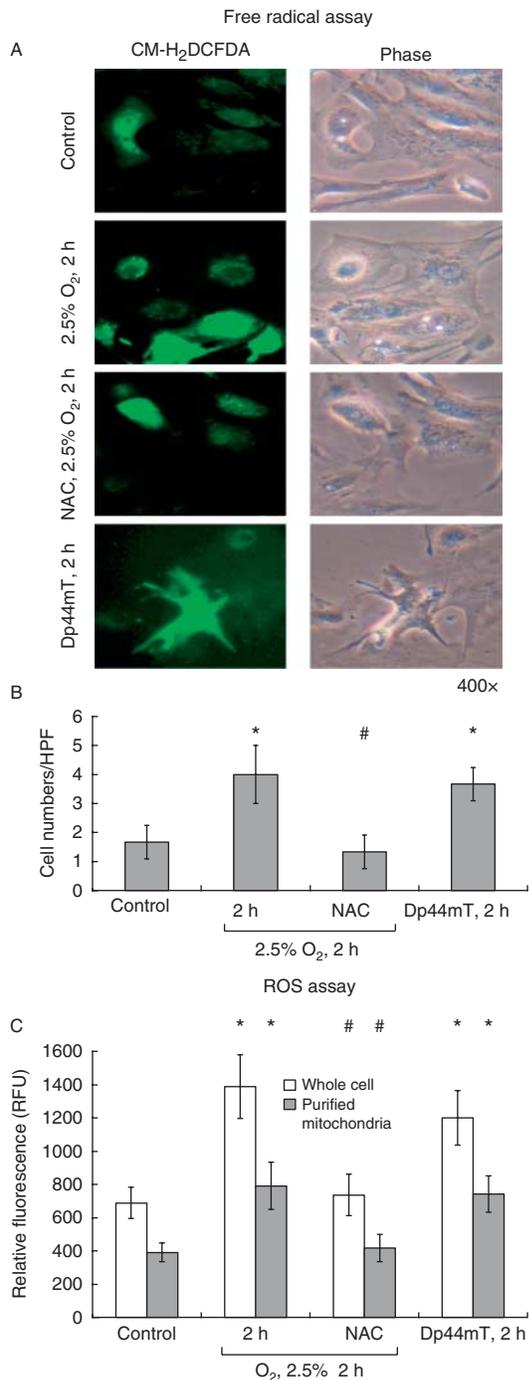


Figure 1

Effect of 2.5% O_2 hypoxia on UII expression in cardiomyocytes. (A and B) UII and UII-R protein levels increased, reaching a peak after 4 h of 2.5% O_2 hypoxia. Neonatal cardiomyocytes were subjected to hypoxia for 1–6 h. Total cell lysates were harvested and western blot was performed using anti-UII and anti-UII-R antibodies. α -tubulin was used as a protein-loading control in each lane. $*P < 0.01$ vs normoxia control ($n = 3$). (C) UII mRNA level reached a peak after 2 h of hypoxia and then declined. GAPDH was used as an internal control. $*P < 0.01$ vs normoxia control ($n = 3$). UII, urotensin II; UII-R, UII receptor.

that ROS is produced by the mitochondria and that changes in mitochondria participated in ROS generation in our experiments.

To define the pathways involved in hypoxia-induced UII expression, different inhibitors were used. Hypoxia-induced UII expression was significantly inhibited by the JNK inhibitor (SP600125). This effect was confirmed using a JNK siRNA or NAC (Fig. 3A and B). The ERK and P38

**Figure 2**

Hypoxia-induced UII protein expression in cardiomyocytes is mediated by (A and B) reactive oxygen species (ROS) assay, using a fluorescence microscope, indicated that hypoxia for 2 h increased ROS production similarly to that of the exogenous addition of ROS generator (Dp44mT). Treatment with NAC (500 μ M) 30 min before hypoxia significantly blocked the induction. (C) Mitochondria isolation from cultured cardiomyocytes. Hypoxia and exogenously added Dp44mT (30 nM) significantly increased the ROS relative fluorescence unit (RFU) after 2 h. Treatment with NAC for 30 min before hypoxia significantly blocked the RFU of ROS by hypoxia. * $P < 0.01$ vs normoxia control. # $P < 0.01$ vs 2 h ($n = 3$).

inhibitors did not attenuate the UII protein expression induced by hypoxia. JNK protein phosphorylation increased to its maximal level after 2 h of 2.5% O₂ hypoxia and then declined gradually. SP600125, JNK siRNA, and NAC could effectively block the phosphorylation of JNK protein (Fig. 3C and D). Exogenously added Dp44mT under normoxia also increased UII expression, which was inhibited by JNK siRNA (Fig. 4B and C).

