Characteristic Expressions of GABA Receptors and GABA Producing/Transporting Molecules in Rat Kidney

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Abstract

Gamma-aminobutyric acid (GABA) is an important neurotransmitter, but recent reports have revealed the expression of GABAAergic components in peripheral, non-neural tissues. GABA administration induces natriuresis and lowers blood pressure, suggesting renal GABA targets. However, systematic evaluation of renal GABAAergic components has not been reported. In this study, kidney cortices of Wistar-Kyoto rats (WKY) were used to assay for messenger RNAs of GABA-related molecules using RT-PCR. In WKY kidney cortex, GABA_A receptor subunits, α1, β3, δ, ε, and π1, in addition to both types of GABA_B receptors, R1 and R2, and GABA_C receptor π1 and π2 subunit mRNAs were detected. Kidney cortex also expressed mRNAs of glutamate decarboxylase (GAD) 65, GAD67, 4-aminobutyrate aminotransferase and GABA transporter, GAT2. Western blot and/or immunohistochemistry were performed for those molecules detected by RT-PCR. By immunofluorescent observation, co-staining of α1, β3, and π1 subunits was observed mainly on the apical side of cortical tubules, and immunoblot of kidney protein precipitated with π1 subunit antibody revealed α1 and β3 subunit co-assembly. This is the first report of GABA_A receptor π1 subunit in the kidney. In summary, unique set of GABA receptor subunits and subtypes were found in rat kidney cortex. As GABA producing enzymes, transporters and degrading enzyme were also detected, a possible existence of local renal GABAergic system with an autocrine/paracrine mechanism is suggested.

Introduction

Gamma-aminobutyric acid (GABA) acts as a major inhibitory neurotransmitter in the central nervous system. GABA is synthesized in the brain from glutamic acid via glutamic acid decarboxylase (GAD). There are two types of GAD with different molecular weights, GAD67 and GAD65. Released GABA acts on two general classes of its receptors, GABA_A/GABA_C and GABA_B. GABA_A and GABA_C receptors are ligand-gated chloride channels composed of a combination of various subunits as pentamers. The unique sensitivity of GABA_A receptor to its agonist such as benzodiazepines is known to depend on the combination of subunits including α, β, γ, δ, ε, π, and θ [1,2]. Meanwhile, GABA_B is a metabotropic G-protein coupled receptor coupling with Gi/o. GABA_B receptor controls neuronal activity via activation of K⁺ channels, inhibition of Gαq channels, or both. Similar to the GABA_A receptor-mediated increase in the conductance of Cl⁻, the activation of K⁺ channels from the GABA_A signaling induces hyperpolarization of the cell membrane, which produces chronic stabilization of neuronal activity [3]. GABA secreted from neurons subsequently undergoes re-uptake via GABA transporters (GAT) and is metabolized by the GABA degrading enzyme, 4-aminobutyrate aminotransferase (ABAT) [4].

Considerable evidence suggests that GABA also mediates diverse responses in non-neuronal tissues. GABA_A receptor subunits have been identified in tissues such as the lung [5], retina [6], uterus [7,8], spermatocytes [9], adrenal medulla [10], pituitary lobes [11], and pancreatic β-cells [12]. Mizuta et al. [13] detected functional GABA_A receptors in airway epithelium and airway epithelial cell lines.

The kidney is a key organ responsible for blood pressure regulation by controlling water and sodium balance. It is previously reported that GABA causes natriuresis and lowers blood pressure in human as well as rodents [14,15]. In addition, GABA_A receptor agonist muscimol and GABA_B agonist baclofen increased fractional excretion of water and sodium in isolated rat kidney perfusion [16]. These suggest that GABA may play a pivotal role in the renal pathophysiology of hypertension. To date, there are some reports of the presence of GABAAergic system in the kidney. GABA immunoreactivity was found in the renal medulla and cortex but not in neuronal structures [17], and depolarizing stimuli such as ouabain increased GABA release from rat kidney slice [18]. Sarang et al. examined the expression of GABA_A receptor α1, α2, α3, α5, β1, γ1 and δ subunits in the kidney cortex from Sprague-Dawley rats and found the expressions of α1, α5, β1, γ1 and γ2 subunits. They also showed that GABA and a

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GABA<sub>A</sub> receptor agonist, muscimol, increased Cl<sup>−</sup> uptake in primary cultured renal proximal tubular cells [19]. The authors also reported the expression of β2 and β3 subunits in the renal proximal tube [20], but some of the GABA<sub>A</sub> receptor subunits such as γ<sub>2</sub>, γ<sub>3</sub>, δ, ε, π and θ have not been examined.

