



The gender-specific expression of neuropeptide Y and neuropeptide Y receptors in human atrial tissue during cardiopulmonary bypass surgery

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Background: Cardiac sympathetic nervous system is usually activated in cardiopulmonary bypass (CPB) surgery, accompanied by excessive release of norepinephrine (NE). Neuropeptide Y (NPY) has been shown to regulate NE release in the terminal of sympathetic fiber, which is a target for regulating heart function. The expression of NPY and NPY receptor (NPYR) genes in the human atrial tissues during CPB in cardiac surgery was investigated in the present study.

Methods: A few discarded atrial tissues before and after CPB were collected in 22 patients with rheumatic cardiac valve diseases. The transcriptional levels of NPY and NPYRs were monitored by real-time quantitative polymerase chain reaction (RT-qPCR) method. Moreover, the correlation between the mRNA levels of NPY/NPYRs and the clinical data were investigated in detail.

Results: The mRNA levels of NPY Y1 and NPY Y5 genes were statistically attenuated in male patients after CPB. Conversely, the expression of NPY, NPY Y1 and NPY Y5 genes were enhanced in female patients. Correlation analysis suggested that there was a significant negative correlation between cardiac ejection fraction (EF) after CPB with the atrial transcriptional level of NPY in male patients.

Conclusions: These results suggested that the expression of NPY/NPYRs in human atrial tissue during CPB was gender specific and activated NPY signaling was only identified in female patients. The elevated expression level of NPY in male patients was correlated with lower cardiac EF after CPB.

Keywords: Neuropeptide Y (NPY); NPY receptor (NPYR); cardiopulmonary bypass (CPB) surgery

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Introduction

Myocardial ischemia-reperfusion (I/R) injury is a well-recognized pathophysiological process, which occurs in several clinical conditions such as infarction, shock, organ transplantation and cardiopulmonary bypass (CPB) surgery (1). Several pathogenetic mechanisms involved have been explored. Among them, the imbalance between sympathetic and parasympathetic system inputs to the

heart, is one of the major reasons which induce myocardial ischemia, arrhythmia and death (2).

Neuropeptide Y (NPY) is a 36-residue peptide amide, co-stored and co-released with noradrenaline in sympathetic nerve fibers of postganglionic neurons innervating the cardiovascular system. NPY is associated with hypertension (3), myocardial infarction (4), cardiomyopathy (5), suggesting that NPY functions in cardiovascular homeostasis. To

Table 1 Clinical characteristics of patients

Characteristic	All	Male	Female	P value
Number	22	9	13	
Age	51.77±2.597	51.78±4.245	51.77±3.414	0.999
EF before CPB	63.14±1.282	62.00±1.708	64.00±1.867	0.454
LVIDd before CPB	49.71±1.890	51.22±3.483	48.58±2.116	0.504
LVIDs before CPB	33.00±1.336	34.11±2.406	32.17±1.542	0.486
Total CPB time (m)	89.45±8.522	92.33±12.501	87.46±11.922	0.786
Ischemic time (m)	62.27±5.868	66.56±11.572	59.31±6.194	0.557
Reperfusion time (m)	27.18±5.175	25.78±2.778	28.15±8.693	0.828
EF after CPB	59.36±1.754	56.00±2.205	61.69±2.400	0.112
LVIDd after CPB	45.41±1.372	48.78±1.722	43.08±1.759	0.038
LVIDs after CPB	31.00±1.208	33.78±1.470	29.08±1.603	0.053

Data are presented as n or mean ± SEM. EF, ejection fraction; CPB, cardiopulmonary bypass; LVIDd, left ventricular internal diastolic diameter; LVIDs, left ventricular internal systolic diameter, SEM, standard error of the mean.

date, six NPY receptors (NPYRs), namely NPYRs Y1–Y6 have been reported (6,7). NPYRs regulate the biological processes through diverse ways. NPY Y1 promoted cellular shortening and calcium transient in rat ventricular myocytes (8). Antagonists of NPY Y2 could prevent the inhibition of vagal bradycardia after sympathetic stimulation (9). The influence of vagal neurotransmission and ventricular myocyte excitability was accomplished even in the presence of β -receptor blockade, which indicated that pharmacologically targeting NPYRs might be a useful therapeutic strategy synergistically with β -receptor blockers (10). Moreover, the expression of NPY was increased in human plasma after CPB (11), implying NPY signaling controlled the pathological progression of CPB. However, the expression changes of NPY and NPYRs in human heart itself during CPB surgery remain elusive. Whether the expression of NPY and NPYRs have any correlations with clinical data needs to be investigated.