AngII mediates the induction of UII by hypoxia in cultured cardiomyocytes

Under 2.5% O₂ hypoxia, the AngII protein level increased and reached a peak after 2 h of incubation (earlier than UII and UII-R protein peak expression after 4 h of incubation) and then declined gradually (Fig. 4A). The UII protein level increased after 2.5% O₂ hypoxia but was suppressed by ARB (losartan: 100 nM) and AngII blocking antibodies (Fig. 4B and C). In addition, exogenous addition of AngII under normoxia also increased UII expression (Fig. 4B and C).

Hypoxia increases the binding between the UII promoter and AP1 transcription factor and enhances genetic transcription activity of AP1 to UII promoter in cardiomyocytes

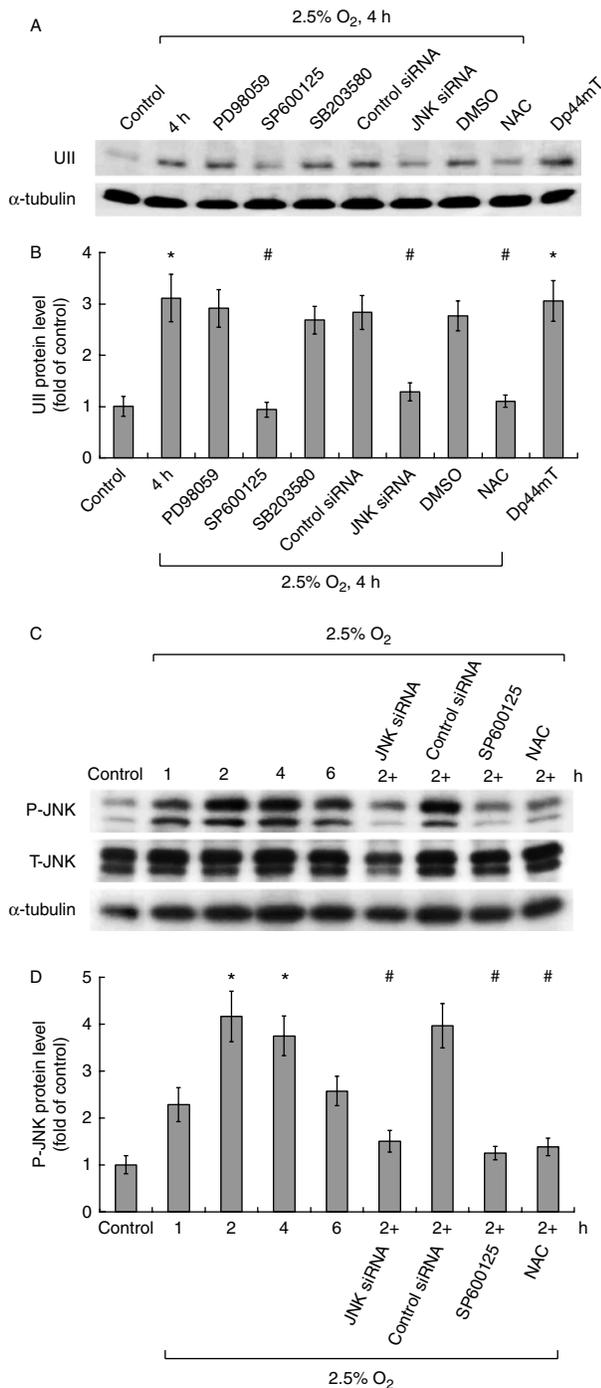
Under 2.5% O₂ hypoxia, EMSA showed increased binding activity between UII promoter and AP1 transcription factor. SP600125 and NAC inhibited the binding activity between UII promoter and AP1 transcription factor (Fig. 5A). This finding indicated that hypoxia in cardiomyocytes may increase the binding between UII promoter and AP1 transcription factor. Hypoxia-induced JNK phosphorylation and ROS may increase UII expression and UII/AP1 binding activity.

The luciferase reporter assay also showed (Fig. 5B and C) that hypoxia increased transcriptional activity of AP1 to UII promoter in cardiomyocytes. Mutation of the UII promoter inhibited this effect of hypoxia. Exogenously added AngII also increased the transcriptional activity of AP1 to UII promoter. SP600125, NAC, and losartan suppressed the transcriptional activity of the UII promoter in cardiomyocytes under hypoxia.

Hypoxia modulates collagen I protein expression in cardiomyocytes

As shown in Fig. 6, 4 h of hypoxia significantly increased collagen I protein expression in cardiomyocytes

(Fig. 6A and B). SP600125, NAC, and *Ull* siRNA effectively attenuated the collagen I protein expression induced by hypoxia (Fig. 6C and D). In addition, exogenous addition of AngII and Ull under normoxia also increased collagen I protein expression in cardiomyocytes. Under normoxia, AngII-induced collagen I protein expression could also be effectively suppressed by *Ull* siRNA (Fig. 6C and D).