GABA may have an important role in the kidney, but a systematic evaluation of GABA-related molecules has not been performed. Therefore, in this report, the expressions of GABA-related molecules, GABA producing enzyme, receptor components, transporters and degrading enzyme, were examined in rat kidney.

Materials and Methods

Animals

All animal experimental procedures were approved by the Fukushima Medical University School of Medicine Animal Committee. Seven-week old Wistar-Kyoto rats (WKY) and spontaneously hypertensive rats (SHR) were obtained from Japan SLC Inc., Sendai, Japan and used for the experiments at 8 weeks of age. Each animal was housed under controlled temperature (21°C) and light conditions (light 7:00–19:00, dark 19:00–7:00) with either 0.3% or 8% NaCl chow (Oriental Yeast Co., Tokyo, Japan) for a week with tap water ad libitum. Rats were anesthetized using pentobarbital (50 μg/kg) intraperitoneal injection. After opening the abdominal cavity and drawing blood from the abdominal aorta, kidneys and brain were harvested and stored for further analyses. The animals were euthanized by pentobarbital overdose (intraperitoneal administration of over 200 mg/kg pentobarbital sodium). Eight rats per group were used for the experiment.

Total RNA extraction and reverse transcription

Total RNA was extracted from the rat kidney cortex and brain preserved in RNAlater (Life Technologies, Carlsbad, CA, USA) using RNeasy plus mini kit (QIAGEN K.K., Tokyo, Japan) following the manufacturer’s instructions. Kidneys from 5 rats were used in the study. Universal RNA from rat normal tissues (BioChain, San Francisco, CA, USA) and brain RNA were used as positive controls. RNA concentration was measured using UV spectrometer, and 0.25 μg of total RNA was reverse transcribed in 20 μl reaction volume using iScript cDNA synthesis kit (Bio-Rad, Hercules, CA, USA). The reverse transcriptase step (5 min at 25°C, 30 min at 42°C, 5 min at 85°C) was carried out using My Cycler (Bio-Rad).

Human RNA samples

Human Kidney Total RNA and Human Brain Total RNA were purchased from Clontech Laboratories, Inc. (Mountain View, CA, USA). The Human Kidney Total RNA was extracted from a normal kidney of a 43-year-old Caucasian female with a sudden death. The Human Brain Total RNA was extracted from a normal whole brain from a 43-year-old Caucasian male with a sudden death. The Human Brain Total RNA was extracted from a normal kidney of a 40-year-old Caucasian female with a sudden death. The Human Brain Total RNA was extracted from a normal whole brain from a 43-year-old Caucasian male with a sudden death. The Human Brain Total RNA was extracted from a normal whole brain from a 43-year-old Caucasian male with a sudden death. The Human Brain Total RNA was extracted from a normal whole brain from a 43-year-old Caucasian male with a sudden death.

