Short communication

THE EXPRESSION OF ENDOTHELIN TYPE A AND B RECEPTORS IN THE LATERAL WALL OF THE MOUSE COCHLEA

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Abstract: Endothelin (ET), originally characterized as a vasoconstrictive peptide, has been found to have many different biological functions, including acting as a local hormonal regulator of pressure, fluid, ions and neurotransmitters in the inner ear. The objective of this study was to examine and quantify the mRNA expression of the endothelin type A and B receptors (ETAR and ETBR) in the strial vascularies (StV) and non-strial tissues (NSt) of the cochlear lateral wall using the real-time quantitative reverse transcription-polymerase chain reaction (RT-PCR) technique. The mouse tissue samples were harvested and RNA was extracted. RT was performed to obtain cDNA, and then the mRNA expression of each gene was measured via real-time PCR. We found that both receptor subtypes were expressed in the cochlear lateral wall, with a predominance of ETAR over ETBR. We showed that the mRNA expression of the two receptor subtypes was higher in the StV with a 1.8 times higher level of ETAR and an 8.1 times higher level of ETBR mRNAs than in the adjacent NSt of the lateral wall tissue. This study shows the existence and the quantity of ET receptor subtypes in the StV and NSt of the mouse cochlea. Our results

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Abbreviations used: ANP – atrial natriuretic peptide; Ct – cycle threshold; ET – endothelin; ETAR – endothelin type A receptor; ETBR – endothelin type B receptor; NO – nitric oxide; NSt – non-strial tissues; PG – prostaglandins; RQ – relative quantity; RT-PCR – reverse transcription-polymerase chain reaction; StV – strial vascularies
suggest that an endothelin-mediated response via two different receptors, ETAR and ETBR, may play an important role in the physiological functions of the cochlear lateral wall by maintaining the homeostatic environment of the cochlea.

Key words: ETAR, ETBR, Real time quantitative RT-PCR, Cochlea

INTRODUCTION

Endothelin, which was first identified as a potent vasoconstrictive peptide secreted from cultured aortic endothelial cells [1], is now known to consist of a family of 3 mammalian vasoactive peptides that includes ET-1, ET-2 and ET-3 [2]. These ETs are synthesized and released by many different cell types. The pathophysiological actions of endothelin are mediated by two different receptors, the endothelin type A receptor (ETAR) and the endothelin type B receptor (ETBR) [3]. Both are heterogeneously distributed throughout many organs. ET-1 and ET-2 bind to ETAR with high affinity, whereas ET-3 displays extremely low binding affinities by comparison. ETBR has similar binding properties for ET-1, ET-2, and ET-3. The pathophysiological actions of ET-1 have been thoroughly investigated. The release of ET-1 from the vascular endothelial cells activates ETAR, which predominates on the underlining vascular smooth muscle cells. The binding of ET-1 with ETAR activates phospholipase C, leading to the formation of inositol 1,4,5-trisphosphate and diacylglycerol [4]. ET-1 has a different function when it binds to ETBR by promoting nitric oxide and prostacyclin release causing vasodilation in vasculum [5]. ET-1 has been reported to have a wide variety of biological functions, not only in the vascular tissues but also in non-vascular tissues [6]. Considerable evidence suggests that ET-1 is a local autocrine and paracrine factor, in addition to its role in vasoactivity [7-9].

Previous studies have demonstrated an extensive distribution of ETs in many organs, including the inner ear. ET-1 activity was detected in the modiolus, spiral ligament, stria vascularis, spiral prominence, Reissner’s membrane, supporting cells of the organ of Corti, and spiral ganglion cells [10]. In the vestibule of normal guinea pigs, ET-like activity was identified in the sensory epithelial cells, supporting cells, dark cells, transitional cells, vestibular membrane, semicircular wall cells and vestibular ganglion cells [11]. ET-1-like activity was identified in the epithelial cells and subepithelial connective tissue of the endolympathic sac and duct of normal guinea pigs [12]. Immunoreactivity for ET-1 and ET-3 was localized preferentially in cisterns of the rough endoplasmic reticulum and Golgi apparatus in the intermediate cells of WBB6F1+/+ mice. Immunoreactivity for ET-1 and ET-3 was also seen in lysosomes which were also occasionally found to contain (pre)melanosomes [13]. It was proposed that ET-1 may play an important role in the regulation of inner ear pressure, fluid volume, and ion balance.