Hypoxia increases collagen I expression in the perinuclei, protein synthesis, and genetic transcription to induce cardiomyocyte hypertrophy

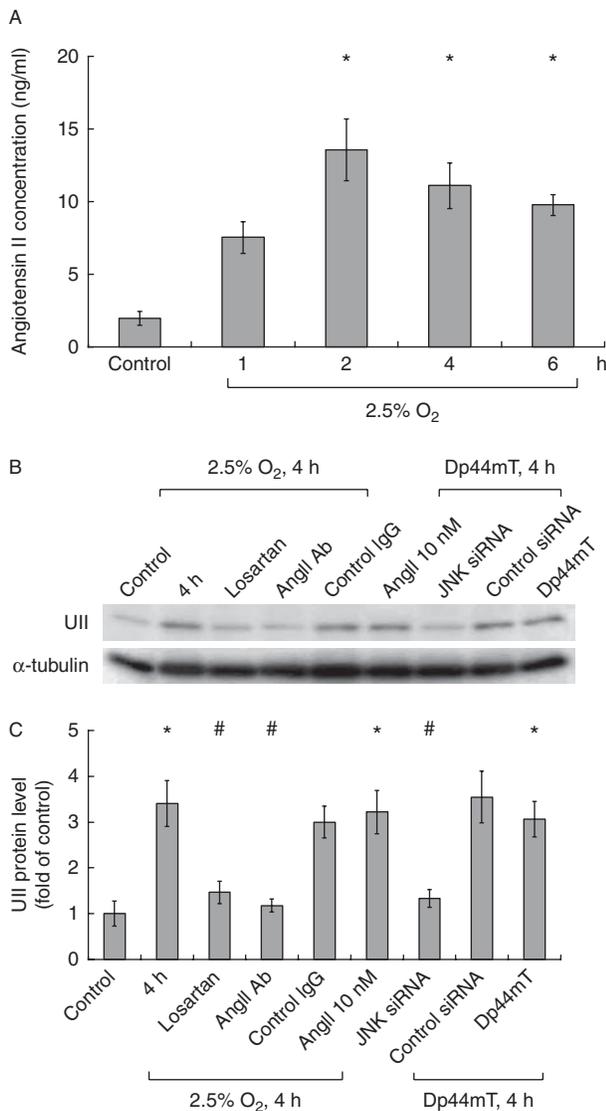
Under hypoxia, Ull increased transcriptional activity in cardiomyocytes. Confocal microscopy (Supplementary Fig. 3, see section on supplementary data given at the end of this article) showed increased collagen I expression (green) in the perinuclei (blue) 4 h after hypoxia, which contributed to hypertrophic changes in cardiomyocytes. Ang II and Ull also increased the expression of collagen I in the perinuclei of cardiomyocytes. Collagen I expression was suppressed by SP600125, NAC, and *Ull* siRNA. Protein synthesis in cardiomyocytes was also evaluated by measuring ^3H -proline incorporation into the cells. Results showed an increase in protein synthesis after 1–4 h of hypoxia, indicative of hypertrophic changes in cardiomyocytes (Fig. 7). Pretreatment with NAC, AngII antibody, losartan, *Ull* siRNA, and SP600125 inhibited the protein synthesis induced by hypoxia. Exogenous addition of Ull, AngII, and Dp44mT under normoxia also increased protein synthesis similarly to hypoxia.

Discussion

Ull plays an important role in cardiac hypertrophy and cardiac fibrosis. Vasoactive effects of mammalian Ull are dependent both on the species and on the regional vascular bed examined. Typical regional variability occurs in the rat in which vasoconstriction to Ull is most robust in thoracic aorta proximal to the aortic arch and decreases gradually toward the distal peripheral arteries. Small peripheral arteries play a major role in regulating peripheral resistance and consequent blood pressure as well as workload on the heart (Ames *et al.* 1999).

Figure 3

The JNK pathway mediates hypoxia-induced Ull expression in neonatal cardiomyocytes. (A and B) Hypoxia-induced Ull expression was blocked by JNK inhibitor (SP600125), JNK siRNA, and NAC. Exogenous addition of Dp44mT (a ROS generator) under normoxia also increased Ull expression. Neonatal cardiomyocytes were pretreated with an ERK pathway inhibitor (PD98059; 50 μM), a JNK inhibitor (SP600125; 25 μM), a p38 MAPK inhibitor (SB203580; 3 μM), NAC, or JNK siRNA prior to hypoxia for 4 h. The neonatal cardiomyocytes were then harvested and analyzed by western blotting using anti-Ull antibody. The results were normalized to α -tubulin. * $P < 0.01$ vs normoxia control. # $P < 0.01$ vs 4 h ($n = 3$). (C and D) Phosphorylation of JNK mediated hypoxia-induced Ull expression in neonatal cardiomyocytes, which was blocked by JNK inhibitor (SP600125), JNK siRNA, and NAC. Neonatal cardiomyocytes were subjected to normoxia or hypoxia for different durations in the presence or absence of inhibitors. Cell lysates were collected for western blot analysis, using antibody for total and phospho-JNK. T-JNK, total JNK; P-JNK, JNK phosphorylation. * $P < 0.01$ vs normoxia control. # $P < 0.01$ vs 2 h ($n = 3$).