Qualitative RT-PCR

Qualitative PCR was performed using the renal cortical mRNA isolated from WKY fed normal chow. Rat GABA<sub>A</sub> receptor α1-6, β1-3, γ1-3, δ, ε, π, θ subunits, GAD65, GAD67, GAT1-3, and ABAT nucleotide sequences were identified in Gen Bank database. PCR primers were designed using Primer Blast (Table 1). One μl of the reverse transcriptase products was amplified by PCR using Go Taq Green Master Mix (Promega, Fitchburg, WI, USA) and 60 nM each of forward and reverse primers. The PCR condition was 2 min at 50.0°C, 10 min at 95.0°C, then 35 cycles of 15 s at 95.0°C and 1 min at 60.0°C, followed by 5 min at 72.0°C. Amplifications for rat GABA<sub>A</sub> R1 and R2 subtypes and GABA<sub>C</sub> p1-3 subunits were carried out using TaqMan Gene Expression Assay (Life Technologies) following the manufacturer’s protocol (Table 1). All PCR reactions were conducted in iCycler (Bio-Rad). Five μl of PCR products were analyzed by electrophoresis in 2.5% TBE Gel (Life Technologies) stained with ethidium bromide, and visualized on a UV transilluminator. Rat brain RNA or commercially available reference RNA were used as positive control.

Quantitative PCR

Quantitative PCR was performed to compare mRNA expression levels in GABA<sub>A</sub> receptor subunits among WKY and SHR fed normal- or high-salt chow. Quantitative PCR was also performed using human kidney and brain RNA commercially available from Clontech Inc. (Mountain View, CA). One μl of reverse-transcription sample was used for real-time quantitative PCR using the iQ<sub>5</sub> Real-Time PCR Detection System and iQ SYBR Green Supermix (Bio-Rad) or StepOnePlus Real-Time PCR System and TaqMan Fast Advanced Master Mix (Life Technologies). PCR reactions were performed in triplicate, and mRNA was quantified based on the Ct value, normalized to GAPDH, and expressed as relative amounts.

Immunoprecipitation

Protein lysate was incubated with Protein G-protein Mag Sepharose (GE Health care Life Sciences, Uppsala, Sweden) clssedlinked to anti-GABA<sub>A</sub> receptor π subunit antibody for 12 hours at 4°C. The immunoprecipitated protein samples eluted from the pellets were suspended in sample buffer and resolved by SDS-PAGE. Western blotting was performed with anti-GABA<sub>A</sub> receptor α1 or β2/3 subunit antibody.

Membrane protein preparation

WKY kidney membrane protein was separated using Mem-PER Eukaryotic Membrane Protein Extraction Kit (Thermo Scientific, Carlsbad, CA) according to the instruction from manufacturer. The hydrophobic fraction containing membrane proteins was diluted to decrease the detergent concentration. The membrane proteins in SDS sample buffer were resolved by SDS-PAGE and detected by immunoblot.

Western Blotting

The rat kidney cortex and brain were immediately snap-frozen in liquid nitrogen and stored at −80°C until use. The tissue was homogenized in lysis buffer (50 mM Tris-Cl pH 8.0, 150 mM NaCl, 0.02% sodium azide and 1% NP-40) containing Complete protease inhibitor mix (Roche, Tokyo, Japan) and centrifuged at 14000 g for 10 min at 4°C. Protein concentrations were determined by BCA protein assay (Thermo Fisher Scientific, Inc., Waltham, MA, USA). Five to 30 μg/lane of protein were separated on NuPAGE Novex 4–12% Bis-Tris Gels (Life Technologies) and transferred onto PVDF membrane (MILLIPORE, Billerica, MA, USA). The membranes were blocked with Immunoblot (DS Pharma Biomedical Co., Ltd., Osaka, Japan). The membranes were probed with a 1:1000 dilution of anti-GABA<sub>A</sub> receptor α1 subunit rabbit polyclonal antibody (MILLIPORE), a 1:500 dilution of β2/3 subunit mouse monoclonal antibody (MILLIPORE), a 1:200 dilution of π subunit rabbit polyclonal antibody (Santa Cruz Biotechnology, Inc, Santa Cruz, CA, sc-25708), p1 antibody (Santa Cruz, sc-25707), p2 antibody
### Immunohistochemistry

Three-μm tissue sections were incubated with primary antibodies including 1:50 to 1:100 dilution of GABA$_A_1$, $\beta_2$, and $\pi$ subunits, GABA$_B$ receptor R1 subtype (Cell-Signaling), R2 subtype (Cell Signaling), GAD67 (Santa Cruz, GAD65 (MILLIPORE) GAT2 (MILLIPORE) or ABAT. Immuno Shot reagents (COSMO BIO Co., LTD, Tokyo, Japan) were used to enhance the immune reaction in some assays. The membranes were washed and probed with 1:5000 goat anti-rabbit or anti-mouse horseradish peroxidase-conjugated IgG (Santa Cruz, Santa Cruz, CA, USA). Proteins were detected by the ECL system (GE Healthcare, Buckinghamshire, UK).

