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Developmental changes in AT1 and AT2 receptor-protein expression in rats

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Abstract

It has long been known that angiotensin type-1 receptors (AT1R) play a critical role in sympathetic regulation, cardiovascular activity, and hormone secretion under physiological and pathological states. On the other hand, the functional significance of angiotensin type-2 receptors (AT2R) is poorly understood. In a recent study we demonstrated that, in rats with chronic heart failure, AT1R protein expression was increased but AT2R expression was decreased in the rostral ventrolateral medulla (RVLM). This imbalance of angiotensin receptors contributed to sympathoexcitation in the heart failure state. In the current experiment, we measured AT1R and AT2R protein expressions in the brainstem, kidney and liver from male foetuses (3 days before birth), male neonates (3 days after birth), male and female adults (8 weeks) and male aged (28 months) rats by Western blot analysis. In the brainstem, we found that the foetuses and neonates exhibited a significantly lower AT2R protein expression compared with adult rats (foetus 0.08 ± 0.01 , neonate 0.12 ± 0.01 , male adult 0.25 ± 0.01 , female adult 0.22 ± 0.02 ; $n = 4$ per group, $p < 0.001$ foetus and neonate compared with male or female adults). In contrast, the foetuses and neonates expressed significantly higher AT1R protein than that of the adults (foetus 0.64 ± 0.09 , neonate 0.56 ± 0.01, male adult 0.13 ± 0.02, female adult 0.08 ± 0.02; *n* = 4 each group, *p* < 0.001 foetus and neonate compared with male and female adults). In the liver, the AT2R protein was also higher in foetus and neonate, than in adult rats. Interestingly, the foetal liver expressed higher AT1R protein compared with that of the neonate. In the kidney, AT2R expression was significantly increased with age (foetus 0.08 ± 0.01 , neonate 0.19 ± 0.02 , male adult 0.49 ± 0.04 , female adult 0.90 ± 0.10 ; $n = 4$ per group, $p < 0.01 - 0.001$). AT1R expression, on the other hand, was higher in the foetuses than that in both neonate and male adults. This study provides data contrary to existing dogma that AT2R expression is higher in foetal life and low in adults, suggesting an involvement of a potentially important functional role for AT2R in adult animals and AT1R in foetal development and/or physiology.

Keywords

angiotensin II receptors; brainstem; development; kidney; liver; protein expression

Introduction

The renin–angiotensin system (RAS) has been identified and investigated for more than 100 years.1 This system, however, is becoming more complex due to the continual discovery of newer functional components.2.3 Angiotensin II (Ang II) is the primary peptide in this

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system. This octapeptide activates two well-known G protein-coupled receptors: the Ang II type 1 receptor (AT1R) and the Ang II type 2 receptor (AT2R).4,5 The AT1R mediates the majority of classical biological functions of Ang II6,7 and plays a critical role in the control of sympathetic nerve activity, the regulation of blood pressure, water and electrolyte balance, thirst, hormone secretion and renal function.⁴ The AT1R has been implicated in a variety of pathologic conditions primarily because its effects are so widespread. Indeed, the AT1R has been regarded as a primary target for cardiovascular and renal pharmacotherapy for common diseases such as hypertension, chronic heart failure and diabetic nephropathy. Antagonists to AT1R and inhibitors of angiotensin-converting enzyme (ACE) have been routinely used to treat patients with these diseases. $8-10$

The AT2R on the other hand, has been viewed, for some time, as being involved only in development and growth, due to the observation of its ubiquitous expression at very high levels in the foetus and its rapid regression to low levels or disappearance after birth.^{4,5,11} This assertion, however, is primarily derived from studies using autoradiography, $12-14$ ligand binding15–17 and *in situ* hybridisation techniques.18–20 Notably lacking from the literature are reports of AT2R protein expression at various stages of animal development and growth. Therefore, in the present experiment, we employed Western blot analysis to measure AT1R and AT2R protein expression in brainstem, kidney and liver from Sprague– Dawley or Fisher 344 rats at various stages of development.

Methods

Animals

A total of 24 rats were used in this experiment. Sixteen Sprague–Dawley rats were purchased from SASCO (Madison, WI): four male foetuses (3 days before birth), four male neonates (3 days after birth), four male adults (8 weeks) and four female adults (8 weeks); eight Fisher 344 rats were obtained from the National Institute of Ageing: four male adults (8 weeks) and four male aged (28 months). The four foetuses were taken from four pregnant female rats, and the four neonates were taken from four litters. The sex of the foetuses and neonates was identified by sex determining region Y (SRY) expression employing reverse transcriptase polymerase chain reaction (RT-PCR; NCBI Reference Sequence: NM_001109181.1; left primer AGGGCTGGGAGAAAGAAGAG and right primer TTGCTGATCTCCGAGTTGTG). All experiments were approved by the Institutional Animal Care and Use Committee of the University of Nebraska Medical Center and were carried out under the guidelines of the American Physiological Society and the National Institutes of Health, *Guide for the Care and Use of Laboratory Animals*.