**Figure 4**

AngII mediates hypoxia-induced UII expression in neonatal cardiomyocytes. (A) AngII was measured in cell lysates and the culture medium by a quantitative, competitive ELISA using a specific anti-AngII antibody. * $P < 0.01$ vs normoxia control. (B and C) Hypoxia-induced UII expression was suppressed by AngII antibody and losartan. Exogenous addition of Dp44mT or AngII (under normoxia) also increased UII expression. UII expression induced by exogenous addition of Dp44mT was suppressed by JNK siRNA. Neonatal cardiomyocytes were subjected to normoxia or hypoxia for 4 h and total cell lysates were immunoblotted with anti-UII antibody. α -tubulin was used as a protein loading control in each lane. * $P < 0.01$ vs normoxia control. # $P < 0.01$ vs 4 h ($n = 3$).

UII-R activation is associated with multiple physiological actions such as vasoconstriction, vasodilation, osmoregulation, inhibition of insulin, and cell proliferation (Zhu *et al.* 2006, Ames *et al.* 2008). Similar to AngII (Chen *et al.* 2004) and ET-1 (Chen *et al.* 2005), UII-induced myocyte hypertrophy may contribute to cardiac remodeling and

deterioration of cardiac function. Tzanidis *et al.* (2003) demonstrated the expression of both UII and UII-R within myocardial tissue, suggesting an important role for UII in cardiac pathophysiology. Zhou *et al.* (2003) reported UII-dependent hypertrophy in isolated cardiomyocyte cultures. Increased UII and UII mRNA as well as greater density of UII binding sites have been found in the myocardium and plasma of patients with congestive heart failure (Douglas *et al.* 2002, Gruson *et al.* 2006) and acute myocardial infarction (Khan *et al.* 2007), as well as in rats with diabetic cardiomyopathy (Dai *et al.* 2008). Bousette *et al.* (2006) reported that blockade of UII attenuated cardiac dysfunction in a rat model of coronary ligation. Furthermore, UII correlated significantly with ET-1 and brain natriuretic peptide, suggesting that UII may play a role in congestive heart failure worsening (Tang *et al.* 2010). UII may contribute to cardiac hypertrophy through upregulation of UII-R and inflammatory cytokines (Tzanidis *et al.* 2003).

Hypoxia is a commonly seen pathological phenomenon in cardiovascular diseases, including coronary artery disease, myocardial infarction, and heart failure. In cardiomyocytes, hypoxia produces stress, inducing the expression of specific cardiac-restricted genes. Tissue hypoxia is a feature of chronic inflammatory disease and may contribute to the development of oxidative stress and cardiomyocyte hypertrophy, as previously reported (Chiu *et al.* 2012). Recent studies have shown that UII stimulates NADPH oxidase, which generates ROS in pulmonary artery smooth muscle cells (Djordjevic *et al.* 2005). Human UII may worsen hypoxia-induced cardiac hypertrophy mainly through a vasoconstriction of the pulmonary artery and partly through the suppression of atrial natriuretic peptide secretion (Gao *et al.* 2010). Hypoxia stimulates UII-R upregulation and UII secretion both *in vitro* and *in vivo* (MacLean *et al.* 2000, Shyu *et al.* 2012). Finally, UII may act as an autocrine and/or paracrine hormone, playing an important role in the development of the ventricular hypertrophy induced by chronic hypoxia. Previous *in vivo* studies associated with pathologies generating ROS (heart failure, post-myocardial infarction, coronary artery ligation, ischemic-reperfusion injury, and diabetic cardiomyopathy) showed increased UII expression, indicating that administration of ROS *in vivo* may activate UII (Douglas *et al.* 2002, Ng *et al.* 2002, Bousette *et al.* 2006, Khan *et al.* 2007, Chen *et al.* 2008, Dai *et al.* 2008).

