Norepinephrine transporter (NET) is expressed in cardiac sympathetic ganglia of adult rat

LI HE^{1, 2}, SUN KAI MA¹, XIAO PING HU³, GOU YUAN ZHANG², JIAN FEI^{1,*}

1 Institute of Biochemistry and Cell Biology, Shanghai Institutes for Biological Sciences, Chinese Academy of Sciences, Shanghai 200031, China

2 Cardiovascular Division of Changzheng Hospital, Shanghai 200003, China E mail: h-li@hotmail.com 3Pathological Division of Pu Nan Hospital, Shanghai 200125, China

ABSTRACT

The sympathetic nervous system plays a cardinal role in regulating cardiac function through releasing the neurotransmitter norepinephrine (NE). In comparison with central nervous system, the molecular mechanism of NE uptake in myocardium is not clear. In present study, we proved that in rat the CNS type of NE transporter (NET) was also expressed in middle cervical-stellate ganglion complex (MC-SG complex) which is considered to control the activity of heart, but not expressed in myocardium. The results also showed that NET expression level in right ganglion was significantly higher than in the left, rendering the greater capacity of NE uptake in right ventricle, a fact which may contribute to the maintenance of right ventricular function under pathologic state.

Key words: Norepinephrine transporter; gene expression; cardiac sympathetic ganglion, rat.

INTRODUCTION

The sympathetic nervous system (SNS) plays an important role in regulating cardiac function in both health and disease through releasing neurotransmitter norepinephrine (NE). In central nervous system (CNS), the neurotransmission of NE is terminated through reuptake of released neurotransmitter by Na⁺, Cl⁻ dependent norepinephrine transporter (NET) on pre-synaptic membrane[1], [2]. Although the NE uptake was well studied in heart, but the molecular basis for that is still less clear[3-5]. Ungerer et.al indicated that they could not detect any specific mRNA of NET in the myocardium investigated [6]. They postulated that somehow norepinephrine transporter protein might be synthesized in neuronal cell situated away from the heart and then transported into the cardiac nerve ending via axon transport. In this study, we proved that NET is expressed in middle cervical-stellate ganglion complex (MC-SG complex) of rat and also showed that different NET expression levels were existed between left and right ganglion.

MATERIALS AND METHODS

Animals

Male Sprague-Dawley rats at 8 w of age and weighing 180-200 g were used. After acclimatization period rats were sacrificed by decapitation and isolated the MC-SG complex of both side and left ventricular myocardium quickly. They were immediately frozen in liquid nitrogen and stored at -70oC for use.

RNA extraction and cDNA preparation

Total RNA was extracted from the MC-SG complexes and left ventricular myocardium using TRIzol (GIBCO). After the treatment of final RNA aliquots with DNase I to eliminate contamination of genomic DNA, 1 μ g of total RNA was used for reverse transcription (RT) into cDNA with M-MLV (GIBCO) according to the manufacturer using random primer.

^{*} Corresponding author, E mail: feijian@sunm.shenc.ac.cn Received May-30-2001 Revised Aug-15-2001 Accepted Aug-25-2001

PCR cloning of rat NET

PCR primers for rat NET are as follows: 5'-CTCAA GGAGG CCACG GTATG GATCG-3' (for the sense primer, gene bank accession number: Y13223.1, the sequence position is from 904 to 928) and 5'-ACCTG GAAGT CATCA GCCAG TCCGG-3' (for the antisense primer, gene bank accession number: Y13223. 1, from 1286 to 1310). The PCR reaction was performed as 35 cycles at 94°C for 1 min, 65°C for 1 min, and 72°C for 3 min. The products were analyzed on 1.8% agarose gel. The DNA band was purified from gel and then cloned into pBluscript KS and sequenced by the dideoxynucleotide method.

For quantitative analysis of NET gene expression by RT-PCR, the expression of β -actin was used as the reference for normalization. The primers for RT-PCR of β -actin are 5' - TCATG AAGTG TGACG TTGAC ATCCG TAAAG-3' (for the sense primer, gene bank accession number: V01217.1, sequence position is from 2727 to 2756) and 5'-CCTAG AAGCA TTTGC GGTGC ACGAT GGAGG-3' (for the antisence primer, from 3106 to3135).