Preparation of tissues

All rats were euthanised by $CO₂$. The whole brain, kidney and liver were removed and immediately frozen on dry ice, then stored at −80°C. The samples were stored for a maximum of 2 weeks. For the brainstem samples, a rostral coronal section 0.5 mm from the obex was taken. For the kidney sample from the foetuses, the whole right kidney was used. For the kidney sample from the neonates and adults, a 0.5 mm horizontal section from the middle of the right kidney was taken, which included renal cortex, medulla, calyx and vessels. The hepatic anterior lobe was used as the liver sample.

Western Blot analysis

The tissues were homogenised in RIPA buffer (50 mM TrisHCl pH7.4, 150 mM NaCl, 2 mM EDTA, 1% NP-40, 0.1% SDS) and total protein was extracted from this homogenate. The protein concentration in each sample extract was measured using a protein assay kit (Pierce; Rockford, IL) and then was adjusted to the same value in all samples with 2X 4%

SDS sample buffer. The samples were boiled for 5 min followed by loading on a 7.5% SDS-PAGE gel (30 μg protein/10 μl per well) for electrophoresis using a Bio-Rad mini gel apparatus at 40 mA/gel for 45 min. The fractionated protein on the gel was transferred onto a PVDF membrane (Millipore) and electrophoresed at 300 mA for 90 min. The membrane was first probed with AT2R primary antibody (AT2R rabbit polyclonal IgG, sc-9040, Santa Cruz Biotechnology Inc, 1:500) and secondary antibody (goat anti-rabbit IgG-HRP, Santa Cruz, 1:2500), and then treated with enhanced chemiluminescence substrate (Pierce; Rockford, IL) for 5 min at room temperature. The bands in the membrane were visualised and analysed using UVP BioImaging Systems. After obtaining the AT2R blot density, the membrane was then treated using Restore Western Blot Stripping Buffer (Thermo Scientific) to remove the AT2R signal, followed by probing with an AT1R primary antibody (AT1R rabbit polyclonal IgG, sc-1173, Santa Cruz Biotechnology Inc, 1:500) and finally with glyceraldehyde-3-phosphate dehydrogenase (*GAPDH*) primary antibodies (GAPDH mouse monoclonal IgG, sc-32233, Santa Cruz Biotechnology Inc, 1:1000) using the same process as the AT2R antibody to get the AT1R and GAPDH blot densities. The final reported data are the normalised AT2R and AT1R band densities by GAPDH.

Statistical analyses

All data are reported as the mean \pm SEM. A one-way analysis of variance (ANOVA) was used followed by the Student–Newman–Keuls *post hoc* analysis, where appropriate. Statistical analysis was performed with the aid of SigmaStat software. *p* < 0.05 was considered statistically significant.

Results

AT2R and AT1R protein expression in the brainstem

Figure 1 shows the AT2R and AT1R expressions in brainstem tissues from foetus, neonate, male adult, and female adult Sprague–Dawley rats. The top panels show the original blots from all samples (four samples for each group) and the bottom depicts the mean data. AT2R expression in the brainstem was significantly increased with age. Foetuses exhibited the lowest AT2R protein levels $(0.08 \pm 0.01, n = 4)$ and male adult rats expressed the highest AT2R protein (0.25 ± 0.01 , $n = 4$). There were no significant differences in AT2R expression between foetus and neonate or between male and female adult tissues. Panel C shows AT1R expression in each group of rats. Mean AT1R expression in foetus (0.64 \pm 0.09, $n = 4$) and neonate $(0.56 \pm 0.01, n = 4)$ were higher than that in male $(0.13 \pm 0.02, n = 1)$ 4; $p < 0.001$ versus foetus and neonate) and female $(0.08 \pm 0.02, n = 4; p < 0.001$ versus foetus and neonate) adult tissues. However, there were no significant differences between foetus and neonate or between male and female tissues.

We also examined the AT2R and AT1R expression in the brainstem of aged rats (28 months), which is shown in figure 2. We did not find any significant difference in AT2R and AT1R protein expression between adult (8 weeks) (AT2R: 0.81 ± 0.09 , AT1R: 1.60 ± 0.41 , *n* = 4) and aged (AT2R: 1.00 ± 0.08 , AT1R: 1.32 ± 0.12 , *n* = 4) Fisher 344 rats.

In order to confirm the specificity of the AT2R antibody used in the above experiment, we employed another AT2R antibody, sc-48452 (AT2R goat polyclonal IgG, Santa Cruz Biotechnology), to repeat this experiment. We obtained similar results (Panel A of figure 3). Panel B of figure 3 shows the negative control using AT2R antibody blocking peptide, sc-48452-P (Santa Cruz Biotechnology).