In this study, we identified some novel findings i) that hypoxia induces UII expression, causing cardiomyocyte hypertrophy; ii) that the JNK pathway specifically mediates hypoxia-induced UII expression in cardiomyocytes;

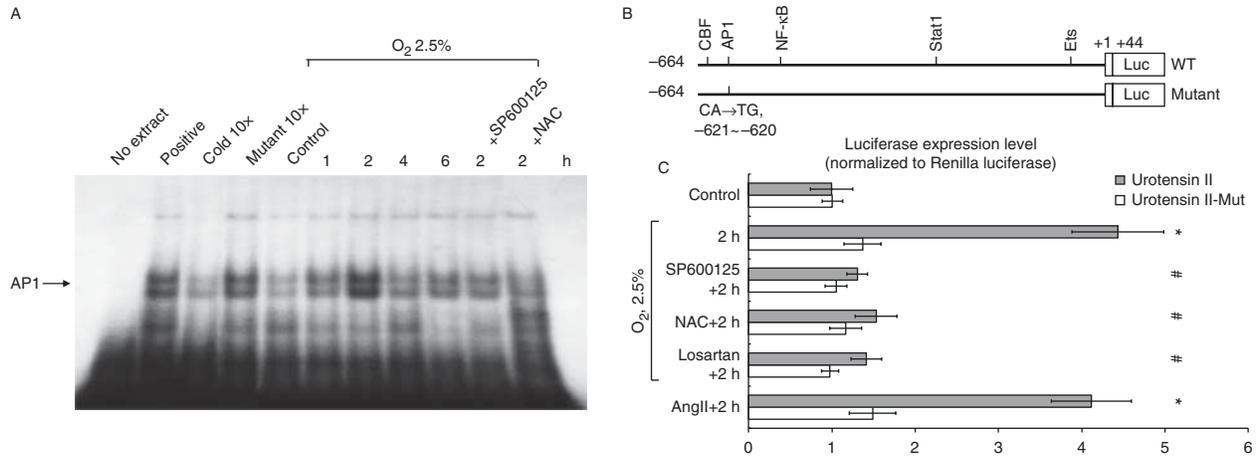


Figure 5 Binding activity between UII promoter and AP1 transcription factor and genetic transcription activity at the AP1-binding site of UII promoter increase in neonatal cardiomyocytes under hypoxia. (A) EMSA showed an increased in binding between UII promoter and AP1 transcription factor under in cardiomyocytes 2.5% O₂ hypoxia. The binding between UII promoter and AP1 transcription factor was suppressed by JNK inhibitor (SP600125) and NAC. (B and C) The luciferase reporter assay revealed that

2.5% O₂ hypoxia increased the transcriptional activity of AP-1 to the UII promoter compared to the UII mutant. Transcriptional activity was suppressed by JNK inhibitor (SP600125), NAC, and losartan. Exogenous addition of AngII (10 nM) increased the transcriptional activity of neonatal cardiomyocytes similar to that of 2.5% O₂ hypoxia. **P*<0.01 vs normoxia control. #*P*<0.01 vs 2 h (*n*=3).

iii) the effect of AngII and ROS on UII expression and genetic transcription in neonatal cardiomyocytes, causing cardiac hypertrophy; and iv) the relationship between AngII, ROS, UII, and collagen I expression in

cardiomyocytes under hypoxia, which results in cardiac hypertrophy and fibrosis.

This study yielded results similar to our previous reports (Chiu *et al.* 2010, 2012, Shyu *et al.* 2012) regarding