### Results

**Expressions of GABA$_A$ receptor $\alpha_1$, $\beta_3$, $\delta$, $\varepsilon$ and $\pi$ subunits were detected in rat kidney cortex**

Messenger RNAs of GABA$_A$ receptor subunits, $\alpha_1$, $\beta_3$, $\delta$, $\varepsilon$ and $\pi$, were detected by RT-PCR in WKY kidney cortex (Figure 1A). All PCR products showed the theoretical molecular weight and their identities were confirmed by sequencing. Clear single bands were detected for all positive controls (brain or universal RNA). However, mRNAs of GABA$_A$ receptor subunits, $\alpha_2$, $\alpha_3$, $\alpha_4$, $\alpha_5$, $\alpha_6$, $\beta_1$, $\beta_2$, $\gamma_1$, $\gamma_2$ and $\theta$ were not detectable in the kidney cortex. Those genes showing clear mRNA signal, namely, $\alpha_1$, $\beta_3$ and $\pi$ subunits, were tested for protein expression by immunoblotting. The major band for $\alpha_1$ subunit in the kidney was around 55 kDa, which was the same as a band of the brain sample on the Alexa Fluor 488 conjugated goat anti-mouse antibody (Life Technologies). GABA$_A_\pi$ subunit antibody (Abcam: ab26055) was used for immunofluorescence studies. Normal rabbit IgG (Santa Cruz: sc-2027) and mouse monoclonal IgG1 (Abcam: ab81032) were used as negative control. Nuclei were stained by DAPI (Life Technologies). GABA-Related Molecules in the Kidney

### Table 1. Primer information.

| Gene   | Forward primer (3’-5’) | Reverse primer (3’-5’) | Accession No. or assay ID | Amplicon size (bp) |
|--------|------------------------|------------------------|--------------------------|--------------------|
| GABA$_A_1$ | tgcagctctcttcacacagt | tgcagctctcttcacacagt | NM_183326               | 87                 |
| GABA$_A_2$ | agctgtctctcttcacacagt | agctgtctctcttcacacagt | NM_00135779             | 92                 |
| GABA$_A_3$ | cctactgtctgtctgcagctg | cctactgtctgtctgcagctg | NM_017069               | 113                |
| GABA$_A_4$ | cttctgtgtgtctggtggncy | cttctgtgtgtctggtggncy | NM_080587               | 68                 |
| GABA$_A_5$ | gctctgtctgtctgcagctg | gctctgtctgtctgcagctg | NM_017029               | 146                |
| GABA$_A_6$ | agtccagactgacggcagag | agtccagactgacggcagag | NM_021841               | 124                |
| GABA$_B_1$ | ccctctgtgtgtctggtggncy | ccctctgtgtgtctggtggncy | NM_012956               | 90                 |
| GABA$_B_2$ | cttctgtgtgtctggtggncy | cttctgtgtgtctggtggncy | NM_012957               | 96                 |
| GABA$_B_3$ | atcgagctctctccagcttcc | atcgagctctctccagcttcc | NM_017065               | 91                 |
| GABA$_A_1$ | tggctctctccagcttcc | tggctctctccagcttcc | NM_017068               | 91                 |

Abbreviations: GABA; $\gamma$-aminobutyric acid, GAD; glutamate decarboxylase, GAT; GABA transporter, ABAT; 4-aminobutyrate aminotransferase.