AT2R and AT1R protein expressions in liver

In the liver, AT2R expression in male $(0.45 \pm 0.06, n = 4)$ and female $(0.65 \pm 0.13, n = 4)$ adults was significantly higher than that in foetal rats $(0.11 \pm 0.03, n = 4)$ and neonates $(0.19$ \pm 0.02, $n = 4$), however, there were no significant differences between foetal and neonate livers or between male and female adult livers (panel B in figure 4). In contrast, we found that foetal rats exhibited a significantly higher AT1R expression in the liver $(1.16 \pm 0.40, n$ $= 4$) compared with the other groups (neonate: 0.17 ± 0.02 , male adult: 0.10 ± 0.02 , female adult: 0.04 ± 0.01 , $n = 4$ each group). AT1R expression in foetal liver was 6.7 times that of the neonate, 11.7 times that of male adults and 28.1 times that of female adults. However, there were no differences in AT1R expression between neonate, male adult and female adult rats (Panel C in figure 4).

AT2R and AT1R protein expression in kidney

As in brainstem and liver, AT2R protein expression in the kidney also significantly increased with age. The lowest expression was in foetal rats $(0.08 \pm 0.01, n = 4)$ and the highest expression was in female adult rats $(0.90 \pm 0.10, n = 4)$ (panel B, figure 5). In renal tissue, foetal rats also had higher AT1R expression $(0.97 \pm 0.08, n = 4)$ compared with neonates $(0.64 \pm 0.06, n = 4)$ and male adult rats $(0.64 \pm 0.11, n = 4)$. There were no significant differences in AT2R expression between foetal and female adult or neonates and male kidneys (panel C, figure 5). Even though there was a tendency for female adult rats to express the highest AT1R protein level $(1.05 \pm 0.30, n = 4)$, the difference did not reach statistical significance.

Discussion

The major new finding from this study is that AT2R protein expression in the brainstem, liver, and kidney of rats increases progressively from foetal life to adulthood. It has been accepted generally that AT2R expression is high in foetal tissue, and then dramatically decreases or even disappears after birth. $6,11,21-29$ The current data clearly challenges that concept. At least in brainstem, liver and kidney tissue, adult rats exhibited a higher AT2R protein level compared with foetus or neonates.

The conventional concept on AT2R expression profiles before and after birth was derived from autoradiographical data,^{12–14} ligand binding experiments^{15–17} and *in situ* hybridisation.^{18–20} Autoradiography and ligand binding provide information on the binding and activity of a ligand to its receptor. The accuracy of these techniques depends largely on the specificity of the agonists and antagonists used. On the other hand, *in situ* hybridisation detects mRNA, and is a good method to locate target mRNA, however it is not a reliable technique to quantify gene expression. To the best of the authors' knowledge the data presented here are the first to directly determine AT2R protein expression in rats at various developmental stages. More importantly, our findings are in direct opposition to the previously reported results using other techniques.12–²⁰

The above discrepancy is not clear, but at least two potential reasons should be considered. First, we used Western blot to measure total AT2R protein, which included both cytoplasmic receptor protein (immature receptor) and plasma membrane receptor protein (mature receptor). However, autoradiography and ligand binding primarily determine mature receptors only on the cell surface. Our data might reflect differences between the total AT2R protein and the mature AT2R protein. Second, different regions, even in the same organ, may exhibit different AT2R expression level. Using competition binding Gwathmey *et al*. 30 demonstrated that AT2R was the predominant receptor subtype in the renal cortex but that AT1R was predominant in the renal medulla of both adult and foetal sheep. However, they

did not find significant differences in AT2R or AT1R binding activity between adult and foetal kidneys. In the foetal ovine brain, on the other hand, Hu *et al*. ³¹ documented a predominance of AT2R in the cerebellum but AT1R in the hypothalamus. In a recent review Mao *et al*. ³² summarised mRNA expression levels for angiotensin receptors in different regions of the foetal rat brain: AT2R mRNA predominated in some brain regions but AT1R in others.