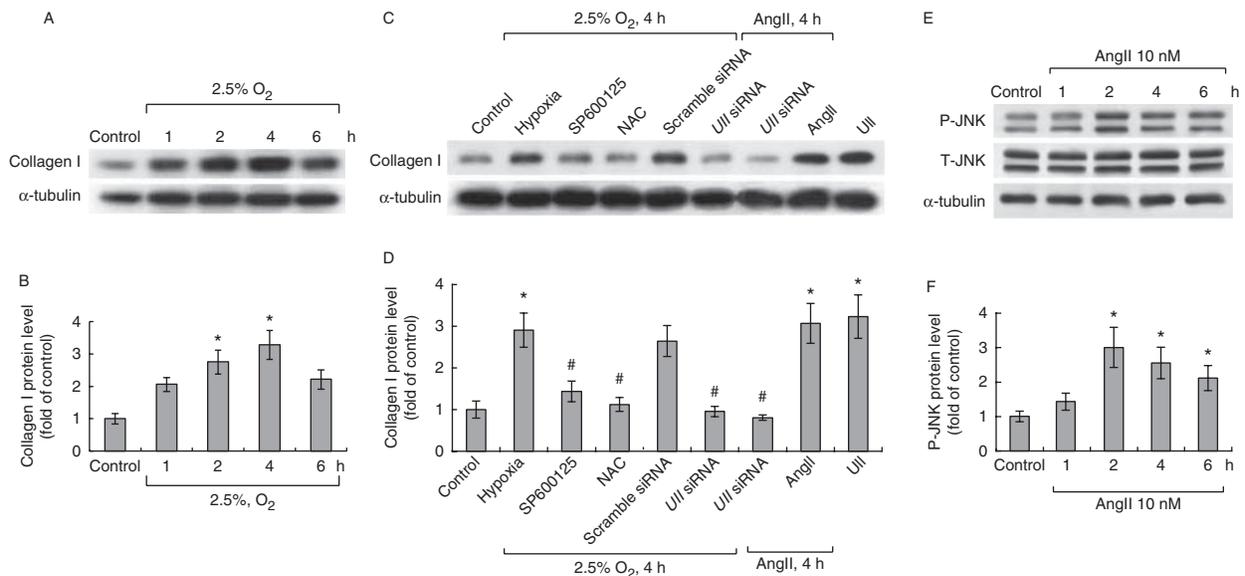


Figure 6 Hypoxia increases collagen I protein expression in cardiomyocytes. (A and B) Hypoxia increased collagen I expression, which reached a peak after 4 h. (C and D) Hypoxia-induced collagen I expression was effectively suppressed by JNK inhibitor (SP600125), NAC, and UII siRNA. Exogenous addition of AngII or UII (under normoxia) also increased the expression of collagen I.

Collagen I expression induced by AngII was effectively suppressed by UII siRNA. (E and F) Exogenously added AngII increased JNK phosphorylation, which reached a peak after 2 h. **P*<0.01 vs normoxia control. #*P*<0.01 vs hypoxia (*n*=6).

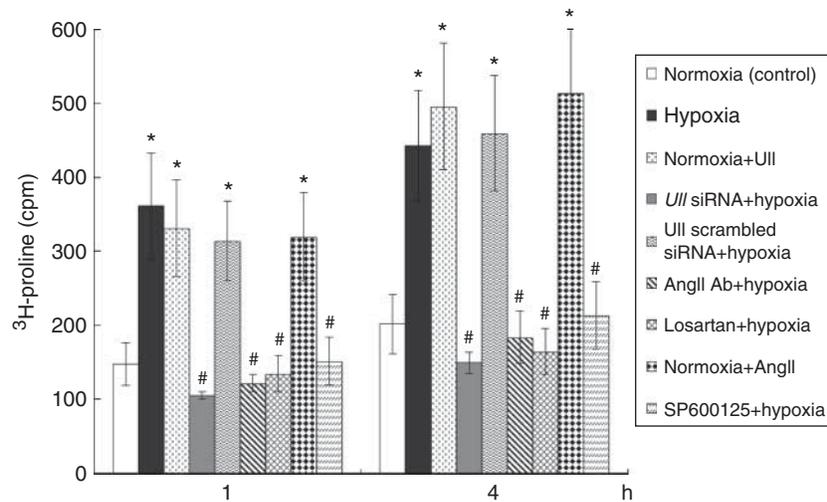


Figure 7

Neonatal cardiomyocytes protein synthesis increases after hypoxia. Incorporation of ³H-proline into neonatal cardiomyocytes increased after hypoxia and exogenous addition of Ull, Angll, or Dp44mT (under normoxia) for 1–4 h. Hypoxia-induced incorporation of ³H-proline into

neonatal cardiomyocytes was suppressed by Ull siRNA, Angll blocking antibody, losartan, NAC, and JNK inhibitor (SP600125). **P* < 0.01 vs normoxia control. #*P* < 0.01 vs hypoxia (*n* = 6).

the putative mechanism involved in the Ull increase observed in cardiomyocytes subjected to 2.5% hypoxia. Mild hypoxia (10 or 5%) did not induce significant oxidative stress to lead to an increase in Ull expression. In addition, extreme hypoxia ($\leq 1\%$) did not provide a suitable environment for cardiomyocytes to survive. Under 2.5% O₂ hypoxia, significant oxidative stress induced both ROS and hypoxic signaling (JNK) and activated Ull expression. Ull binds to a class of G protein-coupled receptor known as Ull-R. Ull and Ull-R are highly expressed in the cardiovascular system and are upregulated in hypoxic myocardium. They also enhance foam cell formation, chemotaxis of inflammatory cells, as well as inotropic and hypertrophic effects on cardiac muscle.

Our study also identified the JNK pathway as the mediator of the hypoxia-induced Ull expression in cardiomyocytes. The JNK pathway has been reported to mediate the oxidative stress induced by Ull (Djordjevic *et al.* 2005). Moreover, hypoxia has been shown to stimulate JNK activation in cardiomyocytes. JNK mediates signals in response to cytokines and environmental stress, including oxidative stress. Previous studies indicated that JNK activation and ROS generation are immediate responses to low oxygen stress. Because the JNK pathway has been previously shown to be responsible for the hypoxic signaling, we successfully identified that either JNK siRNA or NAC could effectively attenuate hypoxia-induced JNK pathway activation in this study.