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Figure 1. GABA$_A$ receptor subunits are expressed in rat kidney. A) RT-PCR results for GABA$_A$ receptor subunits in rat (WKY on normal diet) kidney cortex. Appropriately-sized bands were detected for GABA$_A$ receptor a1, b3 and p subunits in at least five independent experiments. Rat brain RNA or commercial universal RNA was used as positive control. M: molecular marker. B) Immunoblot for GABA$_A$ receptor a1, b2/3 and p subunits in rat kidney cortex.
kidney cortex. B: brain control, K: kidney cortex. C) Immunofluorescent examination of GABA\textsubscript{A} receptor \(\alpha_1, \beta_3\) and \(\pi\) subunits in rat kidney cortex. Staining with normal IgG instead of primary antibody served as negative control.
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**Figure 2. GABA\textsubscript{A} \(\alpha_1, \beta_3\) and \(\pi\) subunits are co-expressed in rat kidney.** A) Immunofluorescent images of GABA\textsubscript{A} \(\alpha_1, \beta_3\) and \(\pi\) subunits with DAPI counterstain. B) Immunoblot of rat kidney cortex protein immunoprecipitated with GABA\textsubscript{A} receptor \(\pi\) subunit antibody. Molecular sizes of \(\alpha_1\) and \(\beta_3\) subunits around 110 kDa suggest dimerization. C) Immunoblot of rat kidney cortex membrane protein for GABA\textsubscript{A} receptor \(\pi\) subunit.
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Immunoblotting using an antibody against β2/3 subunit showed a similar pattern in the brain and kidney, with a major band around the theoretical 55 kDa and a secondary band around 40 kDa. The kidney may express similar or greater amounts of β3 subunit than the brain. Immunoblotting for π subunit detected a major band around 50 kDa for both brain and kidney tissue, with a weak secondary band slightly larger than the major band in the brain and smaller bands for kidney samples (Figure 1B). Then, to elucidate the protein localization, immunofluorescence study was performed for α1, β3 and π subunits (Figures 1C and 2A). Alpha 1 subunit distributed widely in the cortical tubules, especially proximal-like tubules. Specific staining of β3 subunit was observed in the apical region of the proximal-like tubules with tall cells and a brush border-like structure and also in the apical region of distal-like tubules. No staining for β3 subunit was detected in the glomeruli. π subunit staining was present mostly in the distal-like tubules and less in the proximal-like tubules (Figure 1C).

**Figure 3. GABAA receptor R1 and R2 subtypes are expressed in rat kidney.** A) RT-PCR results for GABAA receptor subtypes in rat kidney cortex. Appropriately-sized bands were detected for GABAA receptor R1 and R2 subtypes in at least five independent experiments. Rat brain RNA or commercial universal RNA was used as positive control. M: molecular marker. B) Immunostaining for GABAA receptor R1 and R2 subtypes in rat kidney cortex. Staining with normal IgG instead of primary antibody served as negative control.
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**GABA<sub>A</sub> receptor α1, β3 and π subunits co-localized in the apical membranes of cortical tubules**

By immunofluorescent observation, co-staining of α1, β3 and π subunits was observed mainly on the apical side of cortical tubules (Figure 2A). Kidney protein immunoprecipitated with π subunit antibody revealed α1 and β3 subunits, and the molecular weights of blotted α1 and β3 subunits were close to 110 kDa, suggesting
in the collecting-like tubules, similar to GABA<sub>Α</sub> receptor π subunit, and moderate staining on the apical side of proximal-like tubules (Figure 3B).

GABA<sub>C</sub> Receptor p1 and p2 subunits were detected in rat kidney cortex

Messenger RNAs of GABA<sub>C</sub> receptor subunits p1 and p2 were detected by RT-PCR in WKY kidney cortex, but p3 was not detectable (Figure 4A). Immunoblotting for p1 subunit showed a similar pattern in the brain and kidney, with a major band around the theoretical 48 kDa. The kidney may express similar or greater amounts of p1 subunit than the brain. Immunoblotting for p2 subunit showed a major band around 50 kDa in both brain and kidney tissues (Figure 4B).