Our present finding of lower AT2R protein expression in the foetus and neonate compared with adults suggests a re-examination of the functional significance of this receptor. Even though AT2R has been conventionally thought of as an important modulator of foetal development and growth due to the currently accepted concept that this receptor is expressed transiently and abundantly only during embryonic/foetal development, there is little evidence to support this notion. For example, AT2R knockout mice demonstrate apparently normal embryogenesis.^{33,34} These data therefore fail to support the hypothesis of an essential role of this receptor in growth and development. In contrast, the AT2R has been often documented as a predominantly negative regulator of vascular smooth-muscle-cell proliferation both *in vivo* and *in vitro* experiments.35,36 These anti-proliferative and antigrowth effects of AT2R were also observed in neurons, 37 pheochromocytoma, 38 fibroblasts, 39 renal mesangial cells, 40 coronary endothelial cells⁴¹ and *in vivo* on microvascular growth. 42 These data are consistent with our current findings that AT2R expression is lower in foetal and neonatal tissue. In contrast, compatible with the well-known growth and proliferative function of AT1R,4 7 our findings that foetal and neonatal tissue expresses a higher AT1R protein implies a potential role for this angiotensin receptor subtype in organism growth and development. Interestingly, functional data also appear to support our current finding that, in the foetal brain, the AT1R but not AT2R is the dominant angiotensin receptor subtype. For example, in the ovine foetus, intracerebroventricular (icv) injection of Ang II evoked a significant increase in mean arterial blood pressure, $43,44$ which typifies a central AT1R effect, not that of AT2R stimulation. Indeed, Shi *et al*. ⁴⁵ further demonstrated that the pressor response to Ang II in the ovine foetal brain was completely abolished by the AT1R antagonist, Losartan, but not by the AT2R blocker, PD123319. These functional data strongly suggest that, at least in the ovine foetal brain, AT1R mediates the responses to central Ang II.

It is of interest that the pattern of reciprocal changes in AT1R and AT2R during maturation occurred in all tissues examined. This phenomenon may imply a possible reciprocal inhibition of protein expression between these two angiotensin receptors during development. The underlying mechanisms are not clear, but the interaction between AT1R and AT2R at the transcriptional level has been well documented. AT1R activation enhances AT2R mRNA degradation46 and decreases AT2R mRNA accumulation.47 In addition, AT1R mRNA and protein are increased in AT2R knockout mice.48 Transfection of vascular smooth muscle cells with the AT2R gene results in a decreased expression of AT1R mRNA and protein.49 This possible reciprocal inhibition in gene expression between AT1R and AT2R has also been observed in pathologic conditions. Employing Western blot analysis, we found that in the rostral ventrolateral medulla (RVLM) of rats with chronic heart failure, AT1R protein expression was up regulated while AT2R protein expression was down regulated.50 Using RT-PCR and autoradiography, Peng and Phillips51 demonstrated an increased AT1R mRNA expression and receptor binding, but decreased AT2R mRNA and receptor binding in the hypothalamus and brainstem of rats with cold-induced hypertension.

We recognise that Western blotting is the only technique we used in this experiment, and therefore some limitations of our data should be discussed. Owing to the lack of mRNA data, we do not know whether this change in AT2R/AT1R protein occurs at the transcriptional or translation level. Because we did not determine ligand-receptor binding

activity, we are not sure if the observed AT2R/AT1R protein expression profile represents similar changes in receptor function. However, the current data raise some critical questions concerning angiotensin receptors: does the AT2R really decline after birth? What is the role of AT2R in the adult? Does the AT1R have any functional significance in foetal development and growth?

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Figure 1.

Western blot showing AT1R and AT2R protein expression in brainstem samples of male foetus, male neonate, male adult and female adult Sprague-Dawley rats. Panel A shows the original Western blot images (the upper bands are AT2R and the middle bands show AT1R) from all samples (four rats per group). Panel B shows the mean data for AT2R protein expression. Panel C shows the mean data for AT1R protein expression. ****p* < .001.

Figure 2.

Western blot showing AT1R and AT2R protein expression in brainstem samples of adult and aged Fisher 344 rats. Panel A shows the original Western blot images (the upper bands are AT2R and the middle bands are AT1R) from all samples (four rats per group). Panel B shows the mean data for AT2R protein expression. Panel C shows the mean data for AT1R protein expression.

Figure 3.

A re-evaluation of AT2R protein expression in the brainstem samples of male foetus, male neonate, male adult, and female adult Sprague-Dawley rats using a second AT2R antibody, sc-48452 (panel A) and its negative control by the blocking peptide sc48452-P (panel B).

Figure 4.

Western blot showing AT1R and AT2R protein expression in the liver of male foetus, male neonate, male adult and female adult Sprague-Dawley rats. Panel A show the original Western blot images (the upper panels are AT2R and the middle panels are AT1R) from all samples (four rats each group). Panel B show the mean data for AT2R protein expression. Panel C shows the mean data for AT1R protein expression. $\frac{*p}{<}$.05, $\frac{*p}{<}$.01, $\frac{***p}{<}$.001.

Figure 5.

Western blot showing AT1R and AT2R protein expression in the kidney of male foetus, male neonate, male adult and female adult Sprague-Dawley rats. Panel A shows the original western blot images (the upper panel shows AT2R and the middle panel shows are AT1R) from all samples (four rats each group). Panel B shows the mean data for AT2R protein expression. Panel C shows the mean data for AT1R protein expression. **p* < .05, ***p* < .01, ****p* < .001.