Previous studies have reported that Ull expression and upregulation of Ull-R promote cardiomyocyte hypertrophy via the p38 and ERK pathways in an EGFR-dependent manner (Onan *et al.* 2004). However, in our study, the JNK pathway was the main pathway involved in the hypoxia-induced Ull expression in cardiomyocytes.

With regard to JNK phosphorylation, it means activation of JNK pathway. As seen in previous and our studies, increases in JNK tyrosine phosphorylation and JNK activity occurred at an early hypoxic exposure time. In terms of the downstream significance of JNK activation by hypoxia, it is now well documented that hypoxia leads to the activation of AP1 that participate in the response of cells to hypoxia. Similarly, JNK phosphorylation enhanced oxidative phosphorylation and JNK pathway transduction to facilitate Ull expression, and JNK inhibitor, JNK siRNA, and NAC repressed the expression of Ull. Oxidative phosphorylation is intimately related to redox biology through control of the JNK signaling.

The relationship between AngII, ROS, and Ull, as well as their regulatory effects on genetic transcription in cardiomyocytes, has not yet been defined. AngII has been shown to induce cardiac hypertrophy after hypoxia (Chiu *et al.* 2012). Hypoxia regulates the expression of Ull in cardiomyocytes, and AngII acts as an upstream regulator of Ull. Our study found that both hypoxia-induced AngII expression and exogenously added AngII-stimulated Ull expression and genetic transcription in cardiomyocytes.

These effects could be effectively inhibited by ARB. Infection with an adenovirus encoding the AT₁ receptor was necessary to produce a reproducible hypertrophic response to AngII, strongly implicating receptor upregulation as the determinant for initiating hypertrophy. The interaction between AngII and UII has been shown to increase myocardial dispensability (Fontes-Sousa *et al.* 2009) and vasoconstriction (Wang *et al.* 2007). Previous reports also showed that, with the addition of AngII, the myocardial UII/UII-R system was upregulated in isoproterenol-treated rats, indicating that the UII/UII-R system is a newly identified regulating system involved in cardiac hypertrophy (Zhang *et al.* 2007). In this study, we demonstrated that hypoxia increased the secretion of AngII from cardiomyocytes and that the secreted AngII increased UII expression, resulting in cardiac hypertrophy.

Our data also showed that ROS is involved in cardiomyocyte hypertrophy induced by UII. In fact, hypoxia-induced UII expression, UII promoter and AP1 binding, UII-related genetic transcription in cardiomyocytes, and cardiomyocyte hypertrophy could be effectively suppressed by NAC. Hypoxia has been shown to stimulate oxidative stress and increase ROS formation with subsequent cardiomyocyte hypertrophy (Chiu *et al.* 2012). UII has also been shown to induce ROS and cardiomyocyte hypertrophy (Liu *et al.* 2009). Hypoxic signaling (JNK) modulated by ROS from the mitochondrion to the nucleus controls and promotes cardiomyocyte hypertrophy. Thus, ROS may be a second messenger mediating the effects of growth-promoting neurohumoral agonists and intracellular signaling pathways. 2.5% hypoxia represents a condition of significant oxidative stress associated with subsequent elevation of ROS. Mitochondria have long been considered a likely site of oxygen sensing, and the electron transport chain acts as an O₂ sensor by releasing ROS in response to hypoxia. The ROS released during hypoxia act as signaling agents that trigger diverse functional responses, including activation of downstream transcription factor (AP1) and hypoxic signaling (JNK). The primary site of ROS production during hypoxia appears to be complex III in the electron transport chain in mitochondria. O₂-free radical is generated in the mitochondria and requires coupled electron transport, ROS generation, and calcium flux. So, ROS is a product of mitochondria and mitochondrial ROS signals initiate the activation of JNK (Laderoute & Webster 1997, Dougherty *et al.* 2004, Guzy & Schumacker 2006).

In different cell types, different MAPK pathways may be involved in UII genetic transcription. Our study shows that AP1, a downstream target of JNK, plays a role in the

transcriptional regulation of UII. The gel shift assay showed that UII promoter–DNA binding activity increased after hypoxia. This effect was abolished by the use of a UII-mutant plasmid, SP600125, and ARB. In cardiomyocytes, UII-related genetic transcription occurs through AP1 binding to the UII promoter. The promoter activity assay confirmed the regulation of UII expression by AP1. In fact, the effects of hypoxia and AngII stimulation on UII promoter activity were abolished when the AP1 binding site was mutated. So, the regulatory DNA in UII promoter may participate in the transcriptional activity.