Two types of GABA-producing enzymes were detected in rat kidney cortex

Both types of GABA producing enzyme, GAD65 and GAD67 mRNA, were clearly detected in the kidney cortex (Figure 5A). By immunohistochemistry, GAD65 staining was mostly observed in glomeruli and arterioles. Some weak staining of cortical tubules and stronger staining of vessels and glomeruli were observed for GAD67 (Figure 5B).

One type of GABA transporter, GAT2, was expressed in rat kidney cortex

Messenger RNAs of peripheral-type GABA transporter GAT2, were detected, but not of GAT1 and GAT3 (Figure 6A). To confirm the protein localization, immunohistochemistry was performed for GAT2. Specific staining was observed on the basolateral side of cortical renal tubules (Figure 6B).

GABA degrading enzyme was detected in kidney cortex

Messenger RNA of GABA degrading enzyme ABAT was detected by RT-PCR (Figure 7A) in the kidney cortex using the primer set that amplifies all three known splice variants of the gene. Immunohistochemistry confirmed the protein expression of ABAT with the major band around 50 kDa for both brain and kidney, and a secondary band larger than the major band in the kidney (Figure 7B). Immunohistochemistry revealed that ABAT staining pattern was also similar to those of GABA<sub>Α</sub> π subunit, present mostly in the distal-like tubules (Figure 7C).

Renal GABA<sub>Α</sub> receptor expression in hypertensive model rats with salt loading

To explore the possibility of GABA receptor alterations in hypertensive and/or salt-sensitive phenotype, we performed quantitative mRNA analysis and western blots of GABA<sub>Α</sub> subunits using the kidney cortex of WKY and SHR on normal- and high-salt diet. GABA<sub>Α</sub> receptor α1 and β3 subunit expressions in the kidney were not affected by stain or salt loading (Figure 8). In contrast, renal GABA<sub>Α</sub> δ subunit (Figure 8) mRNA expression was significantly lower in SHR than in WKY on normal salt diet. The level of GABA<sub>Α</sub> δ subunit mRNA in SHR kidney was about half of that in WKY kidney. On high-salt diet, GABA<sub>Α</sub> δ subunit mRNA expression tended to be lower in the SHR kidney compared to WKY kidney, but the difference was not significant. Salt loading did not significantly alter renal GABA<sub>Α</sub> δ subunit mRNA expression in WKY and SHR. We also performed western blot analysis for GABA<sub>Α</sub> δ subunit (WKY normal salt 100±12%, SHR normal salt 121±20%, WKY high salt 100±8%, SHR high salt 95±7%, relative expression normalized to actin, n = 8 rats/group). In contrast to the mRNA difference in WKY and SHR.
kidneys for GABA\(_\alpha\) \(\pi\) subunit, we found no significant difference in the expression level GABA\(_\alpha\) \(\pi\) subunit in the kidney cortex of four groups of rats, WKY and SHR with and without salt loading on the protein level.

**Expressions of GABA-related molecules in human kidney**

Next, we examined species differences on the expression of GABA-related molecules in the kidney using human kidney and brain RNA samples. All of the GABA-related molecules consistently detected in rat kidney, specifically, GABA\(_\alpha\) \(\alpha1\), \(\beta3\) and \(\pi\) subunits, GABA\(_\beta\) R1/R2 subtypes, GABA\(_\gamma\) subunits \(\rho1\) and \(\rho2\), GAD65 and 67, GAT2, and ABAT were also found in human kidney (data not shown).

**Discussion**

In this study, gene expressions of GABA-related molecules were examined in rat kidney cortex. Expressions of GABA\(_\alpha\) \(\alpha1\), \(\beta3\), \(\pi\) receptor subunits, GABA\(_\beta\) receptor R1 and R2 subtypes, GABA\(_\gamma\) \(\rho1\) and \(\rho2\) receptor subunits were found at both mRNA and protein levels. Enzymes involved in GABA synthesis, GAD67 and GAD65, GABA transporter, GAT2, and GABA metabolizing enzyme, ABAT, were also expressed in rat kidney.