Methods

Collection of specimens

Patients were hospitalized for cardiac surgery in Anzhen Hospital, Beijing. According to preoperative Doppler echocardiography, rheumatic mitral stenosis or insufficiency were found in these patients, partly with tricuspid insufficiency. Patients with cerebrovascular accidents,

diabetes mellitus, myocardial infarction and other diseases were excluded. All researches involving human participants were approved by Medical Ethics Committee. Written informed consent was obtained from the patients and data were analyzed anonymously. Patient characteristics were summarized in *Table 1*. All patients underwent superior vena cava catheterization in the right atrial appendage. A few specimens were taken from the right atrium during the intubation of the superior vena cava (before CPB) and the extubation of the superior vena cava (after CPB), which were routinely excised and discarded tissue. The specimens were stored in RNA stabilization reagent (Qiagen) for later analysis.

Total RNA extraction of patients' atrium

Trizol was used to extract the total RNAs from patients' atrium. Trizol (sigma) was added in patients' atrial tissue for lysis, then chloroform and isopropanol were used for purification, finally the supernatant of centrifugation was dissolved in water with RNase free, and stored at -80°C .

Reverse transcription of mRNA

The total mRNA from patients' atrium was reversely transcribed according to SuperScript™ III RT kit (Invitrogen). Appropriate amount of freshly extracted RNAs with Primer, dNTP mix, SuperScript III RT were used.

Table 2 qPCR primers

Primer name	Primer sequence
Human NPY	Forward: AAAACGATCCAGCCCAGAG
	Reverse: GCTGAAAATAGGAAAAGGCCAG
Human NPY Y1	Forward: CAAGCCCAGTCGCATTAAAA
	Reverse: CAGGTAATCAAAGTATGTTGCAGG
Human NPY Y2	Forward: ACATCTTGTTCCGCGTCTC
	Reverse: CCCATTTTCAGTACAGGTCCAC
Human NPY Y5	Forward: CCGAGGTCTGCTCATTGTG
	Reverse: TCTAACTTGTGGCAGGTCAG
Human GAPDH	Forward: TGGGTGTGAACCATGAGAAG
	Reverse: GAGTCCTCCACGATACCAAAG

qPCR, quantitative polymerase chain reaction; NPY, neuropeptide Y.

Table 3 The relative mRNA levels of NPY, NPY Y1 and NPY Y5 genes in patients

Gene	All	Male	Female
NPY	1.0950 (4.99)	0.5500 (1.07)	2.1400 (10.11)
NPY Y1	1.0300 (3.19)	0.3700 (0.56)	2.5100 (6.17)
NPY Y5	1.2250 (9.70)	0.4300 (0.32)	6.6800 (20.68)

NPY, neuropeptide Y.

The resulting cDNAs were preserved at -20°C .

Real-time quantitative polymerase chain reaction (RT-qPCR)

RT-qPCR was carried out with SYBR Green (ABI). The total reaction system was 20 μL , 50°C 2 min, 95°C 10 min, 40 cycles for 95°C 15 sec, 60°C 1 min. GAPDH was set as the internal parameter and the relative mRNA levels were calculated with the $2^{-\Delta\Delta\text{Ct}}$ method. The primers were listed in *Table 2*.

Statistical analysis

Results were presented as means \pm standard error of the mean (SEM) or median (interquartile range). Data were evaluated by one-way ANOVA or nonparametric tests. A value of $P < 0.05$ was considered statistically significant.

Results

General information

Twenty-two patients hospitalized for cardiac surgery due to rheumatic heart disease were randomly enrolled. These patients received cardiac valve replacement surgery by CPB. The mean age of all patients was 51.77 ± 2.597 , total CPB time was 89.45 ± 8.522 min, ischemic time was 62.27 ± 5.868 min, reperfusion time was 27.18 ± 5.175 min. There was no statistically difference between male and female patients in such general information. The cardiac functions such as ejection fraction (EF), left ventricular internal diastolic diameter (LVIDd), left ventricular internal systolic diameter (LVIDs) were detected by echocardiography before and after CPB. After surgery, the patients' cardiac functions were partly recovered, such as LVIDd returned to normal range (*Table 1*).