Hypoxia is a stimulus to collagen synthesis in cardiomyocytes. This study demonstrated that hypoxia increased collagen I expression and the presence of collagen I in the nuclei. This effect was effectively inhibited by NAC, JNK inhibitor, and UII siRNA, indicating a relationship between AngII, ROS, UII, and collagen I. In addition, UII or AngII, added exogenously, increased collagen I expression. AngII-related collagen I expression was also effectively inhibited by UII siRNA. Previous studies indicated that UII has effects on myocardial protein synthesis and was documented by increased ³H-proline into neonatal cardiomyocytes and myosin

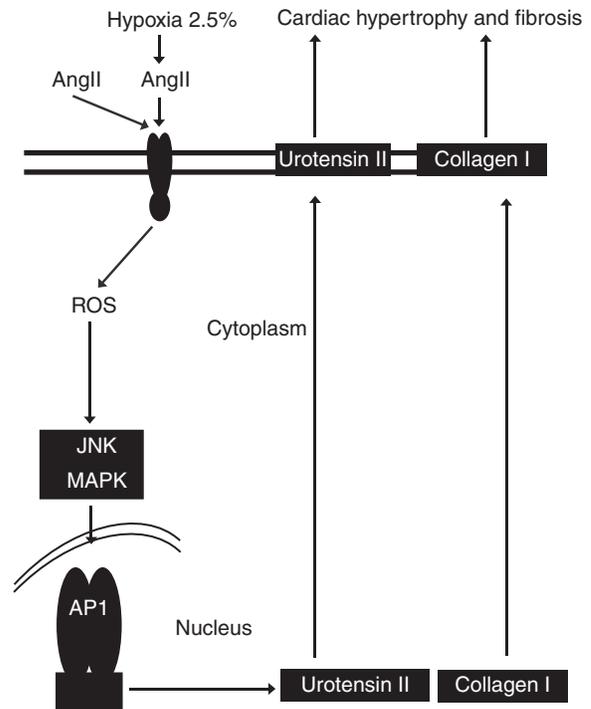


Figure 8 Schematic representation of the AngII/ROS/JNK pathway involved in the hypoxia-induced induction of UII expression leading to cardiomyocyte hypertrophy through increased transcriptional activity.

expression (Tzanidis *et al.* 2003). In our study, UII also increased protein synthesis by increased incorporation of ³H-proline into neonatal cardiomyocytes. In conclusion, UII, AngII, and ROS mediated the expression of collagen I in cardiomyocytes subjected to hypoxia, resulting in cardiac fibrosis in addition to cardiac hypertrophy. The role of UII in the development of cardiac fibrosis results from its ability to increase collagen protein synthesis and secretion in the extracellular matrix. AngII has been shown to induce the secretion of profibrotic factors such as syndecan, transforming growth factor- β 1, and connective tissue growth factor in cardiac fibroblasts. In our study, exogenous AngII stimulation increased UII expression, JNK protein phosphorylation, and collagen I protein expression. These findings indicate that UII is a mediator of AngII-induced cardiac fibrosis. Previous studies indicated that other stress factors, including ischemic cardiomyopathy, postmyocardial infarction, heart failure, ischemic-reperfusion injury, and pressure overload to myocardium also induce UII-related collagen expression (Douglas *et al.* 2002, Ng *et al.* 2002, Tzanidis *et al.* 2003, Boussette *et al.* 2006, Chen *et al.* 2008).

With regard to medical treatment for UII-related cardiac diseases, vasoconstrictor effect of UII results in cardiac hypertrophy, hypertension, and heart failure. Administration of vasodilators, including ARBs, calcium channel blockers, or UII antagonists, may reverse cardiac hypertrophy and fibrosis, heart failure, and hypertension (Boussette *et al.* 2006, Gao *et al.* 2010, Bai *et al.* 2011).

In conclusion, hypoxia in cultured rat neonatal cardiomyocytes induces the expression of AngII and ROS. Ang II, ROS, and the JNK pathway mediate the induction of UII expression with subsequent cardiomyocyte hypertrophy through increased transcriptional activity in cardiomyocytes. Cardiomyocyte hypertrophy after hypoxia can be identified by increased protein synthesis and hypertrophic changes in cardiomyocytes. Hypoxia-induced AngII expression has similar effects to that of exogenously added AngII and may increase both UII expression and transcriptional activity in cardiomyocytes. Cardiomyocyte hypertrophy can be prevented by ARB, which represses the effect of AngII and UII. Thus, UII may be a potential therapeutic target for the treatment of ischemic heart disease with the aim of preventing cardiac hypertrophy and cardiac fibrosis (Fig. 8).

Supplementary data

This is linked to the online version of the paper at <http://dx.doi.org/10.1530/JOE-13-0261>.

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.

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