Our results for GABA\(_\alpha\) receptor subunits differ in some extent to those previously reported by Sarang et al. They found the expressions of \(\alpha1\), \(\alpha5\), \(\beta1\), \(\beta2\), \(\beta3\), \(\gamma1\), and \(\gamma2\) subunits but not of \(\alpha2\), \(\alpha4\), or \(\delta\) subunits in the kidney cortex from Sprague-Dawley rats [20]. In addition, they found the expression of \(\beta2\) and \(\beta3\) subunits in microdissected rabbit proximal tubule S2 segment. We found the expression of \(\alpha1\) and \(\beta3\) subunits similar to Sarang, et al., but we also found the renal expression of \(\pi\) subunit, which was not examined in their reports. Negative expression results for \(\alpha5\), \(\beta1\), \(\beta2\), \(\gamma1\), and \(\gamma2\) in our study may be due to either the difference in the detection methods, as Sarang et al used nested PCR whereas our method utilized single amplification, or the difference in rat strain, as we used WKY instead of Sprague-Dawley.

Our report is the first to show a concrete expression of GABA\(_\alpha\) receptor \(\pi\) subunit in rat and human kidney. To explore the differential expression, Nephromine, a web-based analysis engine freely accessible to academic users at http://www.nephromine.org, was used to study molecular expression in human renal diseases. A search for differential expression of GABA\(_\alpha\) \(\pi\) subunit mRNA in human kidney using Nephromine revealed that it may

**Figure 5. GABA-producing enzymes are expressed in rat kidney.** A) RT-PCR results for GAD65 and GAD67 in rat kidney cortex. Appropriately-sized bands were detected for GAD65 and GAD67 in at least five independent experiments. M: molecular marker. B) Immunostaining for GAD65 and GAD67 in rat kidney cortex. Staining with normal IgG instead of primary antibody served as negative control.

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be overexpressed approximately 2 fold in kidneys with diabetic nephropathy (n = 10) compared to those from healthy donors (n = 12). This difference was seen both in tubulointerstitium (fold change: 2.003, P-value: 4.99E-4) and glomeruli (fold change: 2.392, P-value: 0.008, data source [21]). However, GABAA pi subunit expression in hypertension data set did now show a significant difference between normal tissues from tumor nephrectomy (n = 4) and those with nephrosclerosis (n = 14, data source [22]).

We also found the expression of GABAB receptor R1 and R2 subtypes and GABAC receptor r1 and r2 subunits. As far as we know, this is the first report of GABAB and GABAC receptor protein expressions in the kidney.

It has been shown that GABAA receptor agonist muscimol and GABAB agonist baclofen increased fractional excretion of water and sodium in isolated rat kidney perfusion [16]. Our results suggest the possibility of a novel combination of GABAA receptor subunits, α1, β3 and π, in the kidney, and this combination may be relatively kidney-specific. Muscimol is a potent agonist of GABAC in addition to GABAA receptor [23]. When searching for kidney-specific modulation methods of GABAA receptor, it may be possible in the future to screen for substances that show specific agonistic activity on GABAA receptors with π subunits. Such findings may promote the development of tubule-specific natriuretic agent with less effect on the nervous system. Meanwhile, the function of renal GABAB receptor is not known. As GABAB receptor R1 subtype localizes to glomeruli and arterioles as well as tubules, it may control intra-glomerular perfusion pressure. Also, R2 subtype observed in the distal-like tubules may influence the effects of aldosterone or vasopressin as their co-expression in the principal cells is likely. It has been reported that intrarenal administration of baclofen, a GABAB receptor agonist, has renoprotective effect against ischemia reperfusion injury through the attenuation of renal sympathetic nerve activity [24]. Since overactivation of renal sympathetic nerve activity is associated with tissue renin-angiotensin-aldosterone system overactivity, GABAB activation in renal tubule may alter this phenomenon. Also, the activation of postsynaptic GABAB receptors enhances

Figure 6. GABA transporters are expressed in rat kidney. A) RT-PCR results for GAT1, GAT2 and GAT3 in rat kidney cortex. Appropriately-sized bands were detected for GAT2 in at least five independent experiments. Rat brain RNA or commercial universal RNA was used as positive control. M: molecular marker. B) Immunostaining for GAT2 in rat kidney cortex. Staining with normal IgG instead of primary antibody served as negative control. doi:10.1371/journal.pone.0105835.g006
GABA currents [25]. Chloride handling in distal tubule and collecting duct modulated by GABA-A Cl\(^{-}\) channel may be associated with sodium chloride cotransporter and/or epithelial sodium channel. Further study is required to depict a better picture and the precise mechanism of action on GABA receptors expressed in the kidney.