The expression of NPY and NPYRs in human atrium before and after CPB surgery

Using RT-qPCR, the transcriptional levels of NPY and NPY receptor genes were analyzed. We found NPY and NPY Y1, Y2, Y5 genes could express in human atrium, of them, the transcriptional level of NPY Y2 was the least. The mRNA levels of NPY, NPY Y1 and NPY Y5 were compared before and after CPB in each patient (*Table 3*). For all 22 patients, the relative mRNA levels of NPY, NPY Y1, NPY Y5 were 1.0950 (4.99) ($P=0.236$), 1.0300 (3.19) ($P=0.069$) and 1.2250 (9.70) ($P=0.149$) respectively. There was no statistically difference before and after CPB in all patients. But the relative mRNA levels of NPY Y1 [0.3700 (0.56), $P=0.021$] and NPY Y5 [0.4300 (0.32), $P=0.011$] were statistically down-regulated in male patients after CPB (*Figure 1*). On the contrary, the relative transcriptional levels of NPY [2.1400 (10.11), $P=0.039$], NPY Y1 [2.5100 (6.17), $P=0.019$] and NPY Y5 [6.6800 (20.68), $P=0.004$] were statistically up-regulated in female patients after CPB (*Figure 2*). These results implied that NPY and NPYRs might play a role in CPB surgery, although the mechanism could be different between male and female patients.

Correlation analysis

The correlation between patients' clinical cardiac functions (LVIDd, LVIDs, EF before and after CPB), CPB

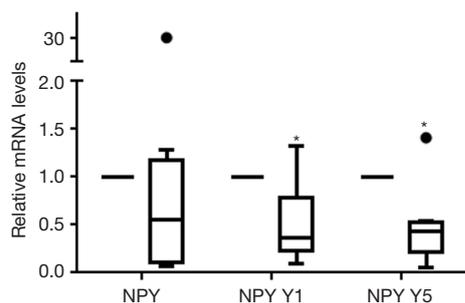


Figure 1 The transcriptional levels of NPY, NPY Y1 and NPY Y5 genes after CPB were compared with before CPB in male patients. Using qPCR, the relative mRNA levels of NPY Y1 and NPY Y5 were statistically down-regulated after CPB. *, $P < 0.05$. NPY, neuropeptide Y; CPB, cardiopulmonary bypass; qPCR, quantitative polymerase chain reaction.

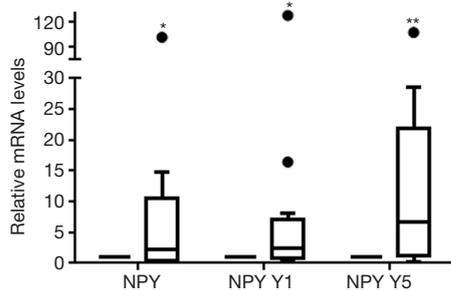


Figure 2 The transcriptional levels of NPY, NPY Y1 and NPY Y5 genes after CPB were compared with before CPB in female patients. Using qPCR, the relative mRNA levels of NPY, NPY Y1 and NPY Y5 were statistically up-regulated after CPB. *, $P < 0.05$, **, $P < 0.01$. NPY, neuropeptide Y; CPB, cardiopulmonary bypass; qPCR, quantitative polymerase chain reaction.

surgery time (ischemia time, reperfusion time), and the transcriptional levels of NPY, NPY Y1, NPY Y5 were all analyzed. There was only a significant negative correlation between cardiac EF after CPB and the relative mRNA levels of NPY (Spearman's correlation coefficient = -0.706 , $P = 0.034$) in male patients. There was no correlation in other factors.

Discussion

CPB needs to block the cardiac blood flow first, myocardial oxygenation is restored after myocardial ischemia for a short period, and ischemia reperfusion injury happens simultaneously (1). In this process, the cardiac function is

regulated by the autonomic nervous system. It is generally accepted that the activated sympathetic system and the suppressed parasympathetic system contribute to the increased blood pressure, myocardial oxygen consumption and cerebrovascular accidents (2). The sympathetic system mainly releases norepinephrine (NE) and activated β adrenergic receptors to increase heart rate and myocardial contraction. Antagonistically, the parasympathetic system release acetylcholine (ACh) and activated muscarinic ACh receptors on the opposite way.