ETAR has been demonstrated to be expressed in the plasma membranes of the margin cells in the StV via immunocytochemical analysis. It was proposed that
ETs, synthesized and released by the intermediate cells, may participate in regulating the function of Na⁺, K⁺-ATPase in the production of the endolymph in the margin cells by mediation of ETAR in a paracrine manner [10]. Recently, another study showed that ETAR was expressed in the plasma of the intermediate cells and the capillary walls of the strial vascularis. ETBR activity was detected in the endothelial cells of the capillaries of the strial vascularis in mice. ET-1 expression was found to be distributed in the margin cells in a strong positive staining, and in the intermediate cells of the strial vascularis in the spiral ligament in a weak positive staining [14].

However, the research thusfar has been limited by the use of the immunohistochemical staining method. The expression of ET receptor subtypes has not been systematically investigated in the lateral wall of the mouse cochlea. There is no available data on the quantity of ET receptors in the StV or NSt of the mouse cochlea. In order to elucidate the importance of the ET receptor in the regulation of inner ear function, it is crucial to quantify the gene expression. Real-time quantitative reverse transcription-polymerase chain reaction (RT-PCR) is a highly sensitive technology that allows high throughput quantification of gene expression [15]. The objective of this study was to determine the levels of mRNA expression of the ET receptor subtypes, ETAR and ETBR, in the strial vasculatures and non-strial tissues of the lateral wall of the mouse cochlea using real-time quantitative RT-PCR techniques. We aimed to establish whether the StV and NSt of mice synthesize ETAR and ETBR. We compared the levels of mRNA expression of ETAR and ETBR in the StV and NSt, which are known to be involved in the regulation of inner ear pressure, fluid volume, and ion balance. The side-by-side comparison of our quantitative data on ET receptor expression clearly highlights the crucial roles of these peptides. Further understanding of the roles of these peptides in the cochlear lateral wall should reveal the importance of endothelin in maintaining the homeostatic environment of the cochlea.

MATERIALS AND METHODS

Animals

For this study, we used adult (c. 30 days old) Kunming mice, an outbreed strain of laboratory animal widely utilized in related pharmaceutical and genetic studies in China. The study was approved by the Animal Care and Use Committee at the West China Medical School of Sichuan University. The temporal bone was removed and StV and NSt tissues were dissected at 4°C, as previously described [17].

RNA isolation and reverse transcription

Total RNA from the tissues of the StV and NSt in the mouse cochlea was isolated using Trizol reagent (MRC USA). RNA was reverse transcribed into cDNA using the Revert Aid™ First Strand cDNA Synthesis Kit (MBI Fermentas...
Inc.) with the addition of random hexamer primers. Briefly, 5 µl of purified RNA was used as template for cDNA synthesis in the presence of 1 µl M-MLV reverse transcriptase (200 U), 1 µl of random hexamer primer, 4 µl of 5 × reaction buffer, 2 µl of hexamer 1x (Roche), 2 µl of dNTP mix (10 mM each) and 6 µl of RNase-free water. After incubation for 60 min at 42°C, the reverse transcriptase was inactivated at 70°C for 10 min, and the cDNAs were stored at -20°C until further analysis.