In addition to the GABA receptors, GABA-related enzymes and transporters were also examined in this study. The expression and activity of GAD in the kidney have been reported in the past. Liu et al. found GAD antigen in mouse proximal and distal tubules but not in glomeruli [26]. However, previous reports did not examine GAD subtypes separately. In this study, we found both GAD67 and GAD65 in the kidney tissue using antibodies specific for each subtype.

For GABA transporters, a high level of human GAT2 was found in the kidney and low levels in the brain and lung [27]. Rat GAT2 has been found not only in the brain and retina, but also in the liver, kidney, and heart [28]. GAT1 and GAT3 were not detected in rat kidney, similar to our results [28]. ABAT expression has been demonstrated in rat kidney, and the molecular mass of kidney ABAT was similar to that of liver ABAT and smaller than that of brain ABAT [29].

We have detected renal expressions of major GABA-related molecules, namely receptors, transporters, and synthesizing and metabolizing enzymes. All three GABA\(\alpha\) subunits examined in this study, \(\alpha_1\), \(\beta_3\), and \(\pi\), showed staining in the renal tubules. Although the staining patterns are not identical, the presence of all three subunits in the same segment is probable, and this suggests that there may be functional receptor composed of \(\alpha_1\beta_3\pi\) subunits, as GABA\(\alpha\) receptors formed by \(\alpha_5\beta_3\pi\) combination has shown to be functional in transfection experiments [1]. We also found expressions of GAD67 and GAD65 in the renal tubules, which suggest the production of GABA in the kidney. Dopamine in the kidney is a locally-functional, autocrine/paracrine factor known to play a role in the pathophysiology of hypertension [30,31]. Proximal convoluting tubules produce dopamine which.

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**Figure 7. GABA degrading enzyme is expressed in rat kidney.** A) RT-PCR results for ABAT in rat kidney cortex. Appropriately-sized bands were detected for ABAT in at least five independent experiments. Rat brain RNA or commercial universal RNA was used as positive control. M: molecular marker. B) Immunoblotting for ABAT in rat kidney cortex. B: brain control, K: kidney cortex. C) Immunostaining for ABAT in rat kidney cortex. Staining with normal IgG instead of primary antibody served as negative control. doi:10.1371/journal.pone.0105835.g007
acts on mainly D1-like receptors in the proximal convoluting tubules to induce natriuresis. Renal dopamine synthesis and release is increased by dietary sodium intake and elevation in intracellular sodium concentration, and locally-secreted dopamine in the kidney is responsible for over 50% of sodium excretion during the state of sodium surfeit. Since the current study revealed that rat renal cortex expresses major molecules to account for the entire life cycle of GABA, locally-secreted GABA, similar to dopamine, may act in an autocrine/paracrine fashion in the kidney to induce natriuresis and to lower blood pressure.

In conclusion, rat kidney was found to harbor all major GABA-related components. This suggests the possible existence of a local GABAcergic system with an intra-renal GABA production, secretion and function in an autocrine/paracrine manner. Further study is necessary to elucidate the precise localization and function of these renal GABA system molecules. As GABA has been suggested to elicit natriuresis and to lower blood pressure, the research on GABA-related molecules in the kidney is promising as a source of new important information on the pathophysiology of hypertension.

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Author Contributions
Conceived and designed the experiments: KT MSY JY. Performed the experiments: KT MSY JY AA YS. Analyzed the data: KT MSY JY AA YS HS TW JK. Contributed reagents/materials/analysis tools: KT MSY JY. Contributed to the writing of the manuscript: KT MSY JY AA YS HS TW JK.

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