Northern blot analysis of NET

The PCR product of NET was labeled with ³²p-dATP and used as probe for Northern blot. Twelve mg of total RNA from the ganglia (n =5) and myocardium of LV(n=5) were run on 1% agarose-formaldehyde gel and transferred to a Hybond-N membrane (Amersham Co). Pre-hybridization and hybridization were carried out according to standard method with the probe of NET. After hybridization, the membrane was washed twice in 2×SSC at room temperature, once in 1×SSC at 42°C for 20 min, then 20 min in 0.1×SSC at 65°C. After autoradiography exposure, the membranes were stripped and re-hybridized with 32P labeled b- actin cDNA probe. Hybridization signal of specific band of NET was measured by densitometry, and was normalized by the corresponding hybridization signal of β actin to correct the differences in loading and transfer.

RESULTS AND DISCUSSION

Neurotransmitter transporters play an important role in controlling of neural signal transmission, dysfunction in expression of neurotransmitter transporter genes may cause different diseases[7-9]. To determine if there is NET expressed in cardiac sympathetic system, RT-PCR was carried out from total RNA of rat myocardium and cardiac sympathetic ganglia. Fig 1 showed that PCR product in expected size was obtained from MC-SG amplified with the primers of NET. The product was gel purified and cloned and partially sequenced. It was identical to the expected sequence in NET (sequence data not shown). In contrast, no specific band could be amplified from rat myocardium (n=3 each), although the b-actin as the control was equally present in all cDNAs.

To determine whether there is difference in NET mRNA expression in left and right MC-SG, Northern blot was performed. Fig 2 showed that NET mRNA level in the right was 2.2 fold higher than that in the left (p=0.01, n=5).

SNS regulates the function of innervated effector organs through releasing neurotransmitter NE [10]. The released transmitter in the cleft binds receptors on the postsynaptic membrane through diffusion, by which the signal is transmitted. After that, NE must be rapidly eliminated from the synaptic cleft. In CNS, this is done by plasma membrane protein of sodium-dependent NE transporter (NET) which has been cloned before[11]. However, the NE uptake mechanism in cardiac system is not clear. Recently Ungerer and co-workers reported that they could not detect NET mRNA expressed in the myocardium. Therefore, it was assumed that







Fig 2. NET mRNA levels in right(R) and left(L) cardiac sympathetic ganglia determined by Northern hybridization analysis. The ratios are expressed as ratio to beta actin mRNA. P <0.05, NET mRNA levels in the right ganglia vs. the left. No detectable NET expression in myocardium of left ventricle (M).

NET mRNA might be expressed, if there is any, in the neuronal cell soma, which is located a considerable distance from the heart[6]. Subsequently, Matsuoka et al also showed high levels of NET mRNA in the cultured superior cervical ganglia (SCG)[12]. Following their works, we investigated the NET expression in cardiac sympathetic nervous system, the experiments clearly showed that NET mRNA expression is present in the MC-SG complex but not in the myocardium. This result supports the contention that NE uptake system in myocardium is formed by NET which is transported from a distance from the heart and also indicates that SNS in heart shares the same mechanism of NE concentration regulation with CNS .This information may be useful for development of new drugs targeting NET to treatment of heart disease.

The cardiac sympathetic ganglia of rat are MC-SG. This was proved by retrograde tracer experiments. The innervation is primarily ipsilateral(approximately 60%) with significant contralateral contributions[5], [13]. Previous studies still suggested that right and left cardiac sympathetic nerves may have different and specific effects on cardaic excitability[14], [15]. The postganglionic fibers from the right stellate ganglia(RSG) appear to play more important role than those from the left stellate ganglia (LSG) in maintaining contractile force, especially heart rate,. The apparent attenuation of heartbeat irregularity was observed when right stellate ganglia (RSG) innervation is interrupted, but the left stellate ganglionectomy (LSG), which is considered to affect cardiac inotropism rather than chronotropism, had no effect on cardiac interval irregularity[16]. MIBG (metaiodobenzylguanidine) is thought to share the same uptake, storage, and release mechanisms as NE in the adrenergic nerve terminals^[17]. Nozawa and colleagues reported that MIBG uptake and accumulation was greater in the right ventricular myocardium than in the left in rats[3]. In our experiment, it was shown that the NET expression is higher in right cardiac sympathetic ganglia of adult rats. This result provides a molecular biological evidence to explain the difference between RSG and LSG and for the greater capacity of NE uptake of the right vs the left myocardium.

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