**Primers**

Primers were designed according to the NCBI Genebank of mice, and were synthesized by the Shanghai SAGON. The primer sequences for RT-PCR were as follows: ETAR, forward primer 5’-CTCCATCTGGATTCTTTTCCTT-3’ and reverse primer 5’-CTTGGTAAAACTCCATGAACT-3’ (a 137-bp product was to be generated); ETBR, forward primer 5’-CAAGTTGCTCGCAGAGGACT-3’ and reverse primer 5’-CAGCTCGATATCTGTCAATACT-3’ (a 128-bp product was to be synthesized); GAPDH, forward primer 5’-TGGGTGTGAACCAGAA-3’ and reverse primer 5’-GGCATGGACTGTGGTCATGA-3’ (a 143-bp product was to be generated).

**Real-time quantitative PCR**

Real-time PCR was used to determine the gene expression profiles of the ET receptor subtypes. cDNAs were amplified by real-time PCR using an FTC-2000 (FUNGLYN, CANADA). Each analysis was performed in a total volume of 30 µl reaction mixture containing 5 µl cDNA sample, 1 µl SYBR Green I, and 2 µl gene-specific forward and reverse primers (10 µM each). Housekeeping genes were included to normalize the data. Amplifications were performed as follows, 45 cycles at 94°C for 2min, 94°C for 20 sec and 54°C for 20 sec, with a final extension at 70°C for 30 sec and at 80°C for 20 sec.

The cycle threshold (Ct) number, defined as the number of PCR amplification cycles required to reach fluorescent intensity above the threshold, was determined for each gene and each developmental time point analyzed. Using serial dilutions of the test sample cDNA, the standard curve was generated on the basis of the linear relationship of existing Ct and the logarithm of the copy number. The slope of the curve was shown to be -3.70, and a strong linear relationship was demonstrated ($R^2 = 1.00$; Fig. 1).

**Analysis**

Samples were normalized internally using the Ct number of the housekeeping gene GAPDH, as follows: $\Delta C_t$ (sample) = (Ct sample) - (Ct GAPDH). The mean Ct of ETBR RNA from the heart was set to a relative quantity (RQ) value of 1 using the $\Delta \Delta C_t$, calculated as follows: $\Delta \Delta C_t$ (sample) = $\Delta C_t$ (sample) - $\Delta C_t$ (Heart), and RQ = $2^{-\Delta \Delta C_t}$ [16]. The data is reported as means ± standard errors (SE). The one-way analysis of variance, ANOVA, was performed using SPSS 11.0 statistical software. P values of less than 0.05 were considered to be statistically significant.
RESULTS AND DISCUSSION

Via real-time quantitative RT-PCR, we demonstrated that both ETAR and ETBR mRNAs were expressed in the cochlear lateral wall (Fig. 2). We used heart tissue samples, which were previously shown to express ETAR and ETBR, as positive controls. GAPDH was examined as a housekeeping gene.

The results of the analysis of ETAR and ETBR gene expression from the StV and NSt are presented in Fig. 3. ETAR mRNA expression predominated in the StV with a relatively lower level of expression of the ETBR receptor mRNA in the StV. The mRNA levels of ETAR and ETBR were relatively low in the NSt as compared to those in the StV; there was a 1.8 times higher amount of ETAR mRNA in the StV than in the NSt of the lateral wall tissue (p < 0.05), and 8.1 times as much ETBR mRNA in the StV as in the NSt (p < 0.05).

In this study, we demonstrated the mRNA expression of the ET receptor subtypes, ETAR and ETBR, in the cochlear lateral wall. Real-time quantitative RT-PCR provides a specific, sensitive and reliable method for the quantification of mRNA expression in the cochlear lateral wall of mice.

The cochlear lateral wall, consisting of the StV and adjacent NSt tissues, is bound by several kinds of parasensory epithelium. These parasensory epithelium cells play an important role in the active ion transportation of K⁺ [18]. The strial vascularis is largely responsible for generating the high K⁺ level in the endolymph, used by neurosensory cells for mechanoelectrical transduction.
Fig. 2. ETAR and ETBR cDNAs amplified from homogenates of the StV, NSt and heart. The amplified fragments were of the expected sizes of 137 and 128 bp, respectively for the ETAR (A) and ETBR gene (B). The amplification of a 143 bp fragment of GAPDH cDNA as a housekeeping control is also shown (C). M – marker, lane 1 – heart, lane 2 – strial vasculatures (StV) of the cochlear lateral wall, lane 3 – non-strial tissues (NSt) of the cochlear lateral wall.