It has been proved that NPY regulates NE release in the terminal of sympathetic fibers. The NPY signaling plays comprehensive roles in all typed cardiac cells. In human right ventricular endocardial endothelial cells, NPY Y1 modulates the cytosolic and nuclear calcium homeostasis, and their secretion of NPY (12). In cardiomyocytes, NPY had been shown to stimulate hypertrophy of rat and chick cardiomyocytes through activation of NPY Y1 receptor (13,14). Pellieux *et al.* found that NPY via NPY Y5 receptor participated in the development of mouse cardiac hypertrophy through MAPK cascade (15). These data suggest NPY Y1 and Y5 have positive chronotropic effects on ventricular cardiomyocytes. With respect to NPY Y2 receptor, it reduces ACh release and vagal bradycardia after sympathetic stimulation (16). These data mentioned above suggest that pharmacologically targeting NPYRs might conduce to cardiovascular homeostasis.

To our knowledge, this is the first reports that the transcriptional levels of NPY and NPYRs genes were investigated in atrial samples from patients with rheumatic diseases before and after CPB surgery. Using qPCR, we found that NPY, NPY Y1, NPY Y2 and NPY Y5 were all expressed in human right atrium, which was consistent with Ejaz's study (17). But the NPY Y2 transcriptional level was so hard to detect in our patients. Since NPY Y2 receptor was thought to be a presynaptic receptor (18), its transcriptional level was little.

In all 22 patients, there was no statistically difference between the transcriptional levels of NPY, NPY Y1 and NPY Y5 before and after CPB. In spite of the sex factor, our data revealed that NPY mRNA level was up-regulated after CPB in female patients. It had been reported that patient plasma NPY concentration was elevated after CPB since cardiac sympathetic nerve activity was increased after surgery (19). In this study the up-regulation of NPY mRNA in female patients in atrium might be attributed to the elevated NPY levels in plasma. In addition, we found that, only four people had atrial fibrillation after CPB in

our 22 patients, and interestingly they were all females. Guler group demonstrated that NPY played a role in the occurrence of atrial fibrillation in the patients after coronary bypass surgery (11). Hence, the atrial fibrillation in female patients might be associated with up-regulation of NPY expression. Since the number of atrial fibrillation patients was small, this manifestation needed to be investigated further. Though there was no statistically difference between the transcriptional levels of NPY in male patients before and after CPB, we found that the relative mRNA level of NPY was negative correlated with cardiac EF after CPB in these patients. Luo *et al.* reported that there was a dose-dependent stimulation of NPY with decreased ATP content and activity in the rat cardiomyocytes (20). Thus, our findings reinforced that the decreased level of NPY mRNA in male atriums might help the patients to recover cardiac functions after surgery.

In this study, the transcriptional levels of NPY Y1 and NPY Y5 were both up-regulated in female patients, but down-regulated in male patients after CPB, which suggested that NPY signaling was activated in female patients. Jackson *et al.* found that NPY Y1 expression level was higher in the male rat skeletal muscle compared with that in females (21). In our atrium muscle tissue, the expression levels of NPY Y1 and NPY Y5 were also stronger in male patients than that of female patients. Musso *et al.* found that estradiol could stimulate NPY Y1 transcription through estrogen receptor alpha (ER- α) (22). However, we could not find significant difference in the mRNA levels of ER- α in female patients before and after CPB (data not shown). Since ER- α moved from cytoplasm to nucleus to regulate target genes (23), the up-regulation of NPY Y1 expression in female patients might be due to the cytoplasm-nucleus translocation. NPY and its receptors had been found gender-specificity in other disease, such as type 2 diabetes (24) and severe periodontitis (25). Since targeting NPYRs pharmacologically may be a useful therapeutic strategy synergistically with β -blockers for cardiovascular diseases, the gender-specificity should be taken into account.

Conclusions

Our results suggested that there was gender-specificity in the expression of NPY and NPY receptor genes during CPB surgery and NPY signaling was activated in female patients. Gender differences should be considered when using NPYRs as pharmacological targets for the treatment of cardiovascular diseases.

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Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

Ethical Statement: The study was approved by the ethics review board of the Capital Medical University affiliated Beijing Anzhen Hospital (No. 2009001X) and written informed consent was obtained from all patients.

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