Fig. 3. The mRNA expression of ETAR and ETBR by real-time quantitative RT-PCR. The quantification of ETAR and ETBR gene expression in the StV and NSt (n = 7) relative to the heart (n = 9). The mRNA expression of ERAR and ETBR was normalized to the expression of ETBR in the heart (control), which was arbitrarily set to a value of 1. Significantly different expression (p < 0.05) is expressed as the average ± SD.

In previous studies, the localization of endothelins was observed in the cochlea of normal guinea pigs [10, 19]. Endothelins are reported to act as modulatory peptides, possibly affecting NO, PG and ANP levels in the lateral cochlear wall [19]. Endothelins may regulate ion transport and influence the production of endocochlear potential. It was proposed that ET-1 and/or ET-3 may play a crucial role in the activation of the sodium pump in the plasma membrane of the strial marginal cells by mediating ETAR in a paracrine fashion. In addition, the immunoreactivities of ETAR along the plasma membrane of the strial middle
cells suggest the possibility of their autocrine action on the middle cells [13]. Jinnouchi et al. [10] further investigated the dynamic changes of ET-1 in the epithelium of the endolymphatic sac of guinea pigs and their relationship to the development of endolymphatic hydrops. Their findings suggest that ET-1 may play an important role as one of the regulators maintaining the fluid balance. Here, our results confirm that ETAR mRNA is present in the StV and NSt of the mouse cochlea. This concurs with earlier results obtained through immunoreactivity studies, suggesting that the StV and NSt probably actively synthesize the peptide ETAR.

ET-1 mediates its effect by acting on either the ETAR or ETBR receptor subtypes. The mRNA of both receptor subtypes is found in the StV and NSt, with a predominance of ETAR over ETBR. ET-1 may reduce its physiological functions in an autocrine and/or a paracrine manner, as has been shown for other cell types [3]. This study demonstrates that ETAR is the predominant endothelin receptor expressed in the StV and NSt. In addition, our studies demonstrate higher expression levels of ETAR and ETBR in the StV than those in the NSt. It was reported that the endothelins are involved in the activation of the sodium pump of the strial marginal cells by mediation of ETAR [13]. In addition, ETAR mediates vasoconstriction of capillaries in the spiral ligament [20]. In this study, we also determined the expression of ETBR in the cochlear lateral wall of mice. However, the function of ETBR in the StV is uncertain. ET-1 has a different function when it binds to ETBR by promoting nitric oxide and prostacyclin release causing vasodilation in the vascellum [21]. Several studies have demonstrated that ETBR also helps to clear ET-1 from the circulation, via a ligand-receptor internalization mechanism [22]. It was suggested that ET plays an important role in regulating the circulation of the cochlea as a local hormone binding to the ETBR of capillary endotheliocytes. In this study, we further demonstrated the presence of ETBR mRNA in the NSt, implying that ETBR is possibly synthesized in these cells.

In summary, we demonstrated that ETAR and ETBR were both expressed in the StV and NSt. We further obtained data suggesting that ETAR and ETBR receptors were heterogeneously distributed in the mouse cochlear lateral wall. Gene expression of the ETAR and ETBR was predominant in the StV over the NSt. This study provides evidence in support of the action of ET in the cochlear lateral wall. In particular, the high expression of these genes in the StV implies an important role in the homeostasis within the microenvironment of the inner ear.

In conclusion, the systematic analysis of ET receptor expression reported here will be helpful in future studies investigating the mechanism of endothelins action. Real-time quantitative RT-PCR provides a specific and sensitive method for the detection and quantification of ETAR and ETBR mRNA. Further studies on the significance of ET receptor-mediated signaling in ET-regulated cochlear function are required.
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